
straditize Documentation

Release 0.1.3

Philipp Sommer

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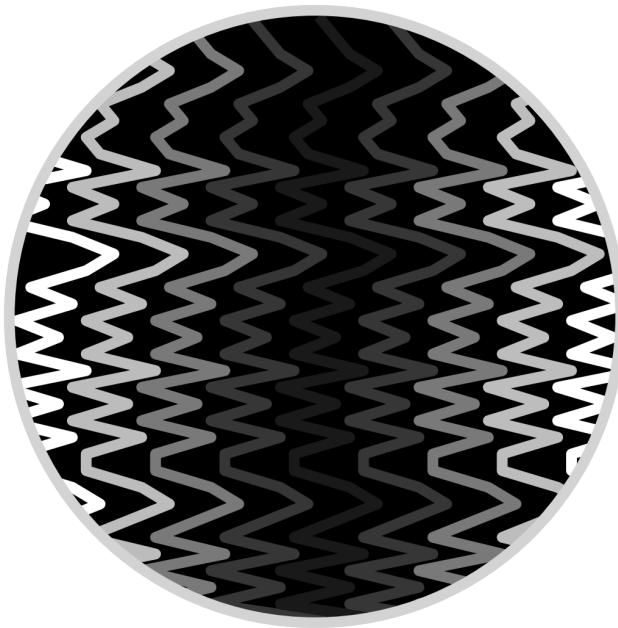
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STRADITIZE (Stratigraphic Diagram Digitizer) is an open-source program that allows stratigraphic figures to be digitized in a single semi-automated operation. It is designed to detect multiple plots of variables analyzed along the same vertical axis, whether this is a sediment core or any similar depth/time series.

Usually, in an age of digital data analysis, gaining access to data from the pre-digital era – or any data that is only available as a figure on a page – remains a problem and an under-utilized scientific resource.

This program tackles this problem by providing a python package to digitize especially pollen diagrams, but also any other type of stratigraphic diagram.

The package is very new and there are many features that will be included in the future. Therefore we would be very pleased to get feedback! To do so, you can contact us or raise an issue on [GitHub](#).

DOCUMENTATION

1.1 About straditize

1.1.1 Why straditize?

In an age of digital data analysis, gaining access to data from the pre-digital era – or any data that is only available as a figure on a page – remains a problem and an under-utilized scientific resource. Whilst there are numerous programs available that allow the digitization of scientific data in a simple x-y graph format, we know of no semi-automated program that can deal with data plotted with multiple horizontal axes that share the same vertical axis, such as pollen diagrams and other stratigraphic figures that are common in the Earth sciences. STRADITIZE (Stratigraphic Diagram Digitizer) is a new open-source program that allows stratigraphic figures to be digitized in a single semi- automated operation. It is designed to detect multiple plots of variables analyzed along the same vertical axis, whether this is a sediment core or any similar depth/time series.

1.1.2 About the author

The code and GUI of straditize was developed by Philipp S. Sommer at the Institute of Earth System Dynamics (IDYST) at the University of Lausanne as part of the SNF funded [HORNET Project](#) (200021_169598).

The other contributors are [Basil A. S. Davis](#), [Manuel Chevalier](#) and Dilan Rech who made significant contributions to the layout, workflow, beta tests and and reviewing of the software.

1.1.3 License

straditize is published under the [GNU General Public License v3.0](#) under the copyright of Philipp S. Sommer, 2018-2019

1.2 Installation

1.2.1 How to install

You can either install straditize through a package manager such as [conda](#) or [pip](#) or install it [from source](#).

Installation using conda

We highly recommend to use [conda](#) for installing straditize. Here you can install it via manually via the [chilipp channel](#)

After having downloaded and installed [anaconda](#), open a terminal (or the *Anaconda Prompt* on windows) and install straditize from the [conda-forge channel](#). You can choose: We recommend to install straditize into its own environment via:

```
$ conda create -n straditize -c conda-forge straditize
```

and then activate this environment via:

```
$ conda activate straditize
```

In that way you do not mess up your base environment. Nevertheless you can also install it into an existing environment via:

```
$ conda install -c conda-forge straditize
```

In the same terminal, now type `straditize` to start the software.

Note: The latest master branch on github is always available under the `master` label on the [chilipp channel](#). Just type:

```
$ conda install -c chilipp/label/master straditize
```

to install the latest version from the master branch. Note that you then have to add the *conda-forge* channel to your default channels via:

```
$ conda config --add channels conda-forge
```

Installation using pip

If you do not want to use conda for managing your python packages and already have python3 installed on your computer, you can also use the python package manager `pip`. To be on the safe side, make sure you have the *Dependencies* installed. If so, open a terminal and install it via:

```
$ pip install straditize
```

To open the software, type `straditize` in the same terminal.

Installation from source

To install it from source, make sure you have the *Dependencies* installed. Download (or clone) the [github](#) repository, e.g. via:

```
git clone https://github.com/Chilipp/straditize.git
```

and install it via:

```
pip install . # or python setup.py install, but pip is recommended
```

from your terminal. To open the software, type `straditize` in the same terminal.

1.2.2 Dependencies

Required dependencies

straditize has been tested for python>=3.6. Furthermore the package is built upon multiple other packages, mainly

- `psyplot-gui`>1.2.0: The graphical user interface for psyplot
- `PyQt5`: Pythons Qt bindings that are required by psyplot-gui (note that PyQt4 is **not** supported!)
- `numpy`, `scipy` and `pandas`: for the data management and computations
- `matplotlib`>=2.0: The python visualiation package
- `pillow`: for reading and writing images
- `scikit-image`: For image recognition features
- `openpyxl`: For exports to Excel files
- `netCDF4`: A library for saving and storing netCDF files.

Optional dependencies

We furthermore recommend to use

- `tesserocr`: for column names recognition. It depends on the `tesseract` OCR and you can install both (on Linux and MacOS) via:

```
$ conda install -c chilipp tesserocr
```

(see *Automatic optical character recognition (OCR)* for more information)

1.2.3 Running the tests

We use `pytest` to run our tests. So you can either run clone out the `github` repository and run:

```
$ python setup.py test
```

or install `pytest` by yourself and run

```
$ py.test
```

Alternatively you can build the recipe in the `conda-recipe` directory via

```
$ conda build conda-recipe
```

which will also run the test suite.

Warning: Running the entire test suite in one single process (such as `python setup.py test`) might be quite memory consumptive because it involves the creation and closing of many PyQt widgets and unfortunately some memory is leaked from one test to another. Therefore we recommend to split the tests into multiple processes, e.g.:

```
# run the test suite but ignore some modules
python setup.py test -a '--ignore=tests/widgets/test_selection_toolbar.py --
→ignore=tests/widgets/test_samples_table.py --ignore=tests/widgets/test_beginner.
→py --ignore=tests/widgets/test_hoya_del_castillo.py'
# run the tests for the previously ignored modules
python setup.py test -a 'tests/widgets/test_selection_toolbar.py
tests/widgets/test_samples_table.py'
python setup.py test -a 'tests/widgets/test_beginner.py'
1.2. Installation
python setup.py test -a 'tests/widgets/test_hoya_del_castillo.py'
```

or equivalently with `py.test` instead of `python setup.py test -a`. Note that `conda build conda-recipe` already splits the session into multiple processes.

Nevertheless, you should expect about ~180 tests to be ran and a total memory usage of about 3 to 4GB RAM.

1.2.4 Building the docs

The online documentation is accessible as PDF, HTML and Epub under <https://straditiz.readthedocs.io> or <https://straditiz.rtd.io>. Thanks to the free services by readthedocs.org, the online documentation is build automatically after each commit to the [github](#) repository.

To build the docs locally on your machine, check out the [github](#) repository and install the requirements in '`docs/environment.yml`' and the `sphinx_rtd_theme` package. The easiest way to do this is via anaconda by typing:

```
$ conda env create -f docs/environment.yml  
$ conda activate straditiz_docs  
$ conda install sphinx_rtd_theme
```

Then build the docs via:

```
$ cd docs  
$ make html # or `make pdf` for a PDF version compiled with Latex
```

1.2.5 Updating straditiz

Updating the software depends on how you installed it on your system.

Updating via conda

If you installed straditiz via conda (see [Installation using conda](#)), you can update it via:

```
$ conda update -c chilipp straditiz
```

Updating via pip

If you installed it via `pip` (see [Installation using pip](#)), you can update it via:

```
$ pip install -U straditiz
```

Updating from source files

If you installed it via `python setup.py install` from the source repository (see [Installation from source](#)), just run that command again after having checked out the latest version from `github`.

1.2.6 Uninstallation

The uninstallation depends on the system you used to install straditiz. Either you did it via `conda` (see [Uninstallation via conda](#)), via `pip` or from the `source files` (see [Uninstallation via pip](#)).

Anyway, if you may want to remove the psyplot configuration files. If you did not specify anything else (see `psyplot.config.rcsetup.psyplot_fname()`), the configuration files for psyplot are located in the user home directory. Under linux and OSX, this is `$HOME/.config/psyplot`. On other platforms it is in the `.psyplot` directory in the user home.

Uninstallation via conda

If you installed straditize via `conda`, simply run:

```
conda uninstall straditize
```

Uninstallation via pip

Uninstalling via pip simply goes via:

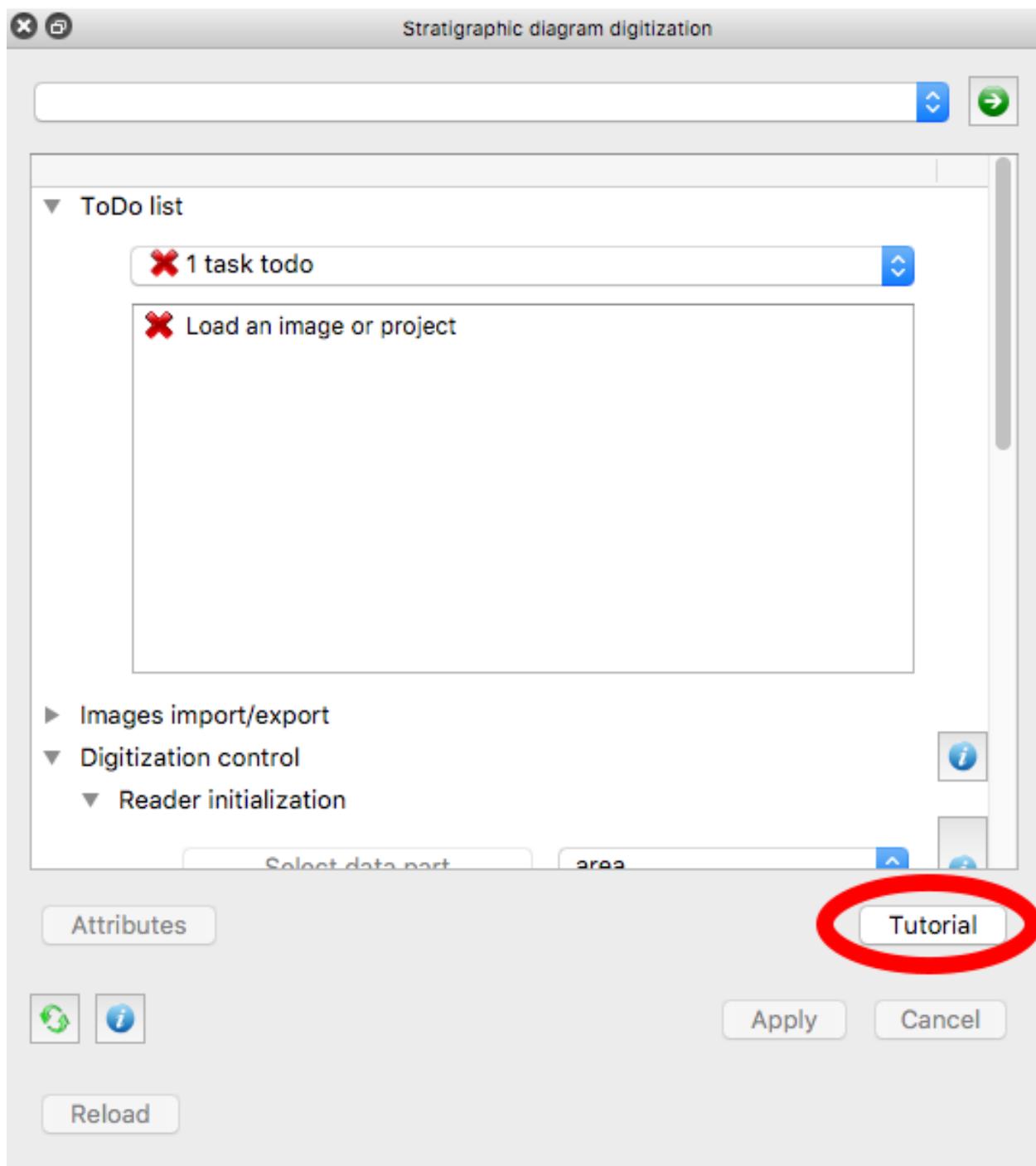
```
pip uninstall straditize
```

Note, however, that you should use `conda` if you also installed it via conda.

1.3 Straditize Tutorial

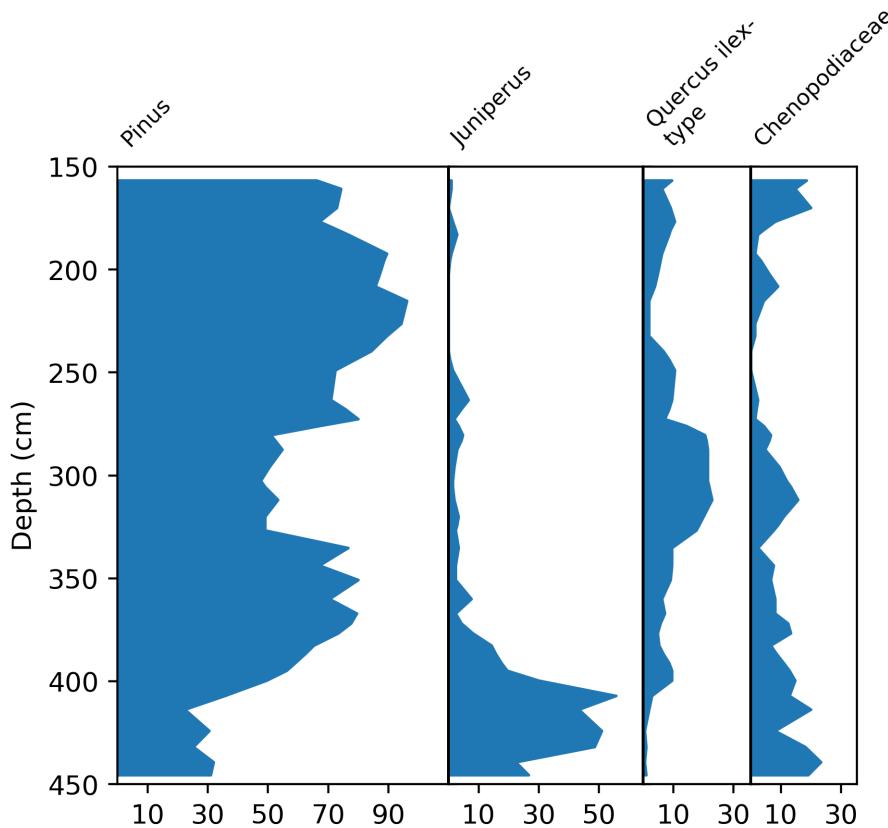
To introduce you into straditize, we implemented a tutorial into the graphical user interface.

To start it in the GUI, click the *Tutorial* button at the bottom of the straditizer control.



1.3.1 Tutorials

Beginners tutorial to introduce straditize



In this tutorial, we digitize a simple small pollen diagram a subset of the Hoya del Castillo dataset from [Basil2007a]. This tutorial will introduce you into the main parts of straditize and the basic workflow.

At the bottom of the straditizer control, you find a navigation panel which helps you to navigate through the tutorial. Click the *Next* button in this control to start the tutorial. You can skip steps by clicking the *Skip* button and you can check what you have done using the *Check* button and then proceed with the next step.

The tutorial will take between 5 and 10 minutes. You can save your current state to a file using *File → Save → Save straditizer* (see [Saving and loading your project](#)) and reload it later via *File → Open project → Open straditizer → Project or image*

If you have questions or troubles with this tutorial, please open an issue on

<https://github.com/Chilipp/straditize/issues>

and we will do our best to assist you.

The straditize workflow

The straditize workflow, i.e. the program for the next few minutes consists of 6 main steps:

1. Load the diagram
2. Select the diagram part (without x-axes labels, y-axes labels, etc.)
3. Clean the diagram part. Only the data parts should be left over, i.e. no y-axes, lines, letters, etc.

4. Digitize the diagram
5. Find and edit the samples
6. Export the data to Excel or CSV

Most of these tasks can be done in an automatic way but you should always review and edit the outcome to make sure that what you do is scientifically reliable.

The interface to these semi-automatic steps is the straditzer control

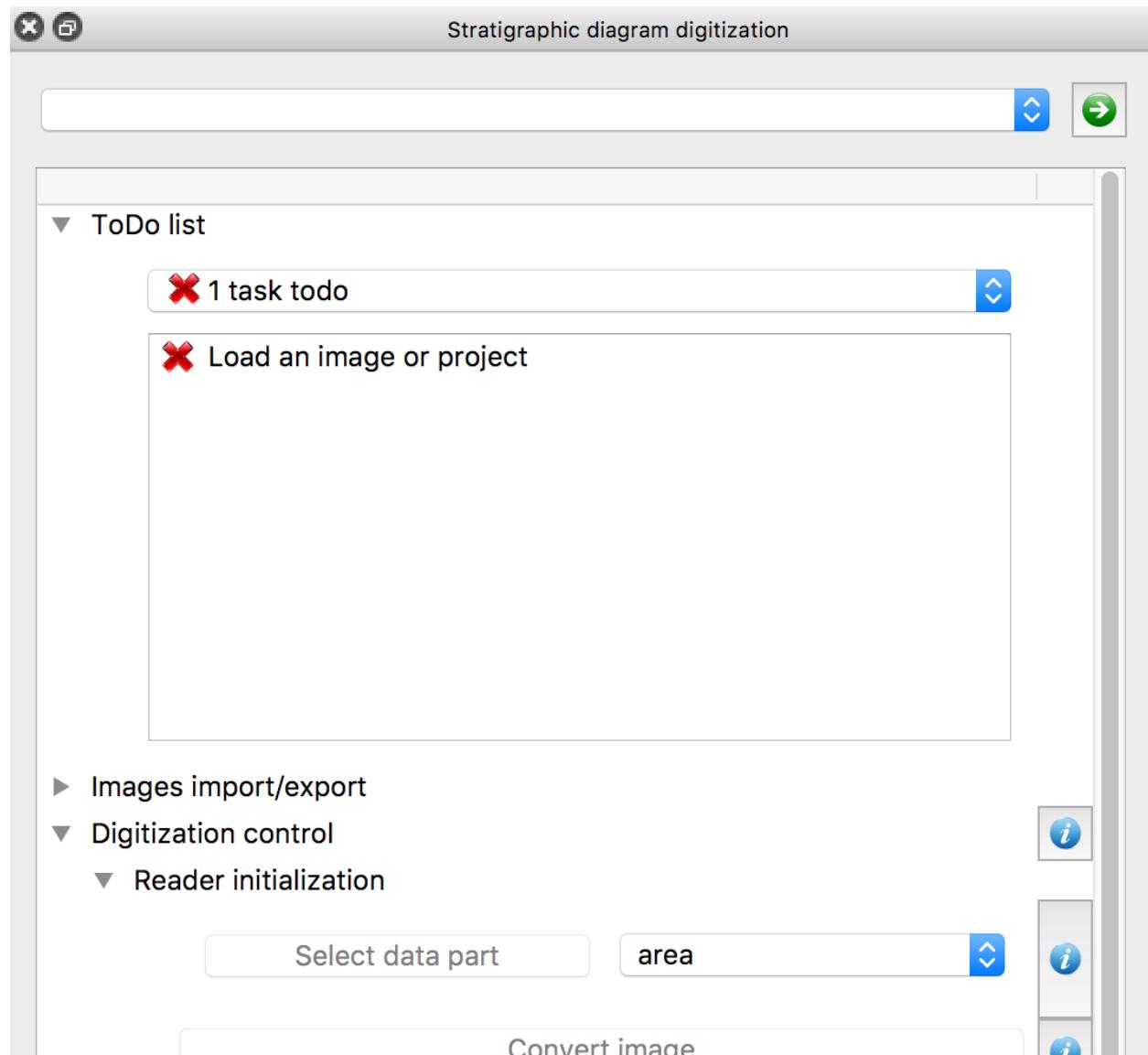


Fig. 1: Click the image to enlarge it

You can expand the submenus using the ► buttons in the control and you can access the user manual using the buttons on the right side.

Before we start with the diagram, we will highlight some of the sections in the GUI:

The ToDo list The list at the top of the control guides you through the *straditization* process. It shows the open tasks you should consider.

Most of them will be marked as done automatically, but if you think you are done with one of the tasks, just right-click the task and mark it as done.

The Plot control This section at the bottom of the control gives you some visual explorations of your diagram and the digitization process. You should, whenever possible, use its functionalities, most of the errors can be spotted visually

The Help explorer This separate widgets can be used to access the user manual and provides guidance for the different functionalities. This should always be your first point of reference if you have any problems.

Load the image of the diagram

The first step is to load the image of the diagram into straditize.

1. Click the  button at the top of the straditizer control.
2. Select the image file `beginner-tutorial.png` from your hard-disk and click Open

Navigation inside the plot

Now you see the stratigraphic diagram (by default) centered at the top of the graphical user interface. Additionally you find a separate window to its right with a magnified version of the diagram. It follows your mouse movements inside the diagram and shows you a magnified version of it.

To navigate inside the plot, you can use the default navigation toolbar that is provided by matplotlib



Especially the Pan/Zoom button  and the zoom-to-rectangle button  are of interest for you. You can enable and disable them by clicking on the corresponding button in the toolbar.

The [matplotlib docs](#) provide further guidance:



The Pan/Zoom button This button has two modes: pan and zoom. Click the toolbar button to activate panning and zooming, then put your mouse somewhere over an axes. Press the left mouse button and hold it to pan the figure, dragging it to a new position. When you release it, the data under the point where you pressed will be moved to the point where you released. If you press 'x' or 'y' while panning the motion will be constrained to the x or y axis, respectively. Press the right mouse button to zoom, dragging it to a new position. The x axis will be zoomed in proportionately to the rightward movement and zoomed out proportionately to the leftward movement. The same is true for the y axis and up/down motions. The point under your mouse when you begin the zoom remains stationary, allowing you to zoom in or out around that point as much as you wish. You can use the modifier keys 'x', 'y' or 'CONTROL' to constrain the zoom to the x axis, the y axis, or aspect ratio preserve, respectively.

With polar plots, the pan and zoom functionality behaves differently. The radius axis labels can be dragged using the left mouse button. The radius scale can be zoomed in and out using the right mouse button.

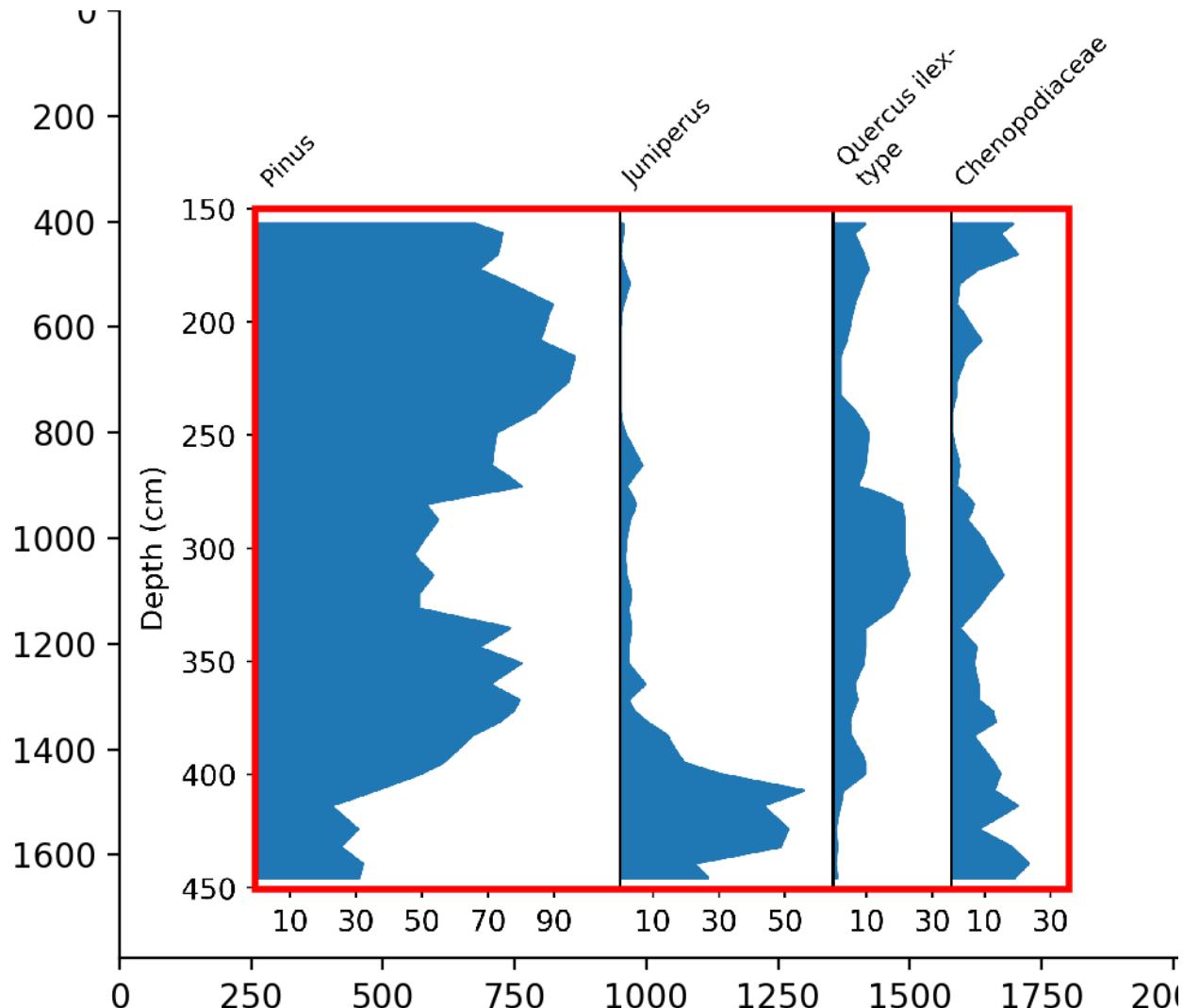


The Zoom-to-rectangle button Click this toolbar button to activate this mode. Put your mouse somewhere over an axes and press a mouse button. Define a rectangular region by dragging the mouse while holding the button to a new location. When using the left mouse button, the axes view limits will be zoomed to the defined region. When using the right mouse button, the axes view limits will be zoomed out, placing the original axes in the defined region.

More information can be found in the matplotlib documentation: https://matplotlib.org/users/navigation_toolbar.html

Select the diagram part

The second step involves selecting the diagram part. This is the part where your data is displayed, but without the vertical and horizontal axes descriptions.



Being exact in this step significantly reduces the later work and simplifies the automatic digitization.

1. Click the *Select data part* button in the digitization control.
2. Straditize automatically recognizes the data part. But you can change this by moving the crosses around with your mouse (see the note below).
3. Click the *Apply* button at the bottom
4. Done! You now see a red rectangle in your diagram the marks the data part. You can hide it using the *Plot control* section in the straditizer control. For the sake of this tutorial, expand the *Plot control* section and remove the rectangle by clicking the  icon for the *Diagram part*

Note:

These marks are very common in straditize. In general, you can left-click on a mark to move it around, right-click to delete it, and Shift+left-click on the plot to create a new one.

Note that you need to disable the  and  tools in the navigation to interact with the marks, by clicking on the corresponding button!

If you want to change the appearance of the marks, see the *Marker control* section in the straditizer control panel.

Create the diagram reader

Now you have to select, which type of diagram you are digitizing. In the example, it is an area diagram which is already selected in the dropdown menu next to the *Select data part* button.

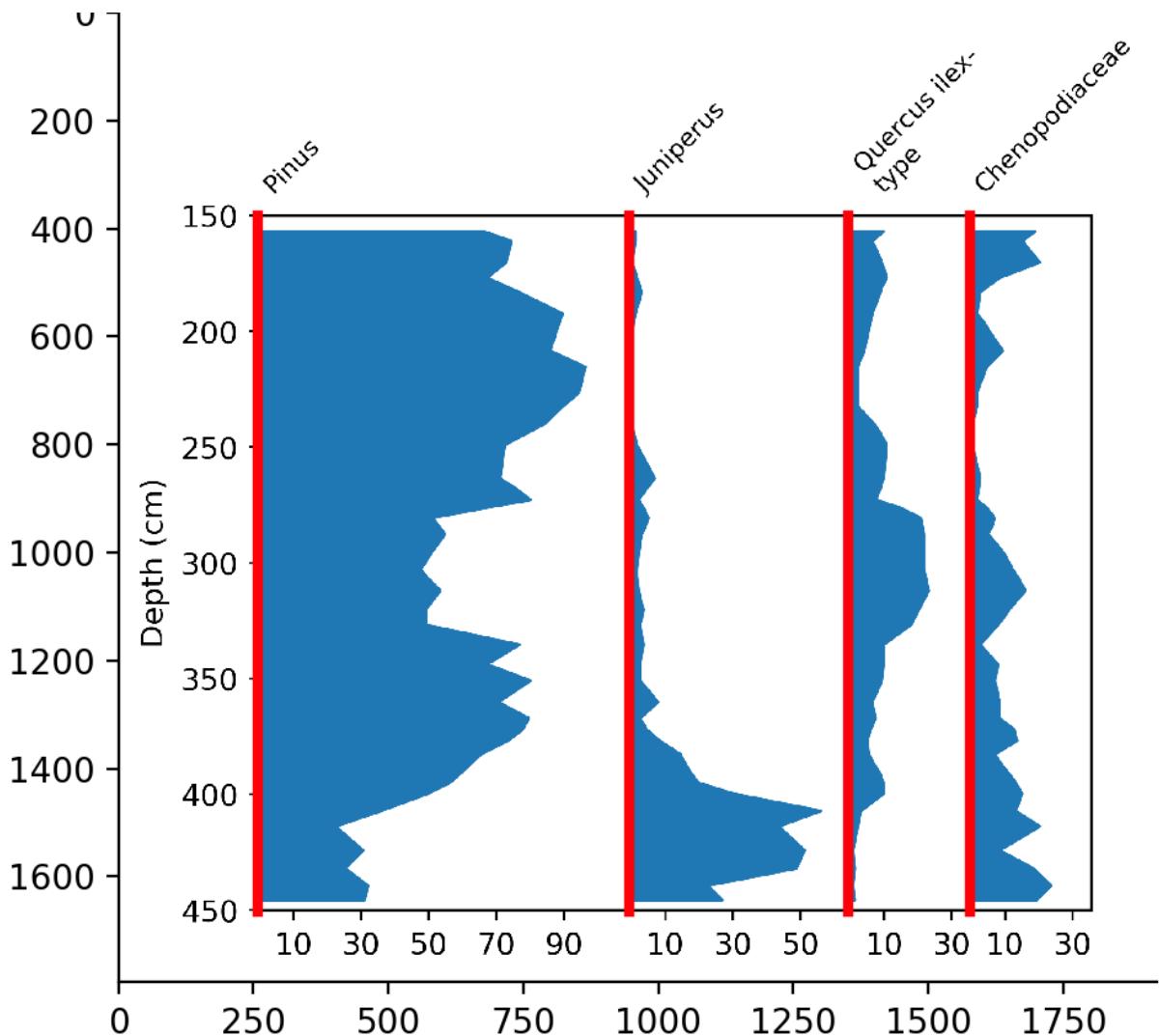
Click on *Convert image* to start the digitization.

See also:

[Selecting the reader](#)

Selecting the columns

The next step is, to separate the columns, i.e. to tell straditize where each of the sub diagrams start.



This is simple in our case, since straditiz can separate them automatically.

1. Click the *Column starts* button. You will see several vertical blue lines appearing on the plot that you can drag and drop such as you did it when selecting the diagram part. You can change their colors using the *marker control*
2. straditiz recognizes these columns automatically and in our example, you do not have to edit them. Therefore hit the *Apply* button at the bottom of the control and you are done.

See also:

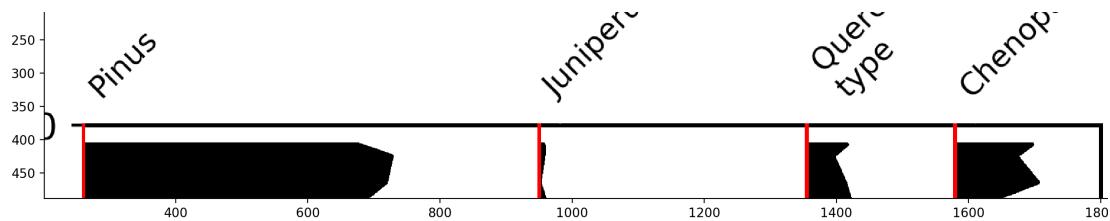
Select the starts for each column

Cleaning the image

The next step is to clean the data image to make sure that everything that is not representing real data is removed. In our case, these are y-axes, x-axes and the outer frame. You can also use an external image editing software such as Adobe Photoshop, but we will do this now inside straditiz and use some of the automatic recognition functionalities.

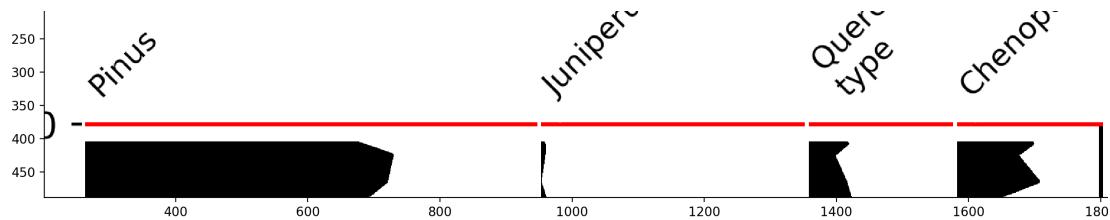
1. Expand the *Remove features* tab

2. Click the y-axes button, the y-axes in the plot will be highlighted



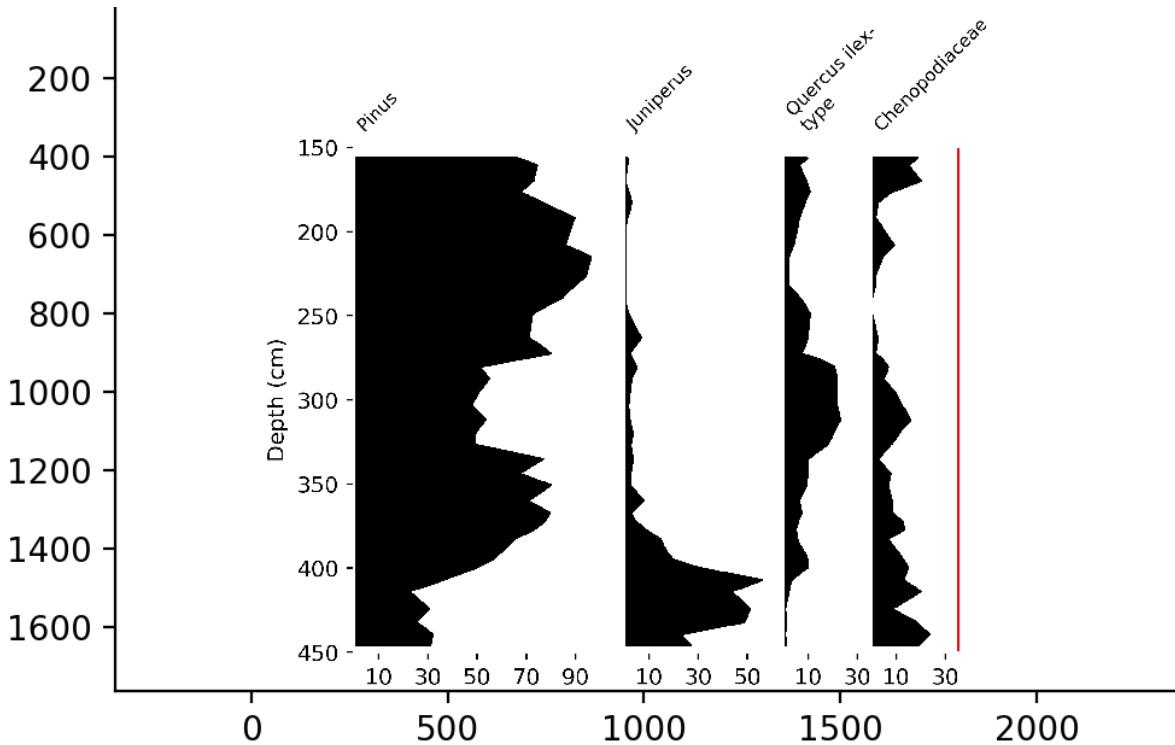
3. Click *Remove* to remove them

4. 2. Click the x-axes button, the x-axes in the plot will be highlighted



5. Click *Remove* to remove them

6. Finally there is the right part of the diagram frame left

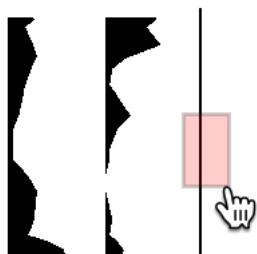


For this, we use the selection toolbar:

- i. from the menu, select the *column selection* tool (click and hold the button to the right of the button in the selection toolbar to open the menu)



- ii. Draw a rectangle around the right line



and make sure you have the mode activated

- iii. Click the *Remove* button when you are done

Digitizing the data

The next step, after we cleaned up the image, is the digitization of the diagram. Click the *Digitize* button.

You now see the lines that result from the digitization. You can remove them in the *Plot control* section by clicking the button of the *Full digitized data* row.

Finding and editing samples

So far, we have one data point per pixel in the image. However, we have to identify the locations of the samples in order to reproduce the original data.

Straditize assists you with this through automatic sample finding algorithms, or you can load the sample locations from an external files or you just add and edit the samples manually.

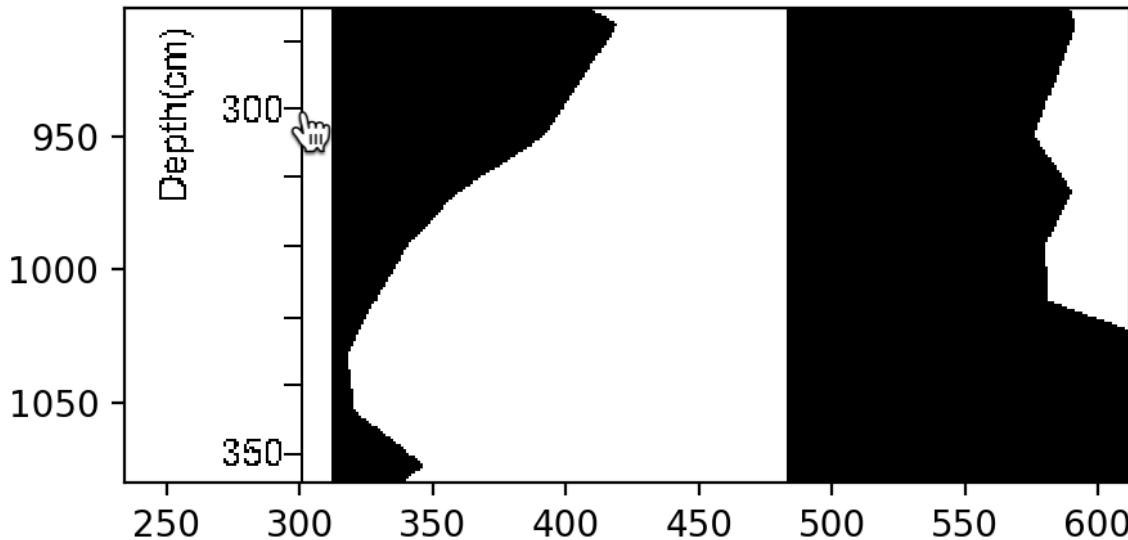
1. Expand the *Samples* item in the digitization control
2. Find the samples by clicking the *Find samples* button. straditize now identified the sample locations based on the extrema in the columns (see [Automatic samples identification](#) for an explanation of the algorithms).
3. To visualize the samples, you can again use the items in the *Plot control* tab. However, you can (and should!) also edit them by clicking the *Edit samples* button.
4. Now you see one horizontal line per sample that you can drag around (left-click), delete (right-click) or you can also add new samples (Shift + left-click). See the [Editing samples](#) section for more details.
5. Finally, click the *Apply* button or the *Cancel* button to stop the editing of the samples.

You're almost done! Your diagram is now digitized and the data could already be exported.

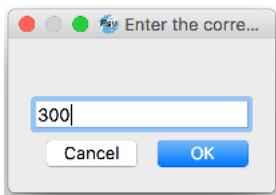
Translating the y-axis

To correctly reproduce the diagram data, there are now only two more things to know, that is the scaling of the y- and x-axes in the diagram.

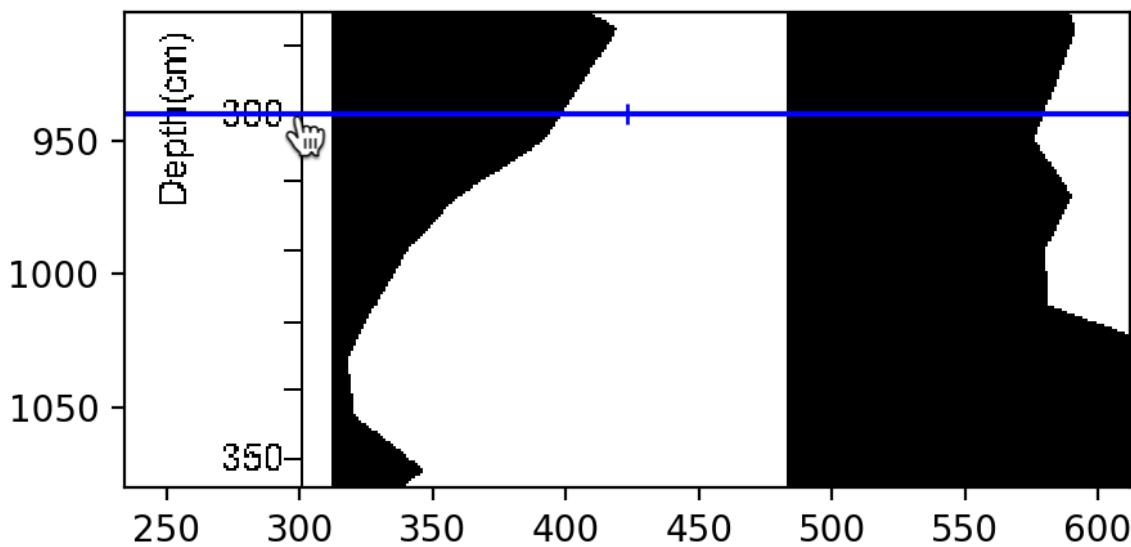
1. Expand the *Axes translations* tab in the digitization control
2. Click the *Insert Y-axis values* button in the *Axes translations* section of the straditzer control
3. Shift-leftclick on the plot to enter the corresponding y-value.



4. A small dialog will appear where you should enter the y-value to use (in this case, 300)

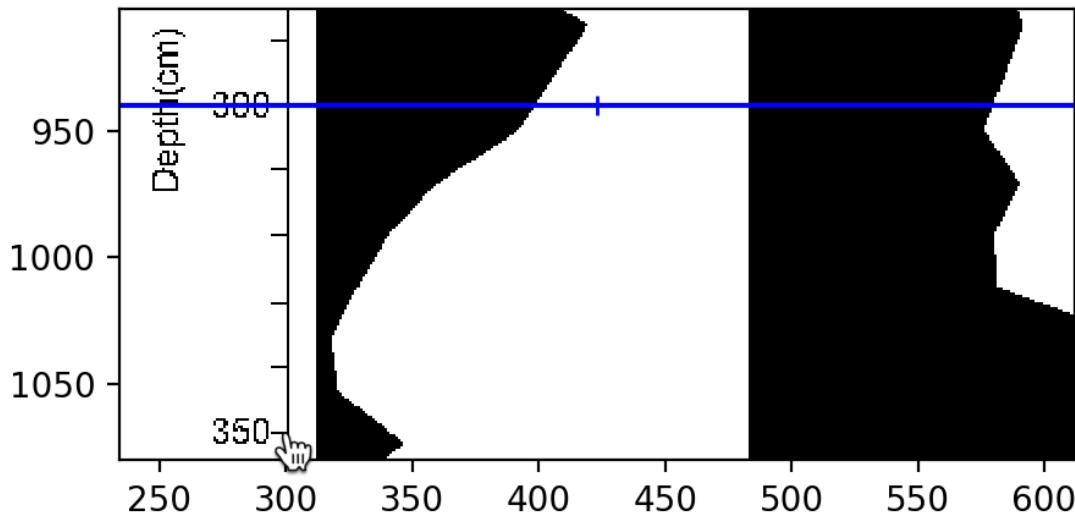


5. After hitting the *Ok* button, you will see a mark on the plot (blue line). You can select the mark via leftclick and drag it to a different location or you can delete it via rightclick.

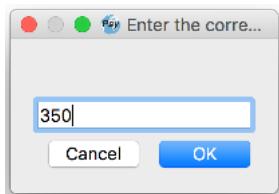


6. now repeat steps 2-4 on a second point on the y-axis

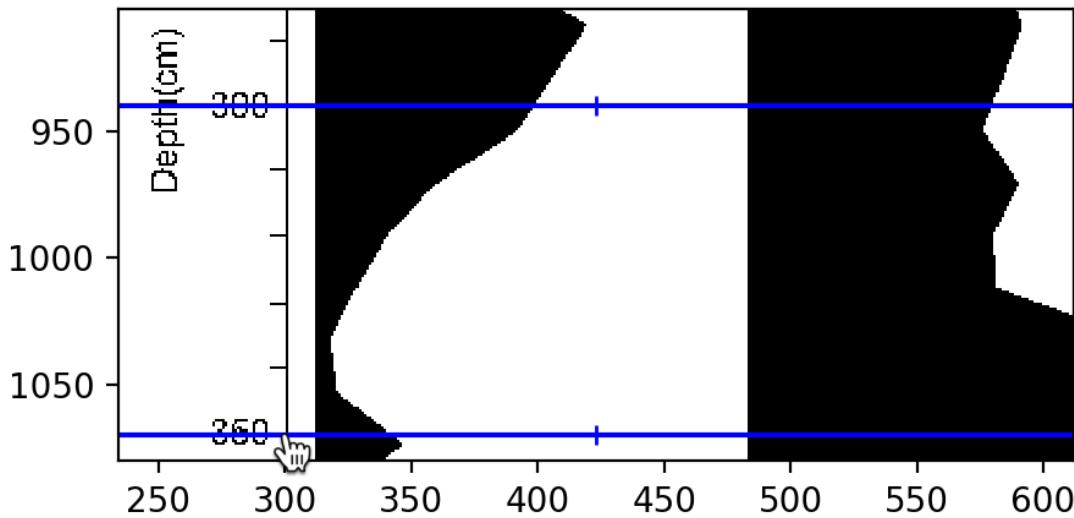
- Select another point



- Enter the corresponding value (here 350)



- A new mark is created that you can modify



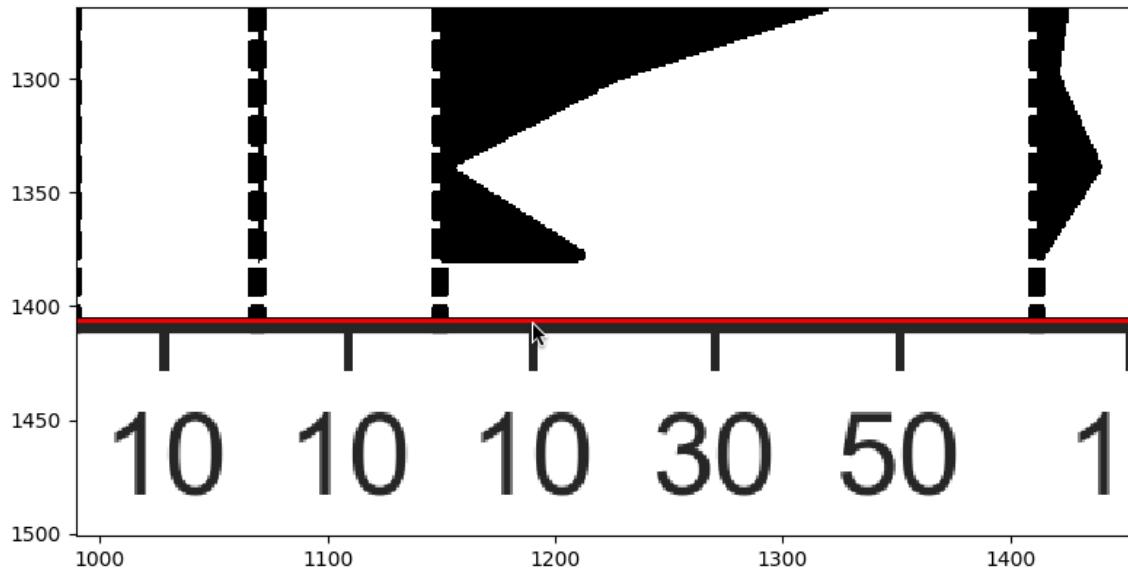
7. Click the *Apply* button at the bottom of the straditizer control when you are done.

Note: If you drag a mark and hold the Shift button while releasing the mouse button, the dialog in point 3 from above will not pop up.

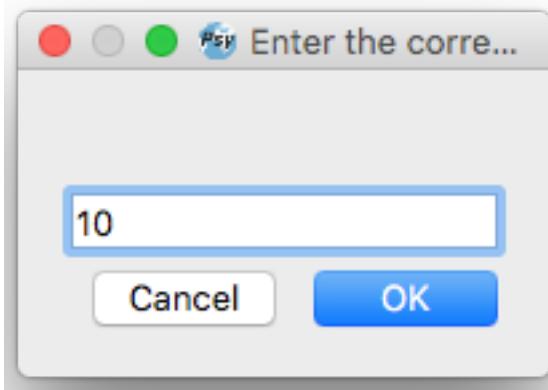
Translating the x-axis

As we did for the y-axes, we do it for the x-axes.

1. Click the *Insert X-axis values* button in the *Axes translations* section of the straditizer control
2. A small dialog will appear where you should enter the value at the column start (in our case, 0)
3. Shift-leftclick on the plot to enter another x-value.



4. A small dialog will appear where you should enter the x-value to use



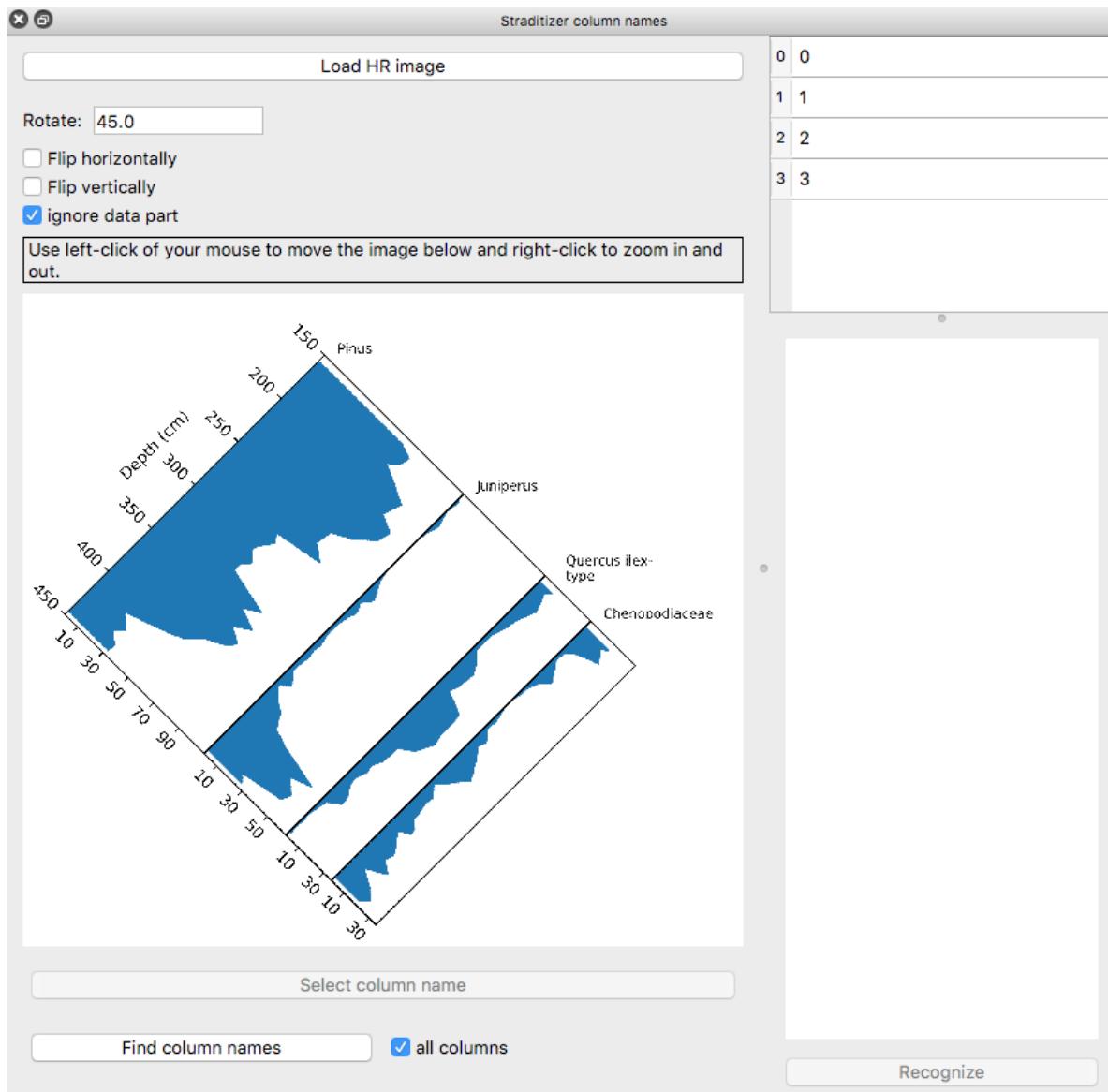
5. Click the *Apply* button at the bottom of the straditizer control when you are done.

Column names

Finally, we select the column titles and save them in the project.

Each of the sub diagrams has a title, in our case, *Pinus*, *Juniperus*, *Quercus ilex-type* and *Chenopodiaceae*. These names should be used in the final digitization result and therefore you can include them in straditize:

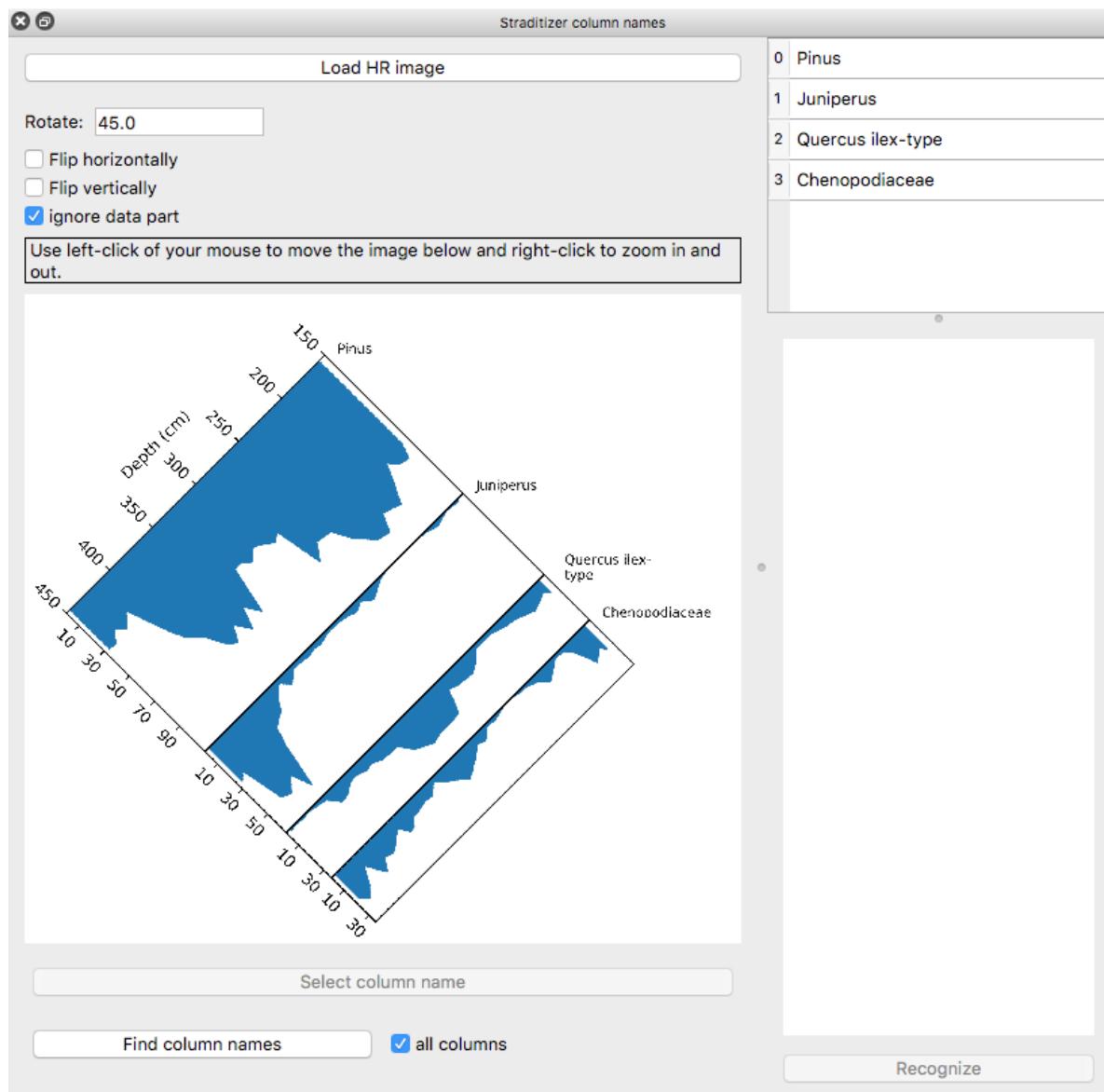
1. Expand the *Column names* item in the straditizer control and click the *Edit column names* button
2. In the appearing widget, the colnames editor



you find a table where you can edit the column names. The plot on its left also shows a rotated version of the diagram, to help you identifying the column names. You can navigate in this plot using leftclick and zoom in and out using right-click (see [matplotlibs docs on interactive navigation with the Pan/Zoom-tool](#)).

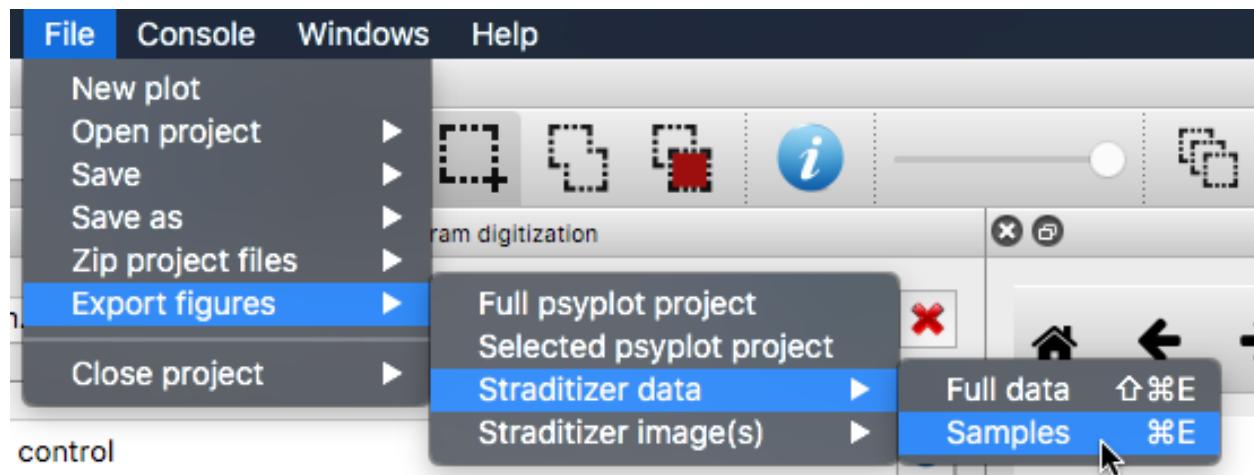
Hint: If you have tesseract installed (see the [user manual](#)), you can also just click the *Find column names* button and you are done.

-
3. When you entered the correct names in the table (see image below), click the *Edit column names* button again to hide the dialog.



Done!

That's it! You can export the samples (see [Export your results](#)) to an Excel or CSV file via
File → Export figures → Straditzer data → Samples



Now start with your own diagram!

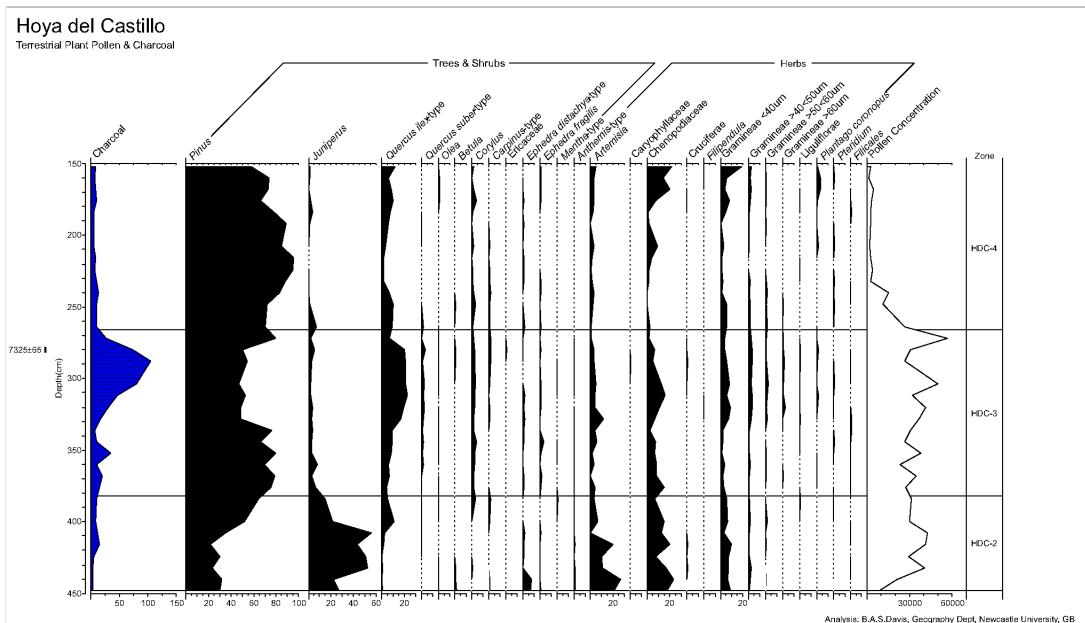
If you have questions or troubles, please open an issue on

<https://github.com/Chilipp/straditize/issues>

Thanks for using **straditize** and happy digitizing!

References

Advanced tutorial: The Hoya del Castillo pollen diagram



In this tutorial, we digitize a stratigraphic diagram based on the data in [Basil2007] and introduce to you how to extract the samples out of an area diagram.

At the bottom of the straditizer control, you find a navigation panel which helps you to navigate through the tutorial. Click the *Next* button in this control to start the tutorial. You can skip steps by clicking the *Skip* button and you can check what you have done using the *Check* button and then proceed with the next step.

The tutorial will take between 15 and 30 minutes. You can save your current state to a file using *File → Save → Save straditizer* (see *Saving and loading your project*) and reload it later via *File → Open project → Open straditizer → Project or image*

If you have questions or troubles with this tutorial, please open an issue on

<https://github.com/Chilipp/straditize/issues>

and we will do our best to assist you.

Load the image of the diagram

The first step is to load the image of the diagram into straditize.

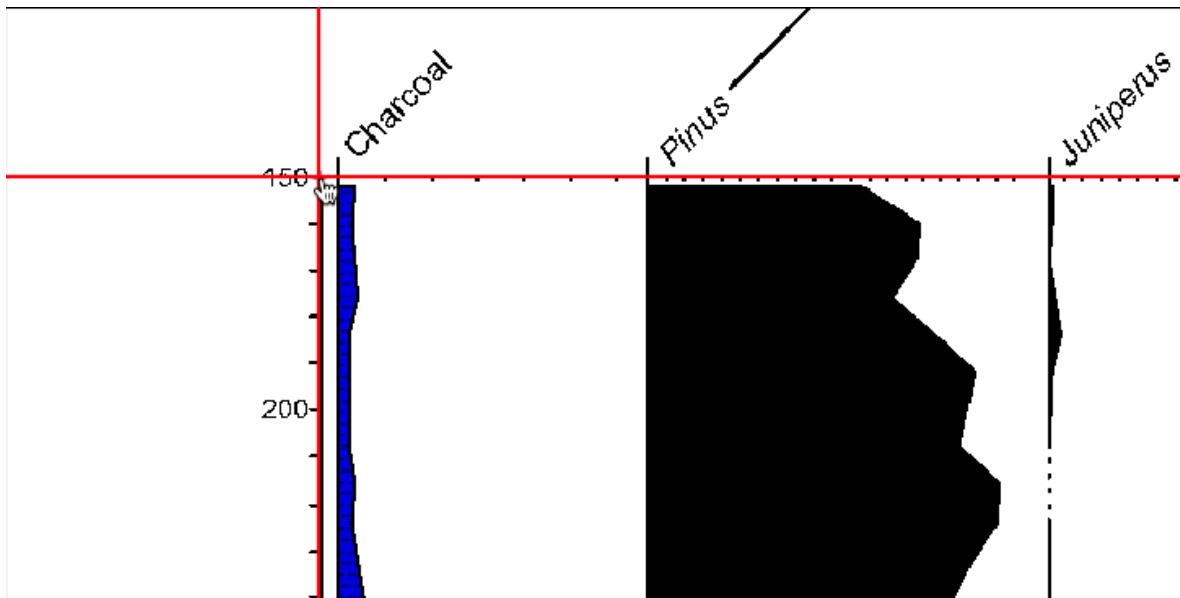
1. Click the button at the top of the straditizer control.
2. Select the image file `hoya-del-castillo.png` from your hard-disk and click *Open*

Select the diagram part

The second step involves selecting the diagram part. This is the part where your data is displayed, but without the vertical and horizontal axes or axes descriptions.

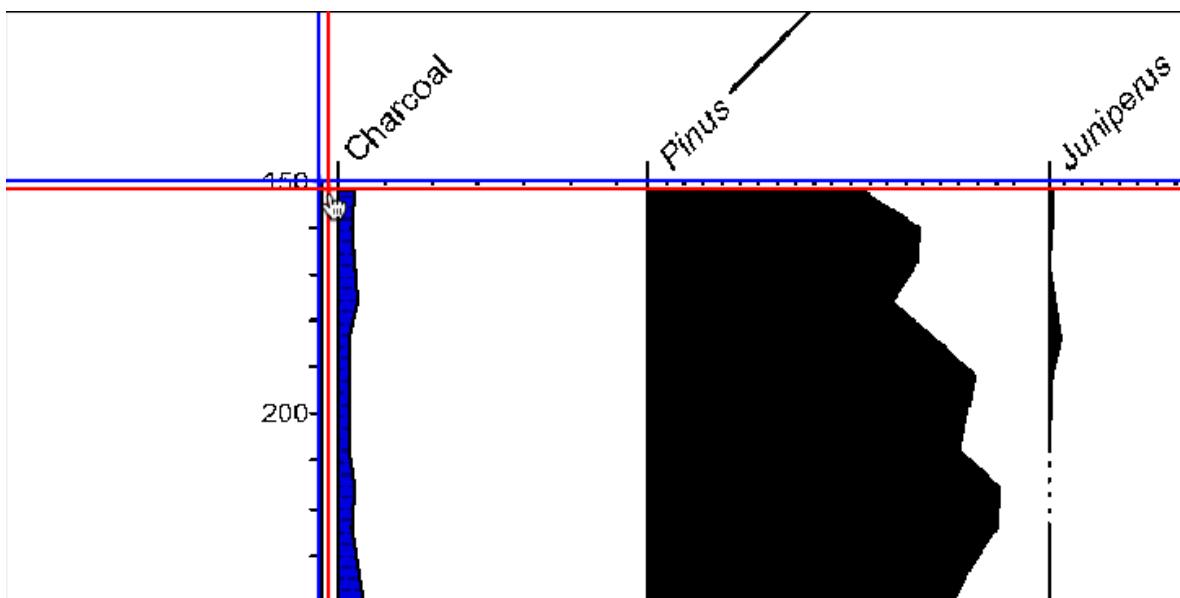
Being exact in this step significantly reduces the later work and simplifies the automatic digitization.

1. Click the *Select data part* button in the digitization control.
2. Click on the upper left cross to select it

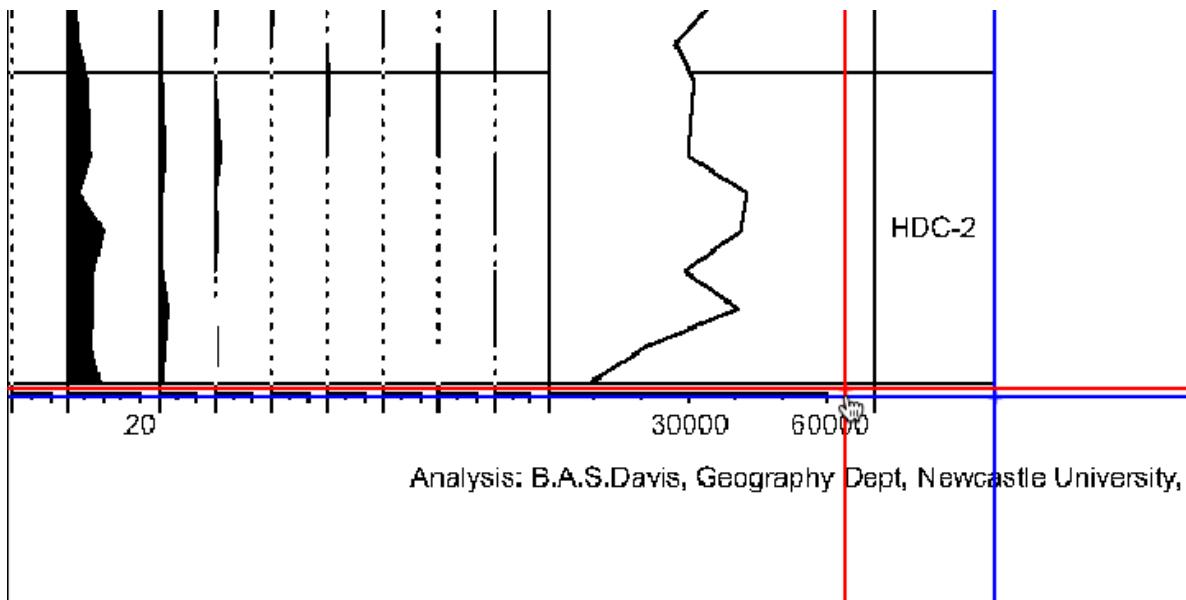


3. Hold the left mouse button and drag the cross to the upper left corner of the diagram. Make sure you don't include the y-axis and x-axis ticks.

You can also use the zoom and navigation tools in the figure toolbar for navigation.



5. Now select the cross at the lower right corner of the diagram and move it such that you don't include the x-axis ticks (i.e. the numbers on the x-axis)



6. Click the *Apply* button at the bottom

If you want to change the appearance of the marks, see the *Marker control* section in the straditizer control panel.

Create the diagram reader

Now you have to select, which type of diagram you are digitizing. In the example, it is an area diagram which is already selected in the dropdown menu next to the *Select data part* button.

Click on *Convert image* to start the digitization.

See also:

[Selecting the reader](#)

Identifying the columns

The next step is, to automatically separate the columns from each other. This is simple in our case, since straditize can separate them automatically.

1. Click the *Column starts* button. You will see several vertical blue lines appearing on the plot that you can drag and drop such as you did it when selecting the diagram part. You can change their colors using the [marker control](#)
2. straditize recognizes these columns automatically and in our example, you do not have to edit them. Therefore hit the *Apply* button at the bottom of the control and you are done.

See also:

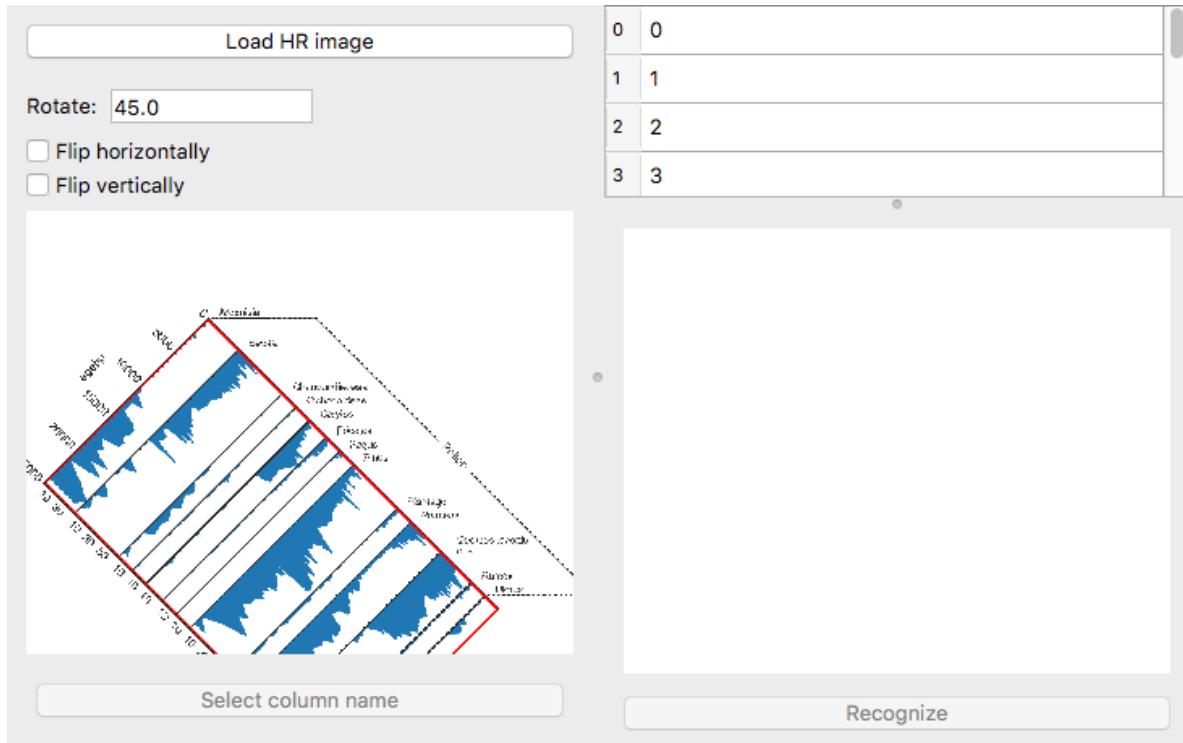
[Select the starts for each column](#)

Specifying the column names

Straditze can handle column names which will then be included in the final export. It interfaces with the `tesseract` package and we will use this to minimize our typing amount. However, we have a complex diagram so manual corrections are unavoidable.

1. Expand the *Column names* item in the straditzer control and click the *Edit column names* button.

In the appearing widget, the colnames editor



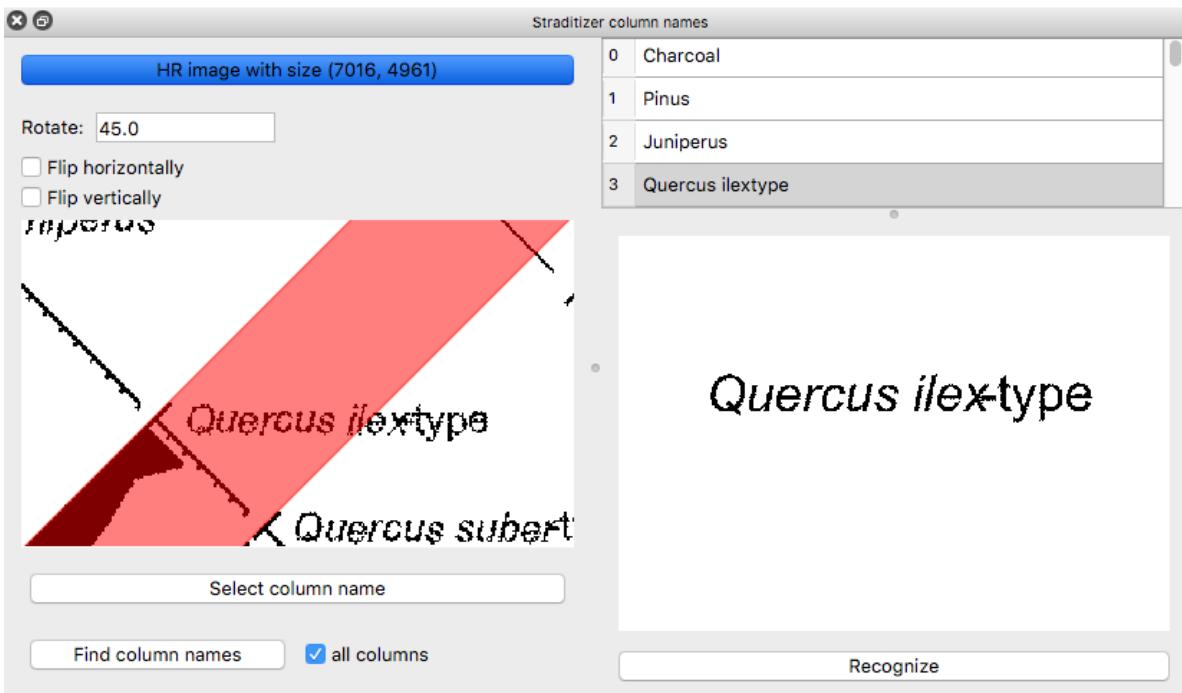
you find a table where you can edit the column names. The plot on it's left also shows a rotated version of the diagram, to help you identifying the column names. You can navigate in this plot using leftclick and zoom in and out using right-click (see [matplotlibs docs on interactive navigation with the Pan/Zoom-tool](#))

2. To improve the text recognition, it is highly recommended to have a clean image with only the column names on it and a sufficient resolution. We have something prepared for you:

The screenshot shows a software interface with a list of column names. The names are arranged in two rows. The first row includes: Charcoal, Pinus, Juniperus, Quercus flex-type, Quercus sub-type, Olea, Betula, Corilus, Ericaceous-type, Ericaea effatae-type, Mentheta rapica, Arthrostipa, Carex, Cyperaceae, Chionopodiaceae, Cruciferae, Filipendula, Gramineas <40um, Gramineas 40-50um, Gramineas >50-60um, Liquidambar, Plantago coronopus, Phragmites, Filaria, and Pollen Concentration. The second row includes: Pollen.

Click the *Load HR image* button and select the *hoya-del-castillo-colnames.png* image file.

3. We will now use the automatic finding of column names. For this, click the *Find column names* button. This will search for the column names in the plot to the left of the table
4. There will probably be some errors in the column names. Therefore, go through each row in the table and check the name. You can use the *Hint* button to help you and/or zoom to the column name to see the name in the original



- When you entered the correct names in the table, click the *Edit column names* button again to hide the button.

Remove artifacts

Stratigraphic diagrams, and especially pollen diagrams, often have a lot of artifacts in it, that are informative for the reader.

In our case, the diagram is splitted into three temporal *zones* (HDC-2, HDC-3 and HDC-4, see the *Zone* column on the right part of the diagram) which are visually separated with horizontal lines. Additionally, the diagram has vertical, dashed lines at each column start (the y-axes for each column).

Before we digitize our diagram, those *informative features* have to be removed. You can do this in an external image editing software (e.g. Photoshop) but we also implemented several automated algorithms to detect common features and remove them easily.

For our tutorial, we use the *Remove lines* feature to detect and remove the vertical and horizontal lines.

Horizontal lines

- Expand the *Remove features* tab in the *Digitization control*
- Expand the item with the *vertical lines* and *horizontal lines* button by clicking on the small arrow on their left
- Set the minimum line width to 1 pixel
- Click the *horizontal lines* button. In the plot you will see, that the horizontal lines are red now (if necessary, go with the mouse over the plot and you will see it in the zoom window). You could edit the selection now using the selection toolbar (see [The selection toolbar](#)), but for our tutorial, this is not necessary.
- Click the *Remove* button to remove the lines

Vertical lines (y-axes)

1. Enable the maximum line width and set it to 2 pixel
2. Set the minimum fraction to 30%
3. Click the *vertical lines* button and the vertical lines turn red and are marked to be removed.
4. Click the *Remove* button to remove the y-axes

Additional automated removal tools are available and fully described in the documentation (see [Removing features](#)). But here, we can continue with the digitization.

Digitizing the data

The next step, after we cleaned up the image, is the digitization of the diagram. Click the *Digitize* button.

You now see the lines that result from the digitization. You can remove them in the *Plot control* section by clicking the  button of the *Full digitized data* row.

Finding and editing samples

So far, we have one data point per pixel in the image. However, we have to identify the locations of the samples in order to reproduce the original data.

Straditize assists you with this through automatic sample finding algorithms, or you can load the sample locations from an external files or you just add and edit the samples manually.

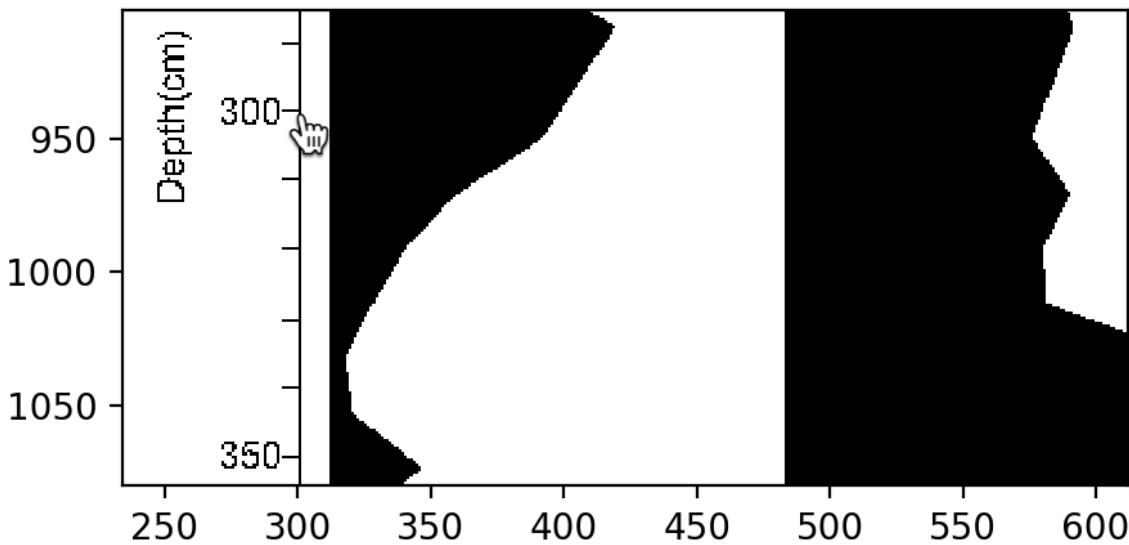
1. Expand the *Samples* item in the digitization control
2. Find the samples by clicking the *Find samples* button. straditize now identified the sample locations based on the extrema in the columns (see [Automatic samples identification](#) for an explanation of the algorithms).
3. To visualize the samples, you can again use the items in the *Plot control* tab. However, you can (and should!) also edit them by clicking the *Edit samples* button.
4. Now you see one horizontal line per sample that you can drag around (left-click), delete (right-click) or you can also add new samples (Shift + left-click). See the [Editing samples](#) section for more details.
5. Finally, click the *Apply* button or the *Cancel* button to stop the editing of the samples.

You're almost done! Your diagram is now digitized and the data could already be exported.

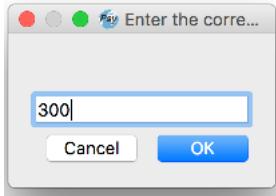
Translating the y-axis

To correctly reproduce the diagram data, there are now only two more things to know, that is the scaling of the y- and x-axes in the diagram.

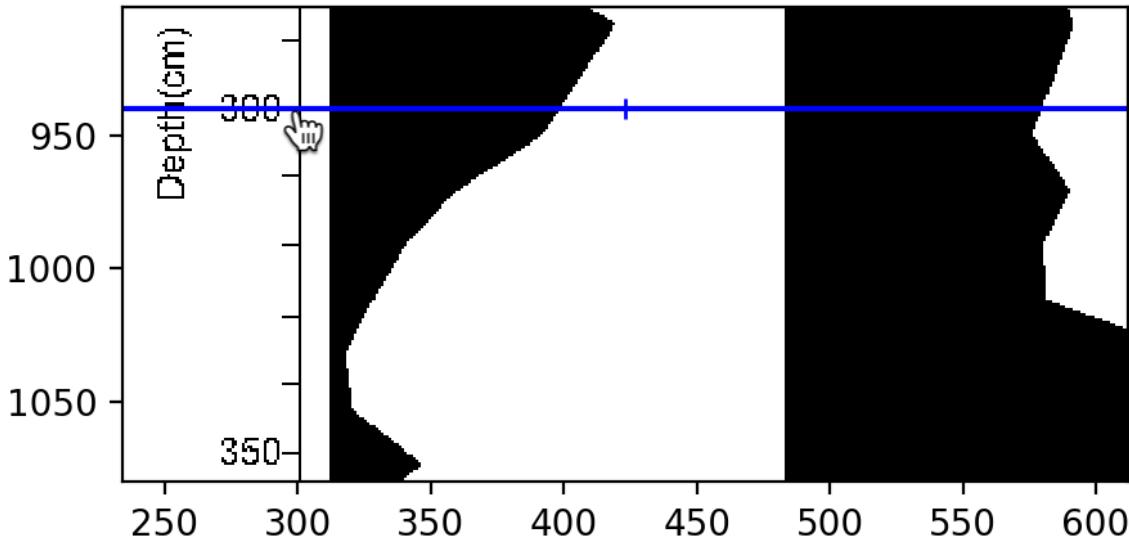
1. Expand the *Axes translations* tab in the digitization control
2. Click the *Insert Y-axis values* button in the *Axes translations* section of the straditzer control
3. Shift-leftclick on the plot to enter the corresponding y-value.



4. A small dialog will appear where you should enter the y-value to use (in this case, 300)

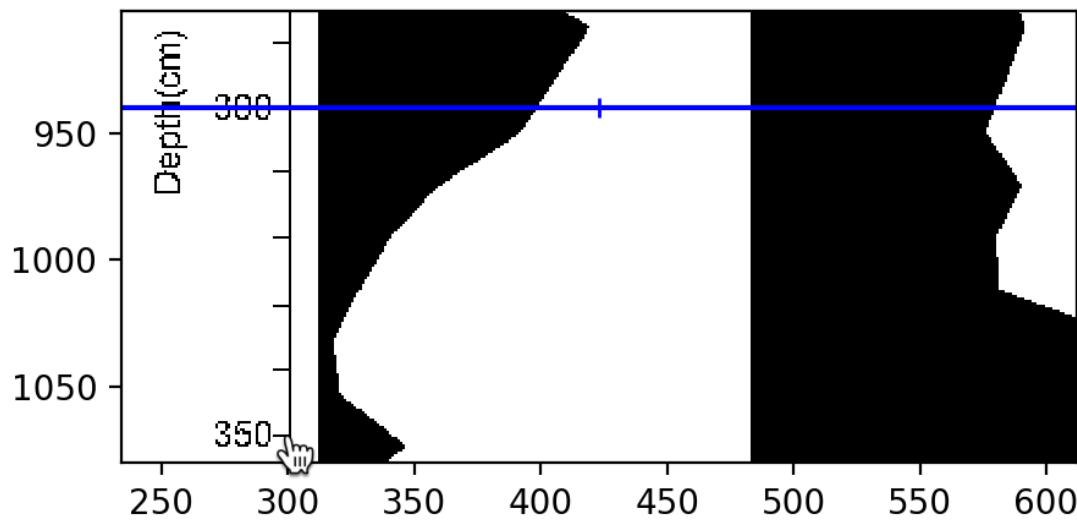


5. After hitting the *Ok* button, you will see a mark on the plot (blue line). You can select the mark via leftclick and drag it to a different location or you can delete it via rightclick.

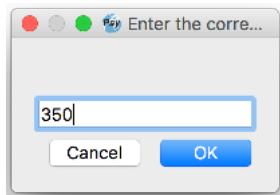


6. now repeat steps 2-4 on a second point on the y-axis

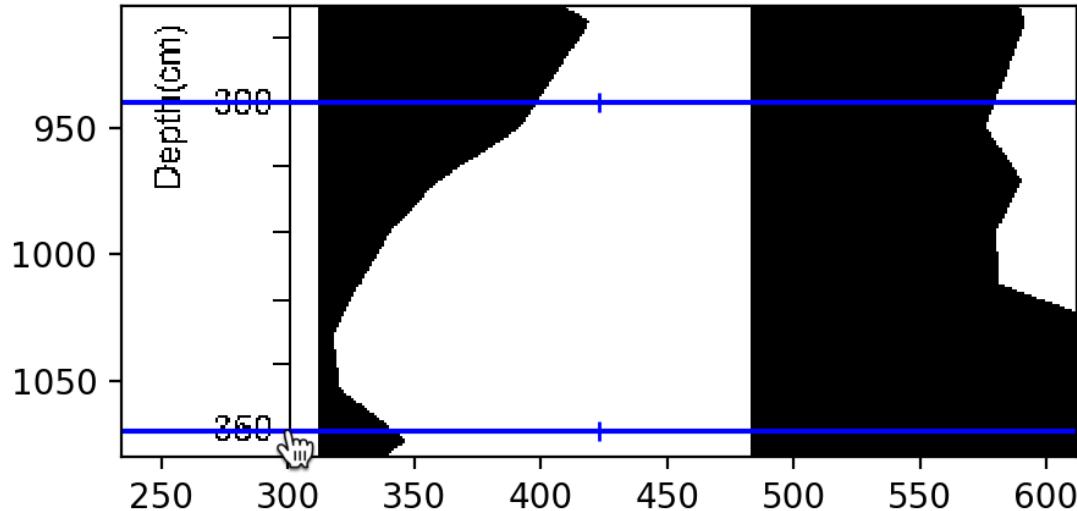
- Select another point



- Enter the corresponding value (here 350)



- A new mark is created that you can modify



7. Click the *Apply* button at the bottom of the straditzer control when you are done.

Note: If you drag a mark and hold the Shift button while releasing the mouse button, the dialog in point 3 from above will not pop up.

Translating the x-axes

Additionally to the vertical axis, we have to tell straditize how to interpret the x-axes of the stratigraphic diagram.

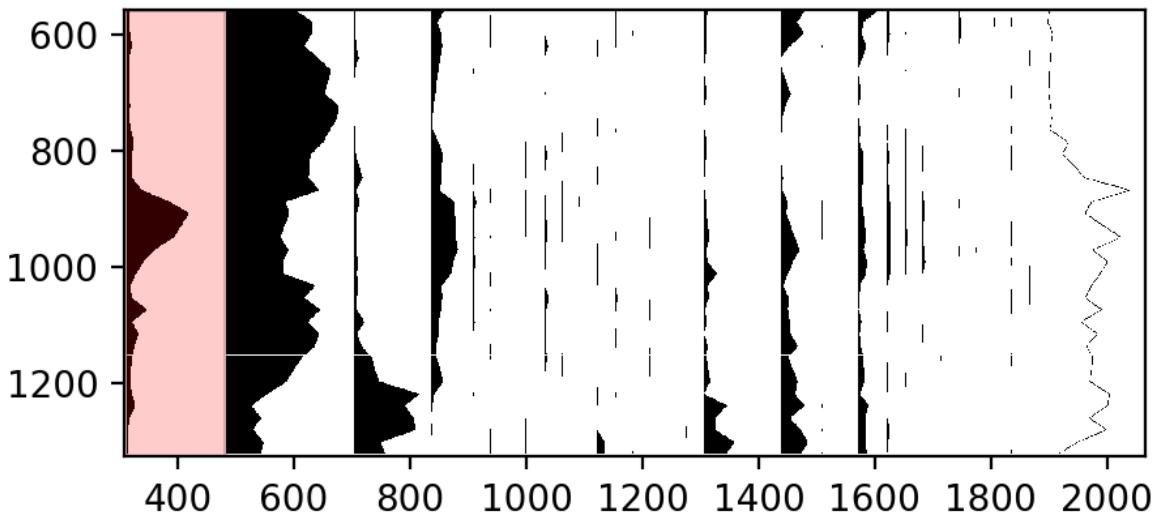
Here we have to deal different units of the x-axes in the diagrams. The first (*Charcoal*) and last (*Pollen Concentration*) columns are counts, the pollen taxa in between are in percent.

Therefore we will create column specific readers, one for the first and one for the last column. Then we will insert the x-axis data.

The steps down below are repetitive, but we will describe them for each step in detail.

The *Charcoal* column

1. Expand the *Current reader* tab in the digitization control
2. Click the + button to start the selection for a new reader
3. Click in the plot on the first column. It will turn red



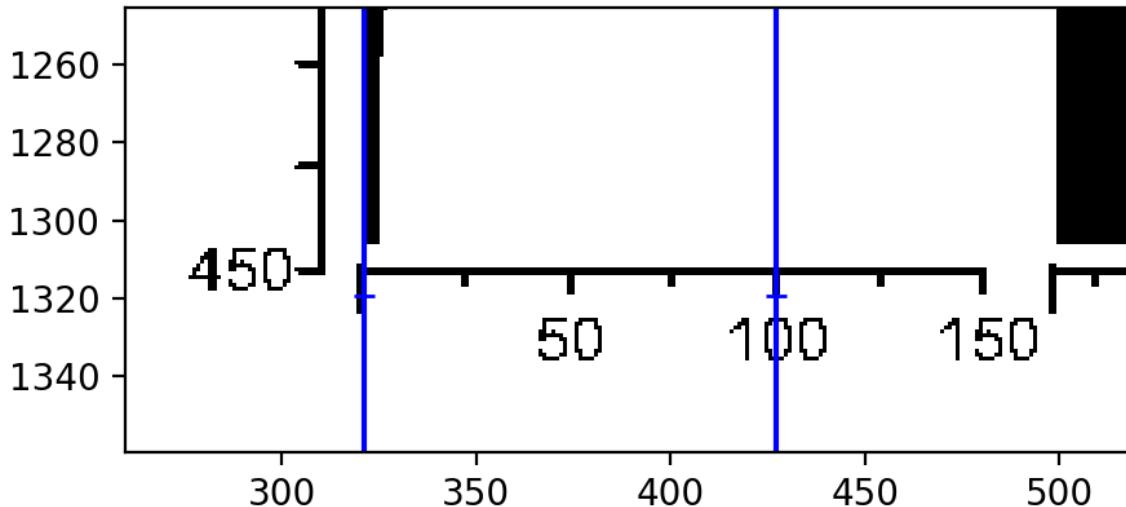
4. Click the *Apply* button and select the *area* reader in the appearing dialog
5. In the dropdown menu in the *Current reader* tab, select the reader for *Column 0* to make it the current reader.

Now, we select the values for interpreting the x-axis of this column. The procedure is more or less the same, as before with the y-axis:

1. Click the *Insert X-axis values* button in the *Axes translations* section of the straditzer control
2. A small dialog will appear where you should enter the x-value of the start of the column (here 0).

Note: The dialog only appears, if the start of one of the selected columns is visible. If this is not the case, Shift-Leftclick on the plot to create a mark.

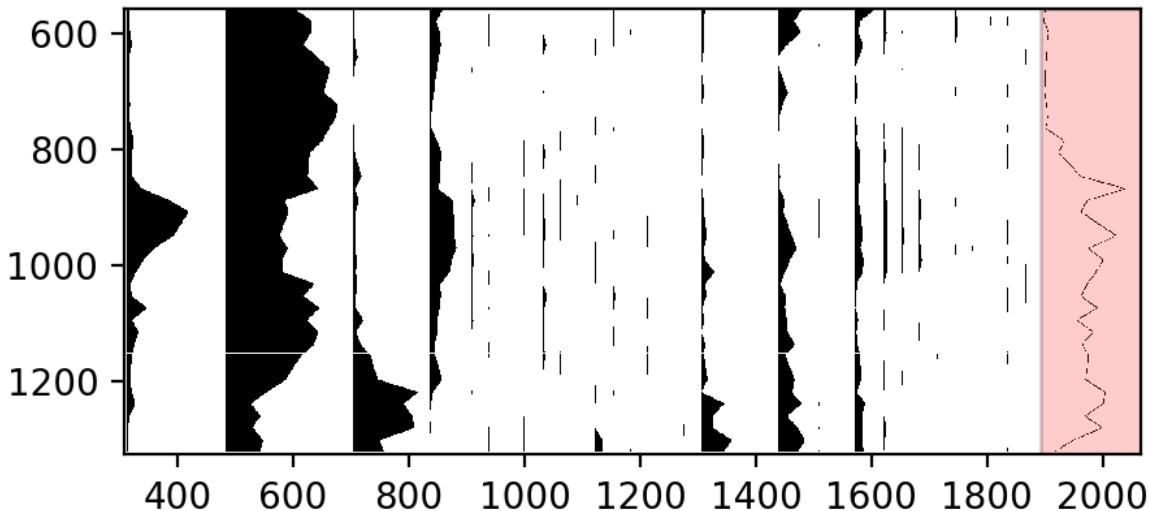
3. After hitting the *Ok* button, you will see a mark on the plot (blue line). You can select the mark via leftclick and drag it to a different location or you can delete it via rightclick.
4. now repeat steps 2-4 on a second point in the same column. Your diagram should now look something like this:



5. Click the *Apply* button at the bottom of the straditzer control and we can continue with the last column

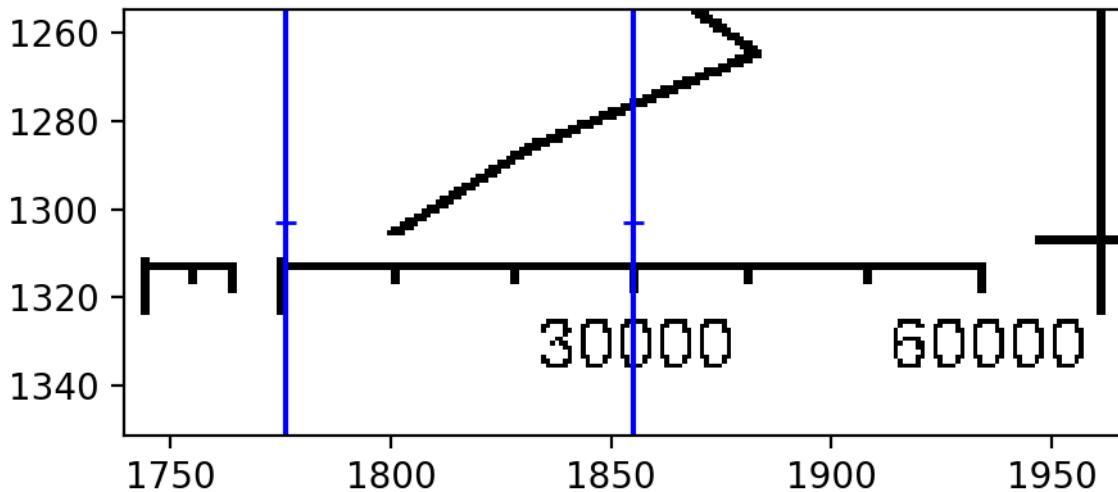
The *Pollen Concentration* column

1. In the *Current reader* dropdown menu, select the reader for the columns 1-27
2. Click the + button to start the selection for a new reader
3. Click in the plot on the last column. It will turn red



4. Click the *Apply* button and select the *line* reader in the appearing dialog.
 5. In the dropdown menu in the *Current reader* tab, select the reader for *Column 27* to make it the current reader.
- Now, we select the values for interpreting the x-axis of this column. The procedure is the same as above:

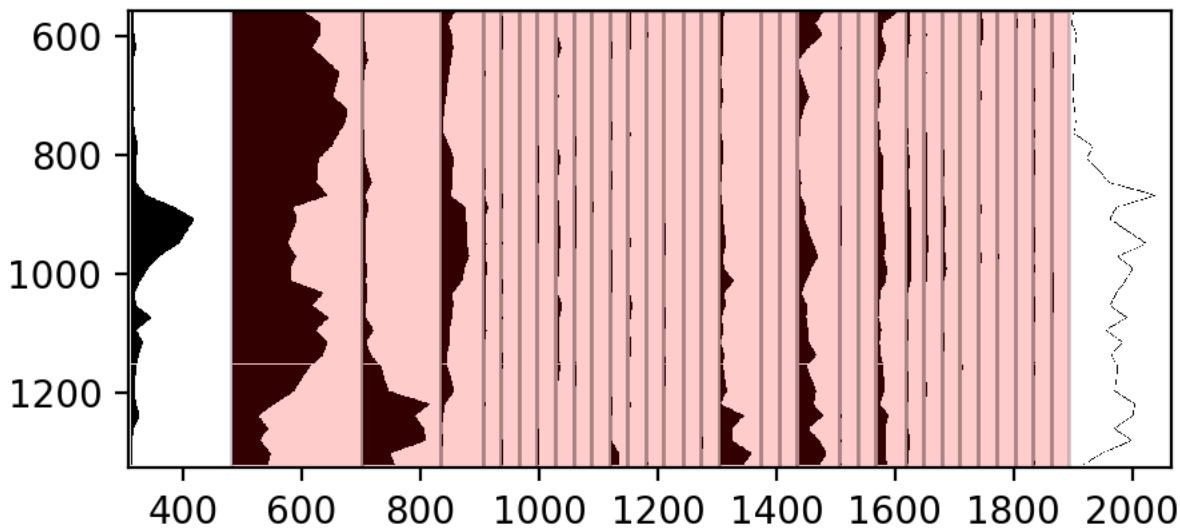
1. Click the *Insert X-axis values* button in the *Axes translations* section of the straditizer control
2. In the appearing dialog, enter the x-value of the column start (here again 0) or Shift-Leftclick on the plot and then enter it (see note above).
3. After hitting the *Ok* button, you will see a mark on the plot (blue line). Again, you can select the mark, drag it or delete it as before.
4. now repeat steps 2-4 on a second point in the same column. Your diagram should now look something like this:



5. Click the *Apply* button at the bottom of the straditizer control and we can continue with the pollen taxa.

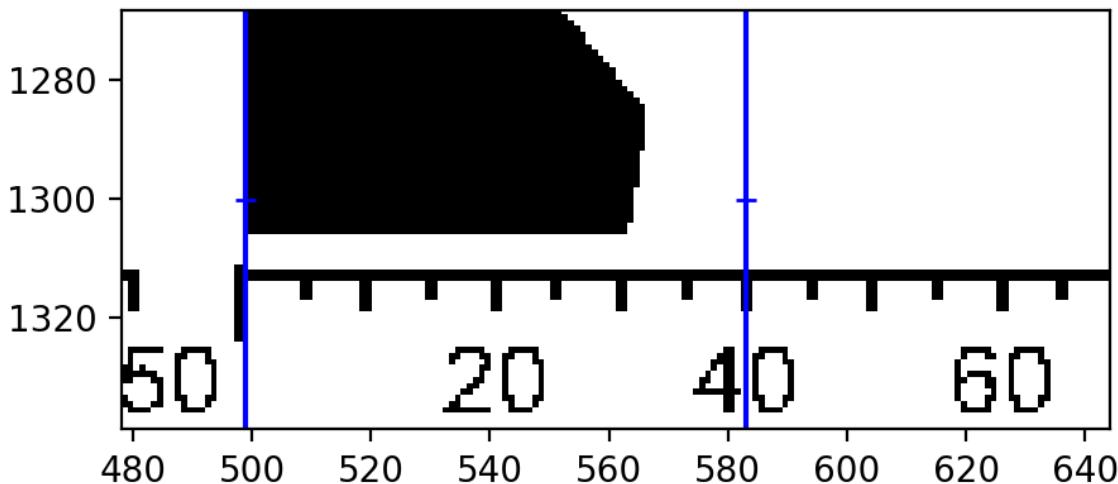
The pollen taxa columns

Last but not least, we translate the x-axes informations for the pollen taxa.



Luckily, as it is common for pollen diagrams, they all have the same scaling. Therefore it is enough to perform the above steps just for one of the columns.

1. In the dropdown menu in the *Current reader* tab, select the reader for *Columns 1-26* to make it the current reader.
2. Click the *Insert X-axis values* button in the *Axes translations* section of the straditzer control
3. In the appearing dialog, enter the x-value of the column start (here again 0) or Shift-Leftclick on the plot and then enter it (see note above).
4. After hitting the *Ok* button, you will see a mark on the plot (blue line). Again, you can select the mark, drag it or delete it as before.
5. now repeat steps 2-4 on a second point in the same column. Your diagram should now look something like this:



6. Click the *Apply* button at the bottom of the straditzer control.

Editing the meta attributes

To keep the overview on your project, you can save some meta informations to it, for example, where you have the diagram from and what it represents.

Here for example, we know that the digitized data represents a pollen core from *Hoya del Castillo* and has been described in detail in [Basil1980].

1. Click the *Attributes* button at the bottom of the straditzer control. You know see a table with attributes, that you can rename, delete and fill according to your needs.
2. Fill the data table:

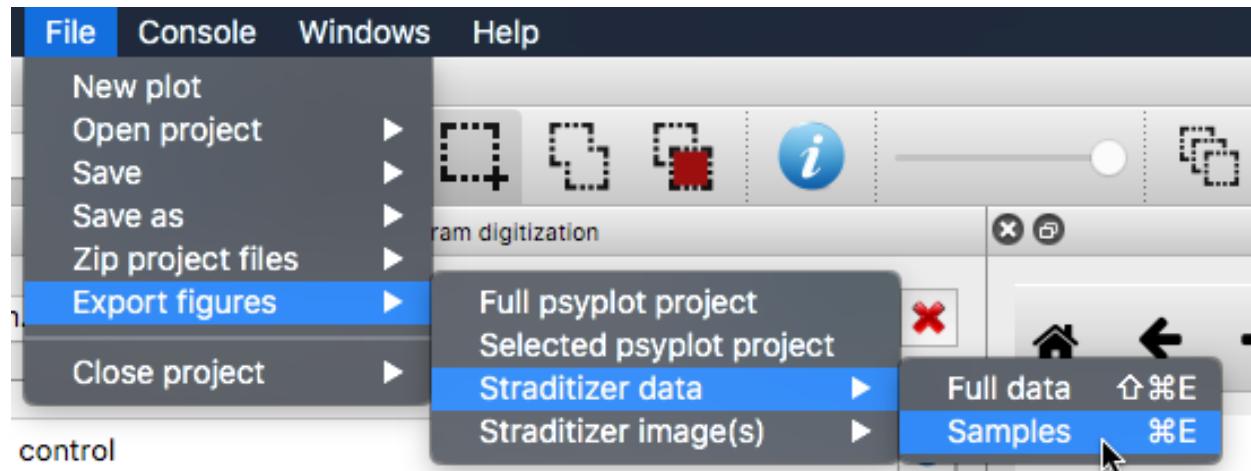
Digi-tized by	Your name
site-name	Hoya del Castillo
Lon	-0.5
Lat	41.25
Archive	Pollen
Country	Spain
Restricted	No
Reference	Davis, Basil A. S., and A. C. Stevenson. "The 8.2ka Event and Early-Mid Holocene Forests, Fires and Flooding in the Central Ebro Desert, NE Spain." Quat. Sci. Rev. , vol. 26, no. 13-14, 2007, pp. 1695-712
DOI	10.1016/j.quascirev.2007.04.007

Note: You can also add new attributes by clicking *Rightclick* → *Insert 1 row* inside the table

Done!

That's it! You can export the samples (see *Export your results*) to an Excel or CSV file via

File → *Export figures* → *Straditizer data* → *Samples*



Now start with your own diagram!

If you have questions or troubles, please open an issue on

<https://github.com/Chilipp/straditize/issues>

Thanks for using **straditize** and happy digitizing!

References

1.4 Contributing, asking for assistance and reporting bugs

First off, thanks for taking the time to contribute!

The following set of guidelines for contributing to straditiz are mostly guidelines, not rules. Use your best judgment, and feel free to propose changes to this document in a pull request.

Table of Contents

- *Contributing, asking for assistance and reporting bugs*
 - *Code of Conduct*
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 - * *Suggesting Enhancements*
 - *How Do I Submit A (Good) Enhancement Suggestion?*
 - * *Pull Requests*
 - * *Adding new examples*
 - *Styleguides*
 - * *Git Commit Messages*
 - * *Documentation Styleguide*
 - *Example*

1.4.1 Code of Conduct

This project and everyone participating in it is governed by the straditiz Code of Conduct. By participating, you are expected to uphold this code.

1.4.2 How Can I Contribute?

Reporting Bugs

This section guides you through submitting a bug report for straditiz. Following these guidelines helps maintainers and the community understand your report, reproduce the behavior, and find related reports.

Before creating bug reports, please check existing issues and pull requests as you might find out that you don't need to create one. When you are creating a bug report, please *include as many details as possible*. Fill out the required template, the information it asks for helps us resolve issues faster.

Note: If you find a **Closed** issue that seems like it is the same thing that you're experiencing, open a new issue and include a link to the original issue in the body of your new one.

How Do I Submit A (Good) Bug Report?

Bugs are tracked as GitHub issues. Create an issue on straditiz repository and provide the following information by filling in [the template](#).

Explain the problem and include additional details to help maintainers reproduce the problem:

- **Use a clear and descriptive title** for the issue to identify the problem.
- **Describe the exact steps which reproduce the problem** in as many details as possible.
- **Provide specific examples to demonstrate the steps.** Include links to files or images, or copy/pasteable snippets, which you use in those examples. If you're providing snippets in the issue, use [Markdown code blocks](#).
- **Upload the project file.** The best is, you only take the part of the stratigraphic diagram that causes the problems.
- **Describe the behavior you observed after following the steps** and point out what exactly is the problem with that behavior.
- **Explain which behavior you expected to see instead and why.**
- **Include screenshots and animated GIFs** which show you following the described steps and clearly demonstrate the problem. You can use [this tool](#) to record GIFs on macOS and Windows, and [this tool](#) or [this tool](#) on Linux.
- **If the problem is related to your data structure,** include a small example how a similar data structure can be generated

Include details about your configuration and environment:

- **Which version of straditiz and psyplot are you using?** You can get the exact version by running straditiz -V and psyplot -aV in your terminal , or by starting the psyplot-gui and open Help->Dependencies.
- **What's the name and version of the OS you're using?**
- **Use conda's diagnostics!** If you installed straditiz through anaconda, include the output of conda info -a and conda list in your description

Suggesting Enhancements

If you want to change an existing feature, use the [change feature template](#), otherwise fill in the [new feature template](#).

How Do I Submit A (Good) Enhancement Suggestion?

Enhancement suggestions are tracked as GitHub issues. Create an issue in the straditiz repository and follow these steps:

- **Use a clear and descriptive title** for the issue to identify the suggestion.
- **Provide a step-by-step description of the suggested enhancement** in as many details as possible.
- **Provide specific examples to demonstrate the steps.** Include copy/pasteable snippets which you use in those examples, as [Markdown code blocks](#).
- **Describe the current behavior and explain which behavior you expected to see instead** and why.
- **Include screenshots and animated GIFs** which help you demonstrate the steps or point out the part of psyplot which the suggestion is related to. You can use [this tool](#) to record GIFs on macOS and Windows, and [this tool](#) or [this tool](#) on Linux.

- Explain why this enhancement would be useful to most straditize users.
- List some other analysis software or applications where this enhancement exists.
- Which version of straditize and psyplot are you using? You can get the exact version by running straditize -V and psyplot -aV in your terminal , or by starting the psyplot-gui and open Help->Dependencies.
- Specify the name and version of the OS you're using.

Pull Requests

- Fill in the required template
- Do not include issue numbers in the PR title
- Include screenshots and animated GIFs in your pull request whenever possible.
- Document new code based on the *Documentation Styleguide*
- End all files with a newline and follow the PEP8, e.g. by using flake8

Adding new examples

You have new examples? Great! straditize can only be improved through new tutorials. You can either implement it in the graphical user interface (see the `straditize.widgets.tutorial` module) or you just add them to the sphinx documentation. We are also looking forward to assist you in the implementation and the sharing of your experiences.

1.4.3 Styleguides

Git Commit Messages

- Use the present tense (“Add feature” not “Added feature”)
- Use the imperative mood (“Move cursor to...” not “Moves cursor to...”)
- Limit the first line (summary) to 72 characters or less
- Reference issues and pull requests liberally after the first line
- When only changing documentation, include [ci skip] in the commit title

Documentation Styleguide

- Follow the numpy documentation guidelines.
- Use reStructuredText.
- Try to not repeat yourself and make use of the `straditize.common.docstrings`

Example

```

@docstrings.get_sections('new_function')
def new_function(a=1):
    """Make some cool new feature

    This function implements a cool new feature

    Parameters
    -----
    a: int
        First parameter

    Returns
    -----
    something awesome
        The result"""

    ...

@docstrings.dedent
def another_new_function(a=1, b=2):
    """Make another cool new feature

    Parameters
    -----
    %(new_function.parameters)s
    b: int
        Another parameter

    Returns
    -----
    Something even more awesome"""

    ...

**Note:** This document has been inspired by `the contribution
guidelines of
Atom <https://github.com/atom/atom/blob/master/CONTRIBUTING.md#git-commit-messages>`__

```

1.5 Command line usage

The `straditize.__main__` module defines a simple parser to parse commands from the command line to load a diagram or a straditize project.

It can be run from the command line via:

```
python -m straditize [options] [arguments]
```

or simply:

```
straditize [options] [arguments]
```

Load a dataset, make the plot and save the result to a file

```
usage: straditize [-h] [-b [backend]] [-ni] [-rc-gui RC_GUI_FILE]
                  [-inc str [str ...]] [-exc str [str ...]] [--offline]
                  [-pwd str] [-s str] [-c str]
                  [-opengl {'software', 'desktop', 'gles', 'automatic'}]
```

(continues on next page)

(continued from previous page)

```

[-o str] [-xlim val val] [-ylim val val] [-f]
[-rt { 'area' | 'bars' | 'rounded bars' | 'stacked area' | 'line' }]
[-V]
[str]

```

1.5.1 Positional Arguments

str	Either the path to a picture to digitize or a previously saved straditzer project (ending with '.pkl')
------------	--

1.5.2 Output options

Options that only have an effect if the `-o` option is set.

-o, --output	The path to the csv file where to save the digitized diagram
---------------------	--

1.5.3 Gui options

Options specific to the graphical user interface

-b, --backend	The backend to use. By default, the 'gui.backend' key in the <code>rcParams</code> dictionary is used. If used without options, the default matplotlib backend is used.
	Default: False
-ni, --new-instance	If True/set and the <code>output</code> parameter is not set, a new application is created
	Default: False
-rc-gui, --rc-gui-file	The path to a yaml configuration file that can be used to update the <code>rcParams</code>
-inc, --include-plugins	The plugin widget to include. Can be either None to load all that are not explicitly excluded by <code>exclude_plugins</code> or a list of plugins to include. List items can be either module names, plugin names or the module name and widget via ' <code><module_name>:<widget></code> '. Default: None
-exc, --exclude-plugins	The plugin widgets to exclude. Can be either 'all' to exclude all plugins or a list like in <code>include_plugins</code> . Default: []
	Default: []
--offline	If True/set, psyplot will be started in offline mode without intersphinx and remote access for the help explorer
	Default: False
-pwd	The path to the working directory to use. Note if you do not provide any <code>fnames</code> or <code>project</code> , but set the <code>pwd</code> , it will switch the <code>pwd</code> of the current GUI.
-s, --script	The path to a python script that shall be run in the GUI. If the GUI is already running, the commands will be executed in this GUI.
-c, --command	Python commands that shall be run in the GUI. If the GUI is already running, the commands will be executed in this GUI

-opengl, --opengl-implementation Possible choices: software, desktop, gles, automatic

OpenGL implementation to pass to Qt. Possible options are ‘software’, ‘desktop’, ‘gles’ and ‘automatic’ (which let’s PyQt decide).

1.5.4 Straditizer options

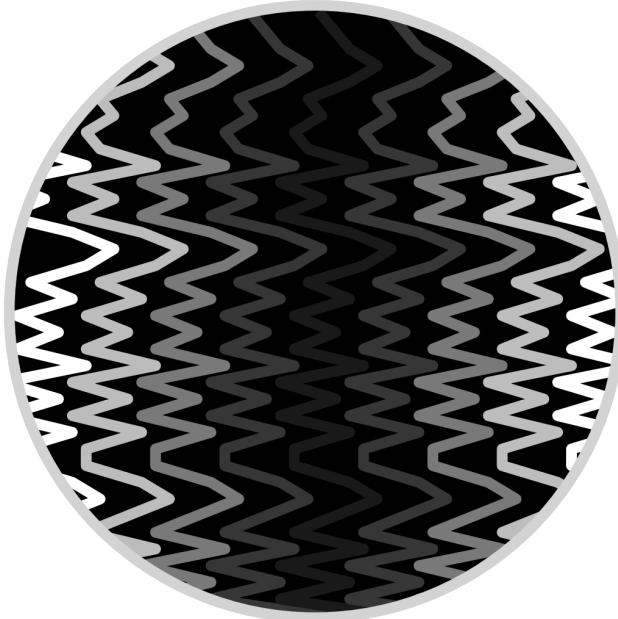
Options specific pollen diagram digitization

-xlim	The x-limits of the data part of the diagram
-ylim	The y-limits of the data part of the diagram
-f, --full	If True, the image is digitized and x- and ylim are set to the entire share of the array Default: False
-rt, --reader-type	Possible choices: area, bars, rounded bars, stacked area, line Specify the reader type. Default: “area” Default: “area”
-V, --version	show program’s version number and exit

STRADITIZE Copyright (C) 2018-2019 Philipp S. Sommer

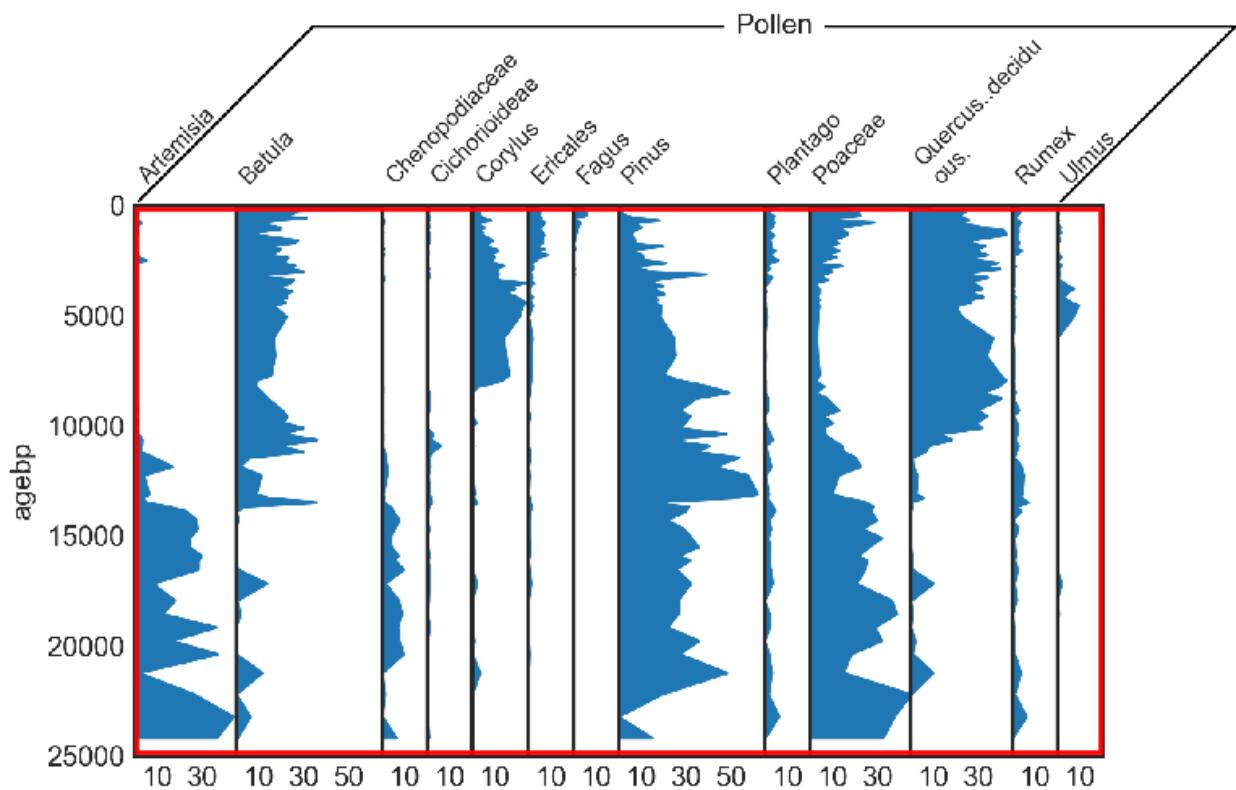
This program comes with ABSOLUTELY NO WARRANTY. This is free software, and you are welcome to redistribute it under the conditions of the GNU GENERAL PUBLIC LICENSE, Version 3.

1.6 Straditize GUI



Welcome to the help explorer of the **Stratigraphic Diagram Digitizer straditize**. Here we guide you through the main functionality of the graphical user interface (GUI) and the necessary steps to digitize your stratigraphic diagram.

1.6.1 Basics and Terminology



A stratigraphic diagram consists of multiple diagrams that are vertically or horizontally aligned, i.e. that have either the same x- or y-axis.

This software assumes, that all the subdiagrams share the y-axis, i.e. the vertical axis. If this is not the case for your specific diagram, just rotate your image by 90°.

column One column in the diagram means one of the subdiagrams (e.g. *Betula* in the example diagram above). The total number of columns is the number of subdiagrams in your plot.

pixel column One pixel column is the horizontal pixel index (x-axis), ranging from 0 to the width of your image. Pixel column 0 represents all pixels on the very left side.

row or pixel row One pixel row (or sometimes only *row*) is the vertical pixel index (y-axis), ranging from 0 to the height of your image. Row 0 represents all pixels at the top.

diagram part This is the part where the data is shown (red rectangle in the figure above).

binary image The binary image is the diagram part converted to black (1) and white (0). Every black pixel in this binary image represents data that should be digitized.

digitize Digitization means transforming each pixel into a value (see the blue line in the image above). Therefore, in contrast to the samples, you get one data value for each pixel row in the binary image. We also refer to this as the *full data*.



sample The samples (red lines in the image above) are a subset of the digitization result. In a very basic sense, the samples represent the original data that was used to create the diagram. In a sediment core for a pollen diagram, it would be one slice of the core that was analyzed under the microscope.

straditizer Everything that refers to the straditizer refers to the full image of your stratigraphic diagram. Speaking in a programmatically sense, it is the top-level interface to digitize the diagram, an instance of the `straditize.straditizer.Straditizer` class. You can always access the current straditizer through the `stradi` variable in the ipython console of the GUI

reader or data reader This is the object that digitizes the diagram part (red area above). For different diagram types (area plot, bar plot, etc.) you have different readers with different functionalities.

The reader of your diagram can be accessed via typing `stradi.data_reader` in the ipython console of the GUI.

1.6.2 Stratitization steps

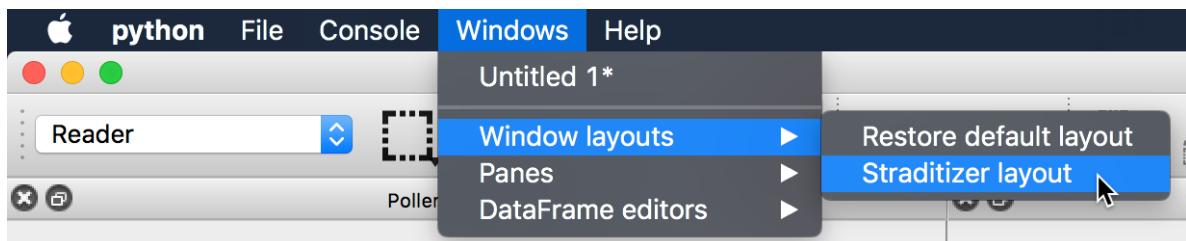
Straditize provides you lot's of possibilities for customizing the semi-automatic digitization of your diagram. However, there are certain steps that you should follow. You may not need all of them and you may exchange them to a certain degree as well.

Load a diagram

The first step is to load the picture of your diagram.

1. If you are starting straditize from the psyplot GUI (i.e. not via typing `straditize` in the terminal or through the app), you should first switch to the straditizer layout via

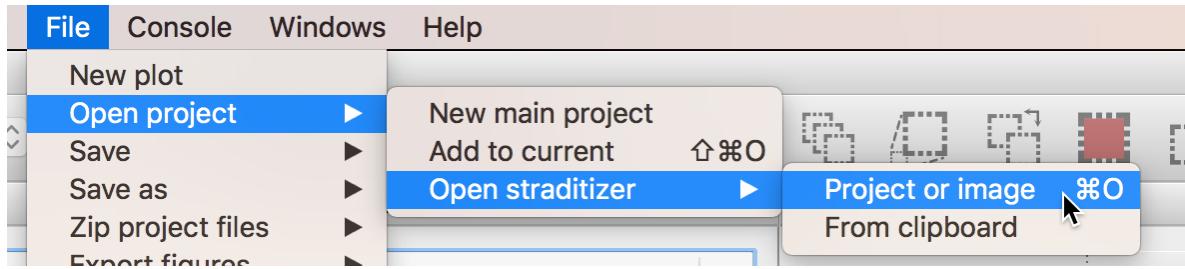
Windows → Window layouts → Straditizer layout.



This will show you the straditizer control panel.

2. To load the picture of your diagram, select

File → Open project → Open straditizer → Project or image



and select the image file you want to load.

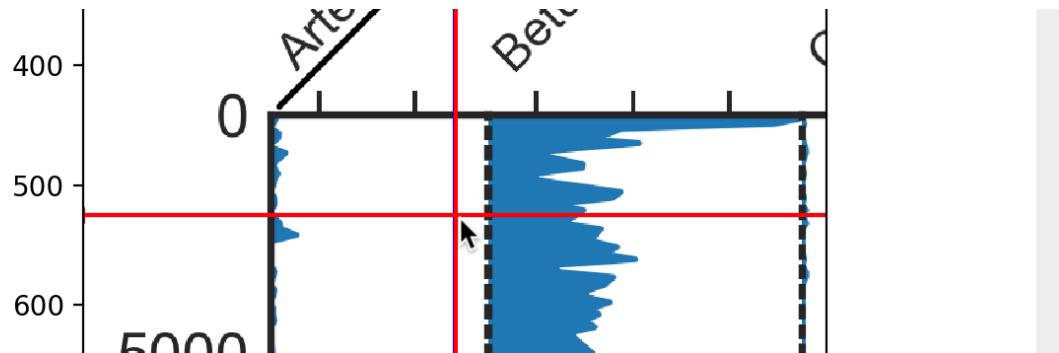
Alternatively, you can use the Control-O (Command-O on MacOS) keyboard shortcut or the  button at the top of the straditizer control.

Select the diagram part

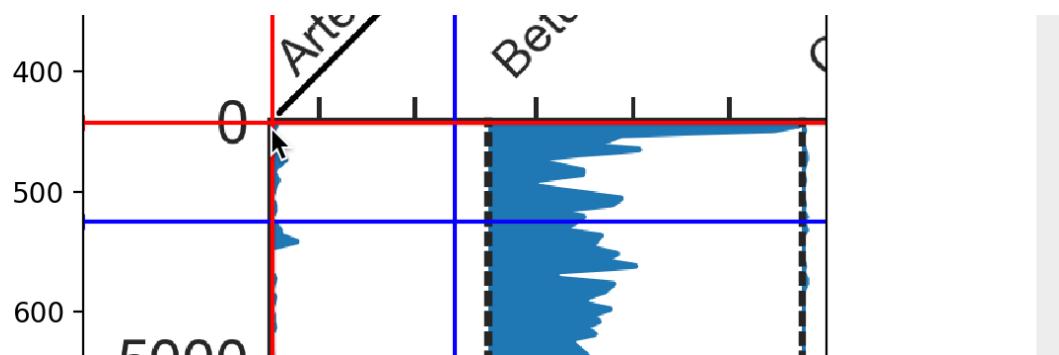
The second step involves selecting the diagram part (see the *Basics and Terminology*) by clicking the *Select data part* button in the digitization control.

The goal is to select the two outer corners of the diagram and try to avoid vertical and horizontal axes or axes description.

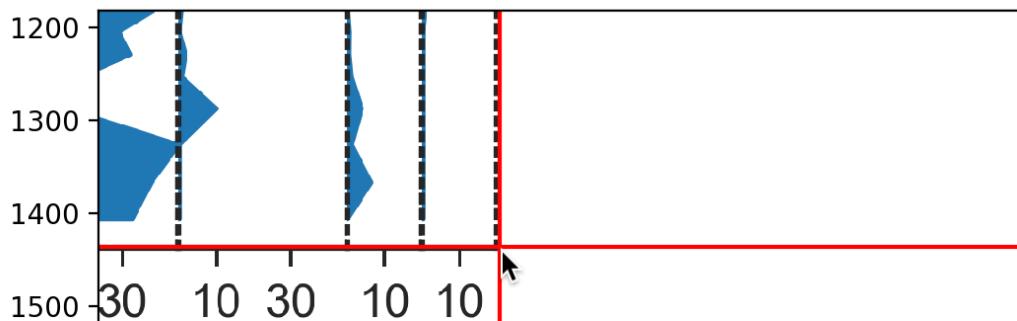
1. Shift - leftclick on the data diagram. This will create a mark at your current mouse location.
2. Click on the cross to select it



3. Hold the left mouse button and drag the cross to the upper left corner of the diagram. Make sure you don't include the y-axis ticks (i.e. the numbers on the y-axis)



4. Create another mark by another Shift - leftclick
5. Move the cross to the lower right corner of the diagram. Make sure you don't include the x-axis ticks (i.e. the numbers on the x-axis)



6. Click the *Apply* button at the bottom

If you want to change the appearance of the marks, see the *Marker control* section in the straditzer control panel.

Selecting the reader

Stratigraphic diagrams and especially pollen diagrams come in different styles and formats.

Straditize supports the following:

area	bars	rounded bars
line	stacked area	

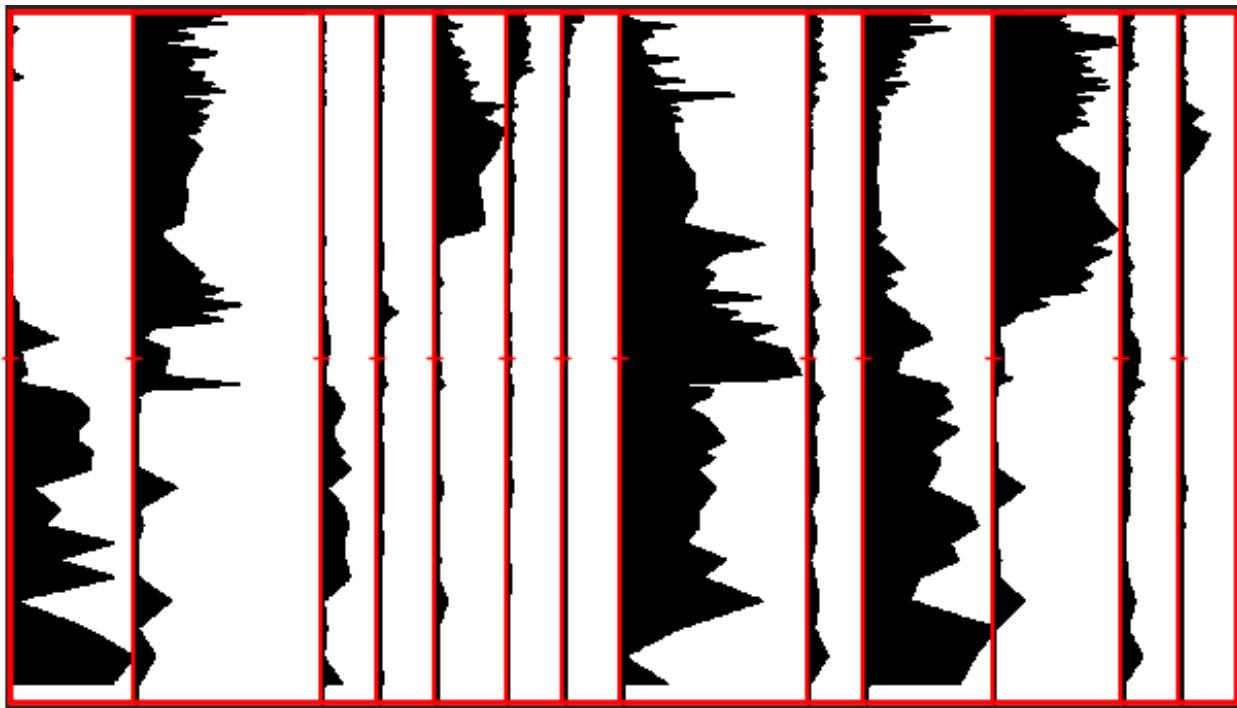
For each type there exists a specific reader that you can select from the drop down menu in the *Reader initialization* section of the straditzer control. If your diagram contains a mixture of types, just choose one type for now and you can then later *create column specific readers*.

After you chose the *diagram part* and selected the reader type (see the table above), you can now click the *Convert image* button. This initializes the reader and converts the data image to a binary image, i.e. black and white.

Everything that is black in this image will be considered as data by the reader. So the next steps will include, cleaning up the image and removing unimportant parts.

Select the starts for each column

A stratigraphic diagram is defined through different columns, each column represents one variable (or for pollen diagrams, one taxa, species, PFT, etc.). To digitize the data, we have to know, where each of these column starts exactly.



After having clicked the *Column starts* button, the software automatically estimates the starts of the columns. However, you should check whether they are all recognized correctly.



There are several possibilities to edit the automatically estimated columns:

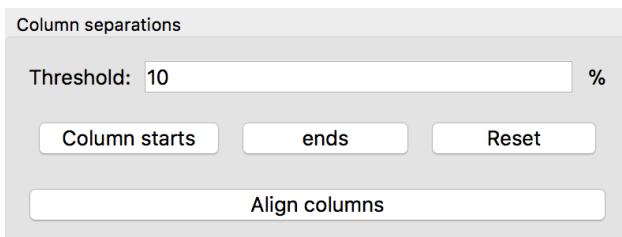
1. Hold the Shift button and left-click on the plot to create a new vertical mark.
Then Left-click the mark and drag it to the start of one column.
2. Delete a column start by right-clicking the mark.
3. Modify the *threshold*. It defines the percentage of a pixel column that has to be covered to assume a valid column start. If you have columns that only contain a very small amount of data, you should lower it

See also:

Select the ends for each column

A pollen diagram is defined through different columns, each column represents one taxa, species, PFT, etc.. To digitize the data, we have to know, where each column starts and ends exactly.

After having clicked the *Column starts* button, the software automatically estimates the starts of the columns and, based on that, the column ends as well. Clicking the *ends* button however lets you modify the automatically estimated column ends.



As for the *column starts*, there are two ways to modify the column ends:

1. Hold the *Shift* button and left-click on the plot to create a new vertical mark.
Then Left-click the mark and drag it to the start of one column.
2. Delete a column start by right-clicking the mark.

See also:

- [Align the columns vertically](#)
- [Select the starts for each column](#)

Align the columns vertically

Sometimes the columns are not aligned properly which makes it hard to identify where the exact sample is located and might lead to misinterpretations of samples. To correct this, we implemented a methodology to align the columns.

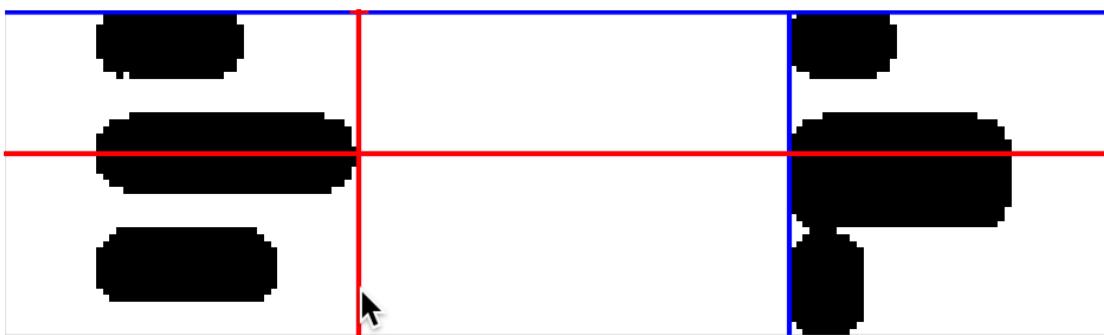
After having clicked the *Align columns* button, multiple marks are created. To correct the vertical displacement, move those marks to the vertical position in each column, that should be on the same vertical level.

You can select the horizontal lines for moving multiple marks and the vertical line for selecting only one mark.

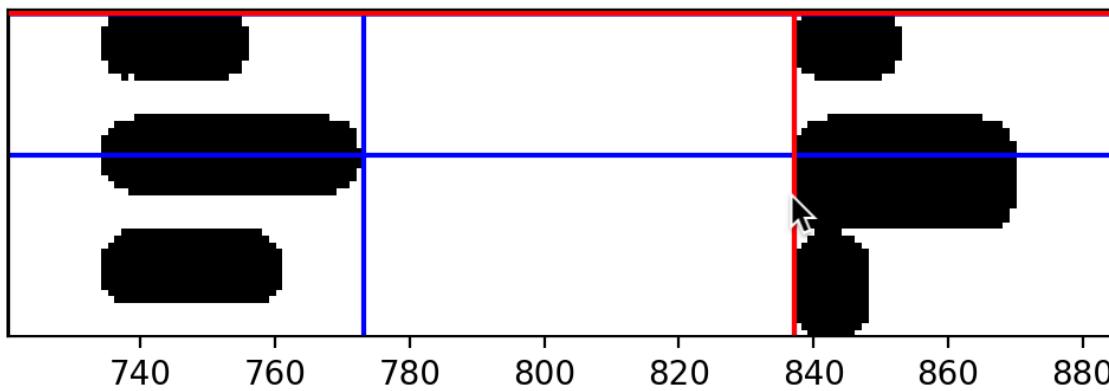
1. Select one mark on one column by clicking on its vertical line



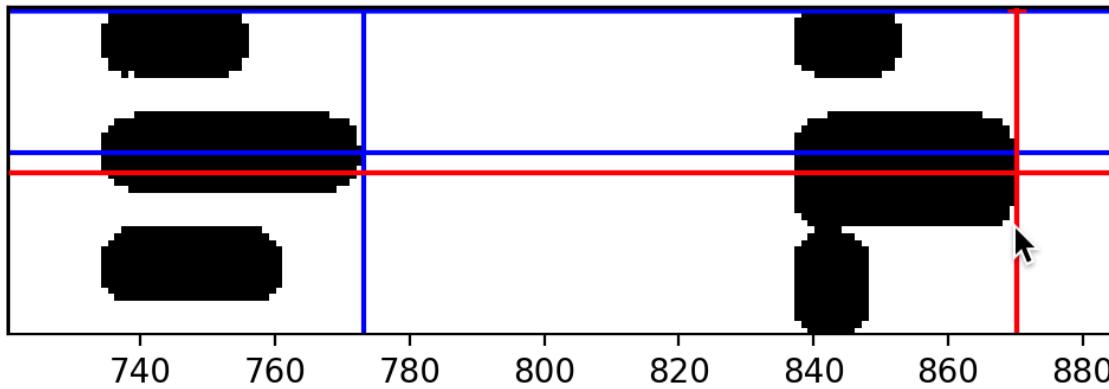
2. Move the mark vertically (the horizontal line does not matter)



3. Select another mark of another column



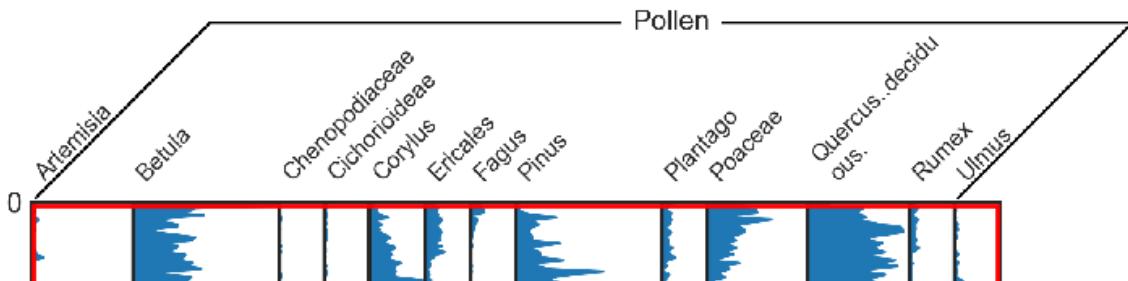
4. Move this mark to a point that should have the same y-value as the first mark



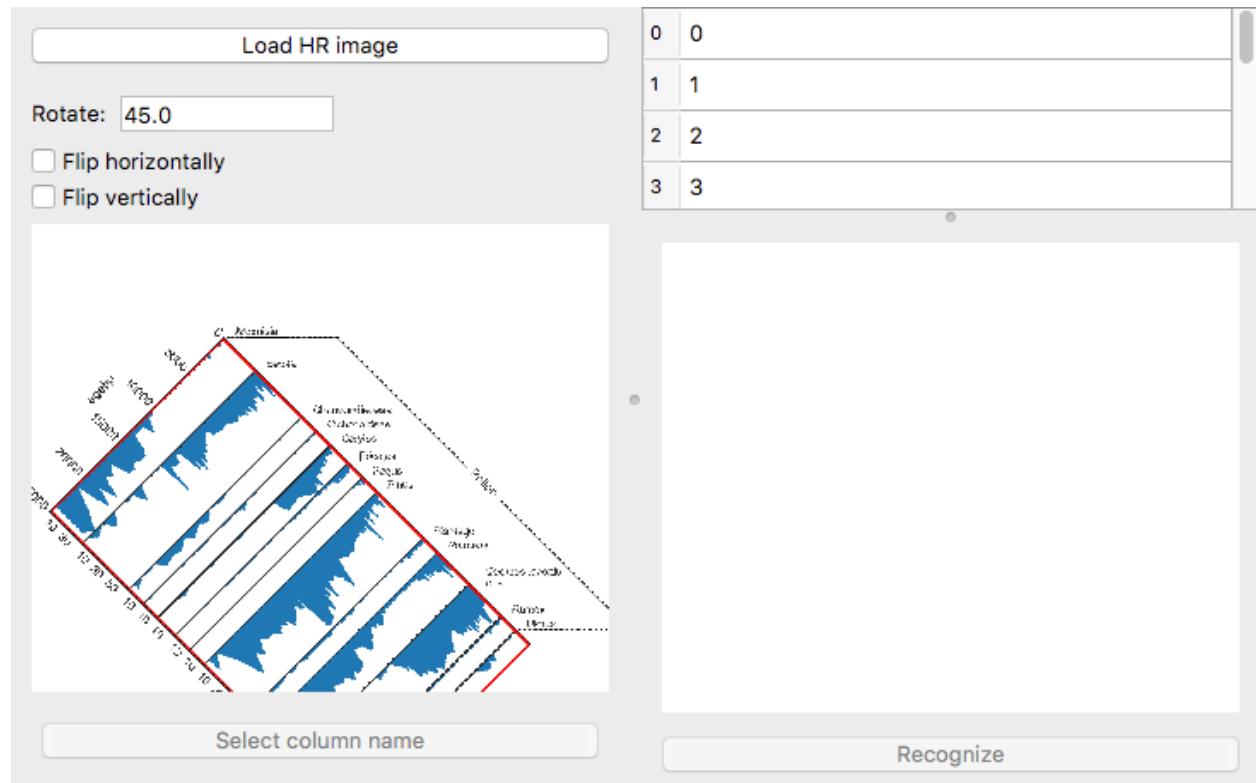
5. Click the *Apply* button

Handling column names

Within straditize, you can assign a name to each of the columns in your diagram, which will then turn up as the header in your final export. Usually these names are the titles of your columns, e.g. such as here



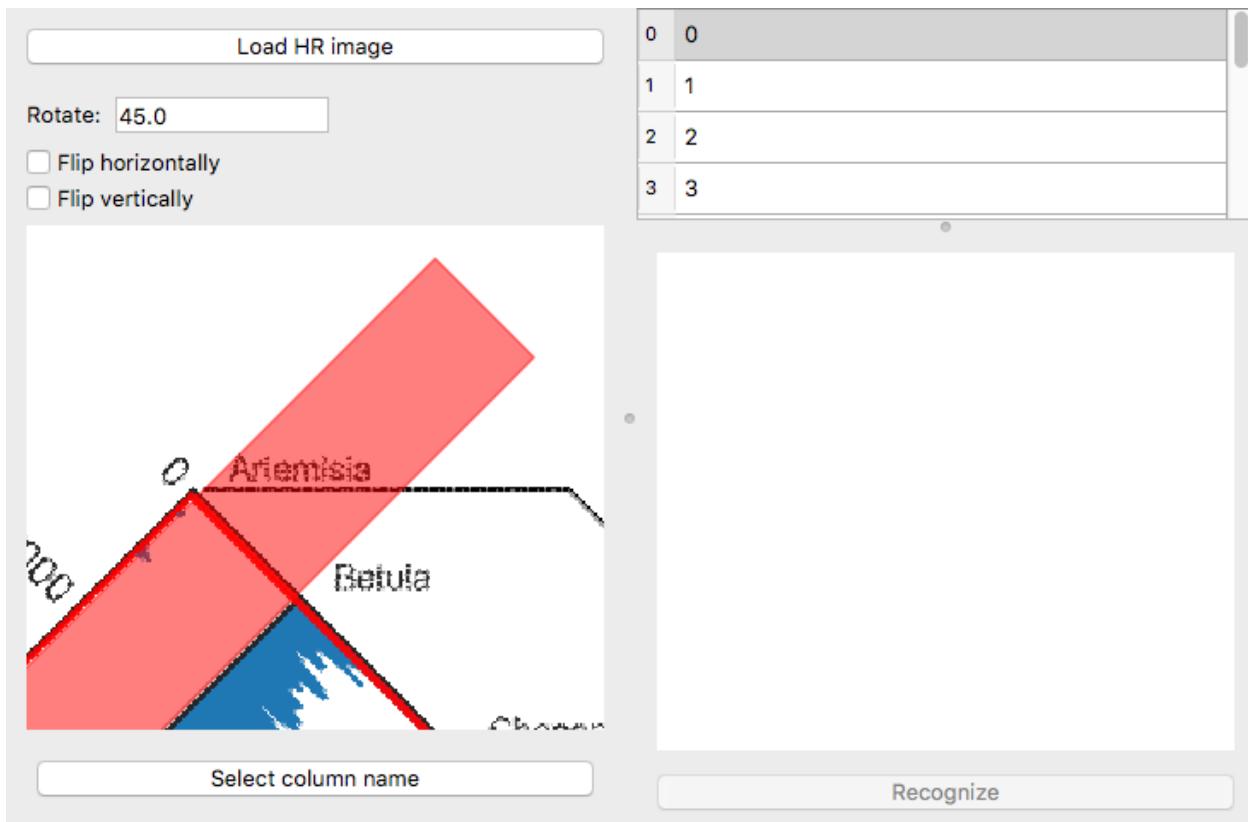
To enter the names of the columns, hit the *Edit column names* button in the *Column names* section of the straditizer control. This will open a dialog to handle the column names:



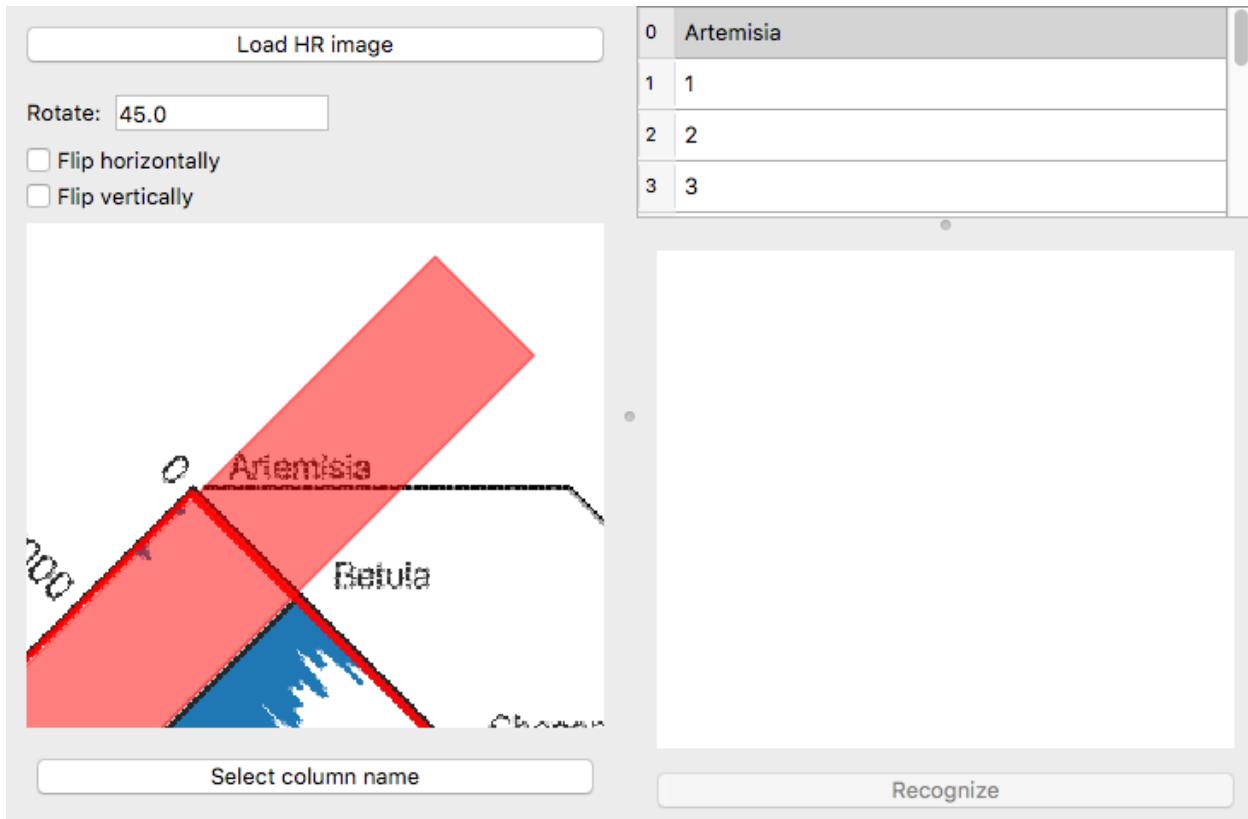
On the left side, you have a rotated version of the image that you can control with the options above it. To navigate in this window, place your mouse over it and then you can move (pan) the image with left-click.

Zooming in and out is done via right-click. To zoom in, hold your right mouse button and move your cursor to the upper right corner of your diagram. To zoom out, move it to the lower left corner.

On the right side you have a table with the column names. Initially, these are just increasing numbers starting from 0 for your first column. If you select one column, it will be highlighted in the left image:



You can now enter the name of the column, in this case Artemisia



Automatic optical character recognition (OCR)

You don't have to type the column name, you can also use the builtin text recognition. For this, we rely on the tesseract software that can be installed (on linux and MacOS) via:

```
conda install -c conda-forge tesseract
```

see the [tesseract feedstock](#) on conda-forge. If you want to automatically find the column names, you also need `tesserocr` to be installed which can be done via:

```
pip install tesserocr
```

or:

```
conda install -c chilipp tesserocr # on linux/MacOS
```

(see <https://pypi.org/project/tesserocr/> for more installation options, in particular for Windows).

1. Use a high resolution version of the image

To improve the result of the text recognition, we recommend to use a sufficient resolution of about 600 dpi. If the image that you are digitizing does not have this resolution, you can optionally load a higher resolution version of it using the *Load HR image* button at the upper left of the dialog.

We also recommend to remove everything in this file but the column names to improve the text recognition.

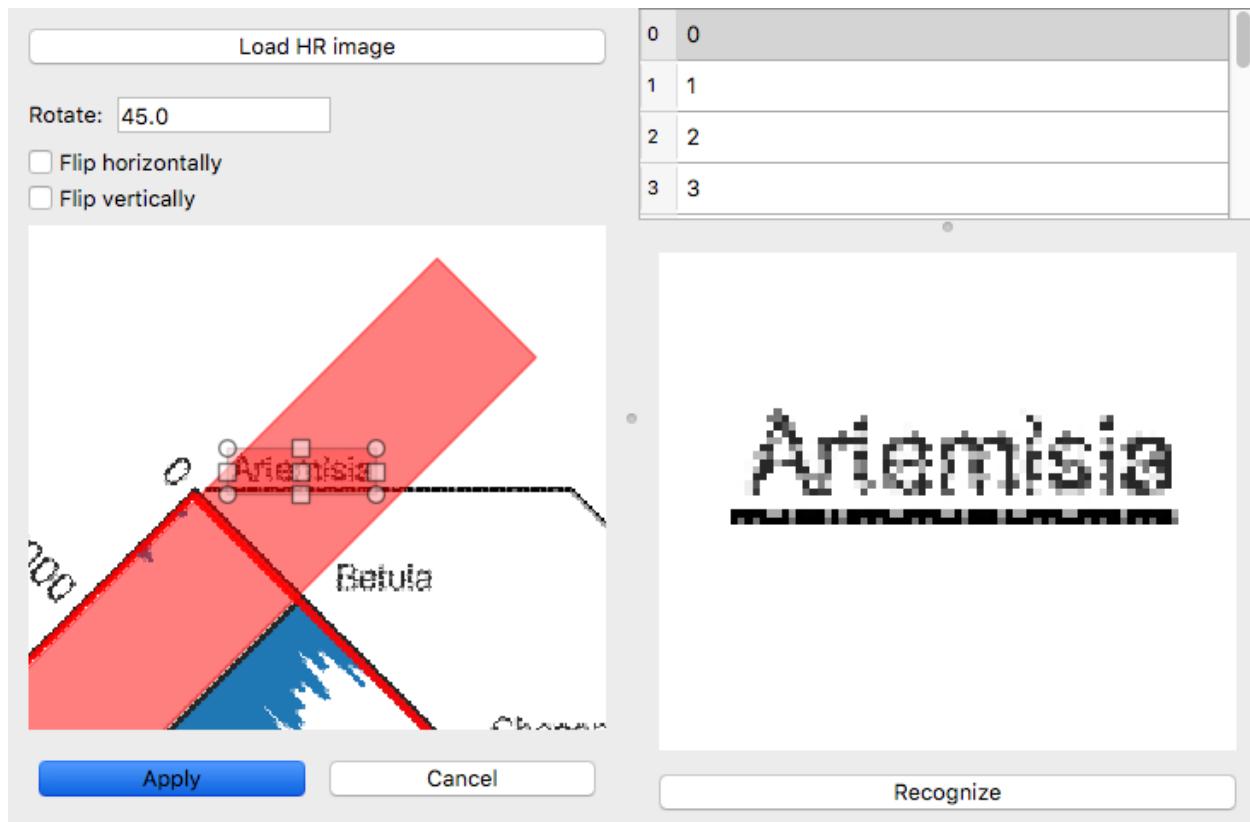
2. Automatically find all column names

Check the *all columns* checkbox and click the *Find column names* button. This then will look for column names in the image that is displayed on the left of the table and insert them into the table.

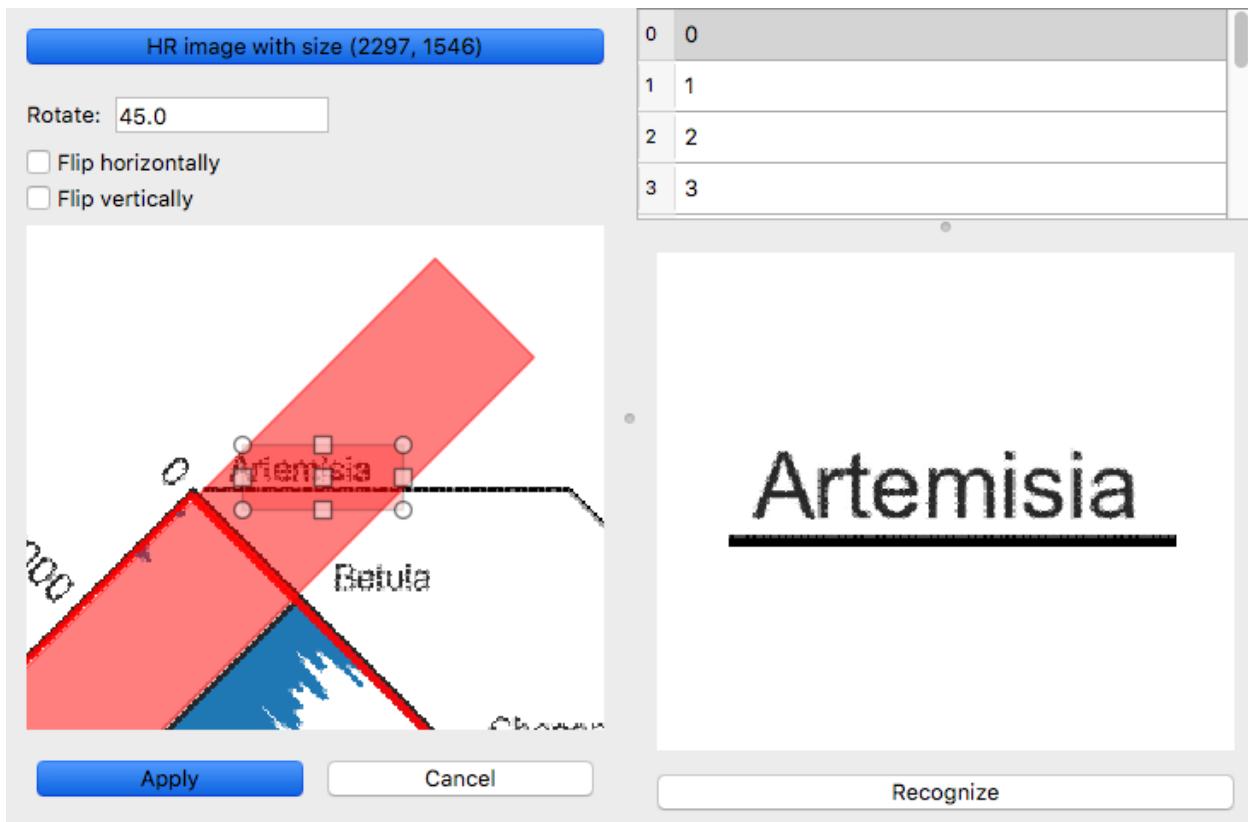
3. Separate treatment of column names

Now, select the first column in the table, zoom in such that you can see the name of the first column and click the *Select column name* button.

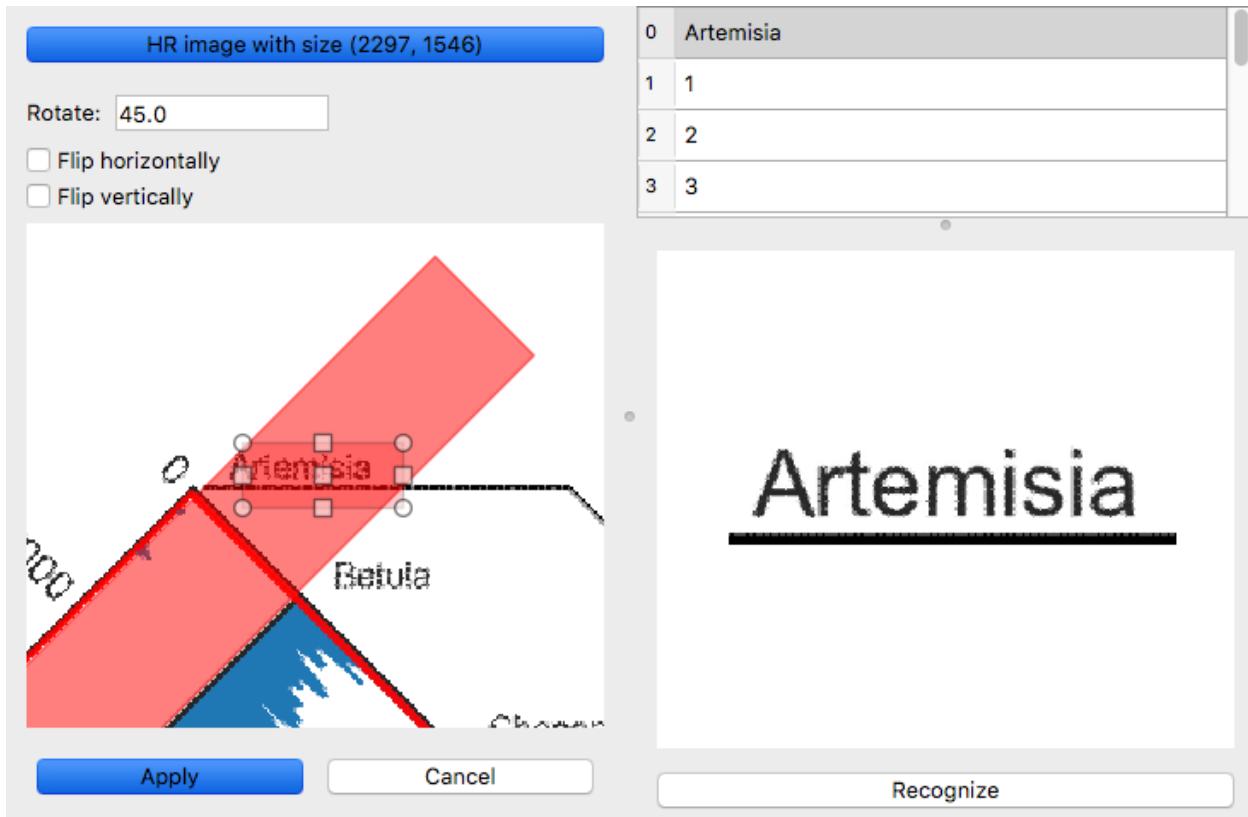
You won't be able to zoom or pan in your plot anymore, but you can now draw a rectangle in the plot around the name of the first column or use click the *Find column names* button (without having the *all columns* checkbox checked) and it will show up in the plot on the lower right of the diagram.



Here you see the result of the low resolution. If we instead specify a higher resolution image (see above), this can look much better



Now click the *Recognize* button at the bottom and it will use tesseract to read the text in the image and fill in the table



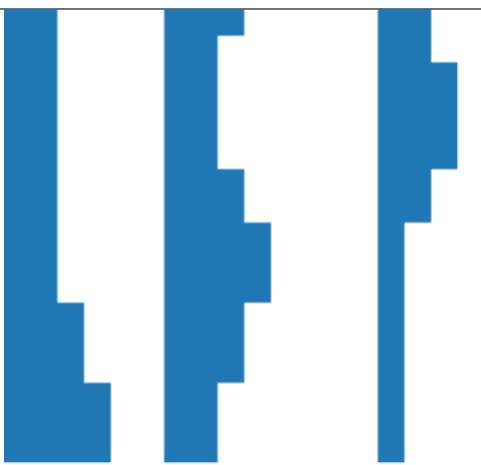
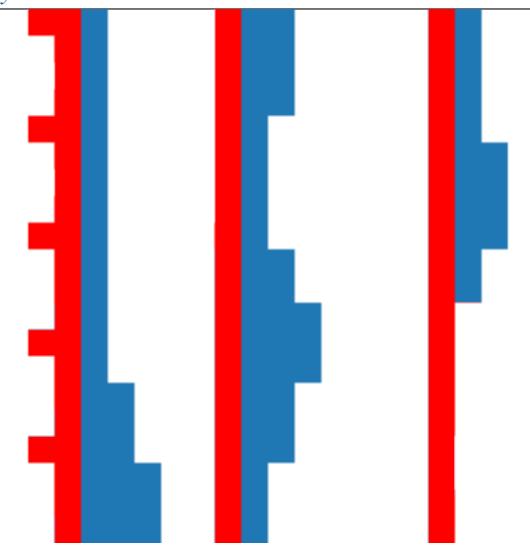
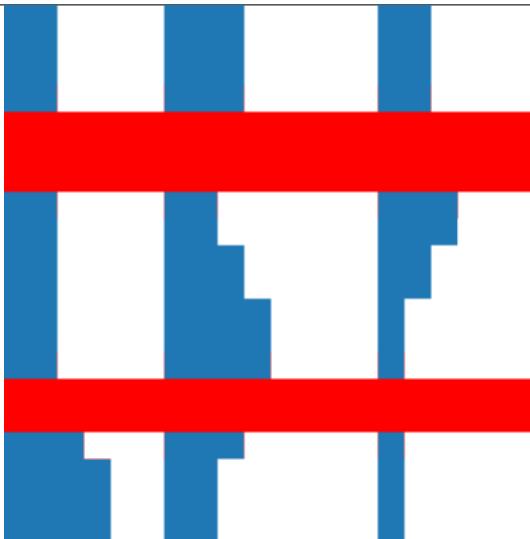
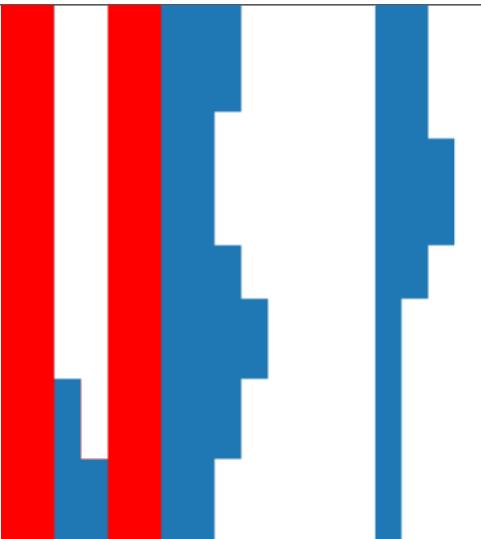
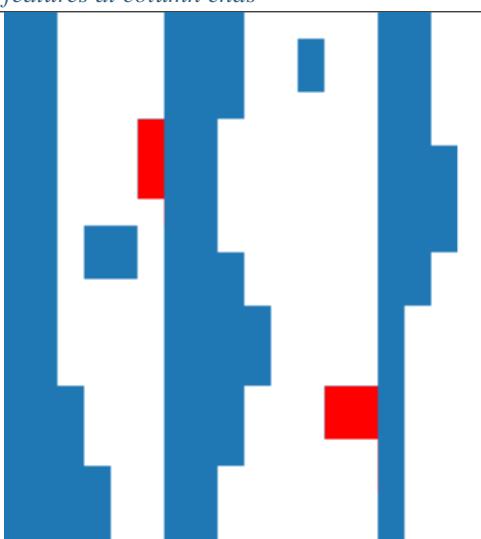
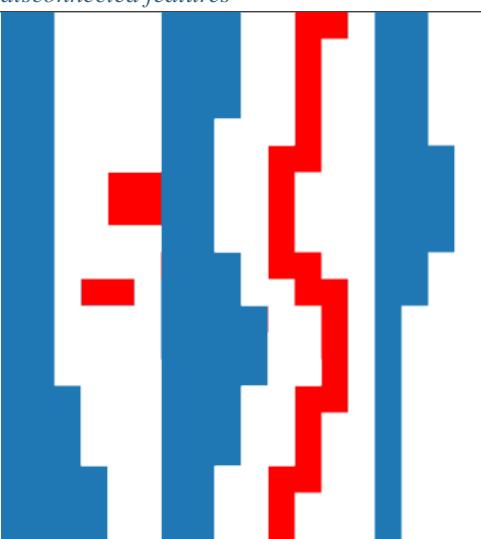
You can now select the next column or click the *Apply* button on the lower left. The latter will reenable the navigation (pan and zoom) in the plot.

Removing features

The diagram part is converted to a binary image where everything black (i.e. 1) represents data (see [Selecting the reader](#)). So before the data is digitized, features that do not represent data have to be removed.

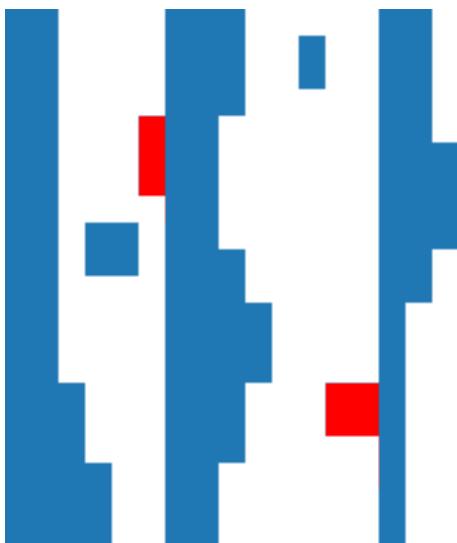
Here, you can either use the [features of the selection toolbar](#), or you use one of the automatic removal tools we provide here. Note that you can edit every selection using the [mouse selection tools](#) or the [automatic tools](#) from the selection toolbar.

After you selected the features to remove, click the *Remove* button at the bottom of the straditizer control panel.

<i>x-axes</i>	<i>y-axes</i>
	
<i>horizontal lines</i>	<i>vertical lines</i>
	
<i>features at column ends</i>	<i>disconnected features</i>
	
<i>cross column features</i>	<i>small artifacts</i>
1.6. Straditize GUI	

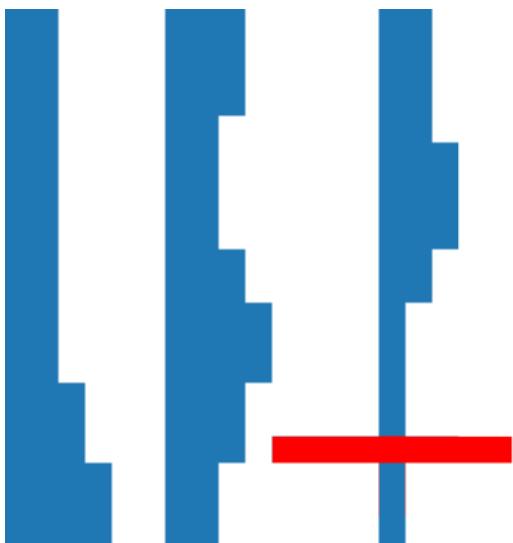
Remove features at column ends

This function automatically finds the features that touch the end selects them for removal.



Remove cross-column features

This function removes features that have a certain amount of pixels in multiple columns. Only those features are selected for removal, that have at least the given *Number of pixels* in more than one column of the stratigraphic diagram.



Note: This feature should be called before initializing column specific readers.

Remove disconnected features

This function automatically recognizes features that are disconnected either from the column start or from the previous feature in the column.

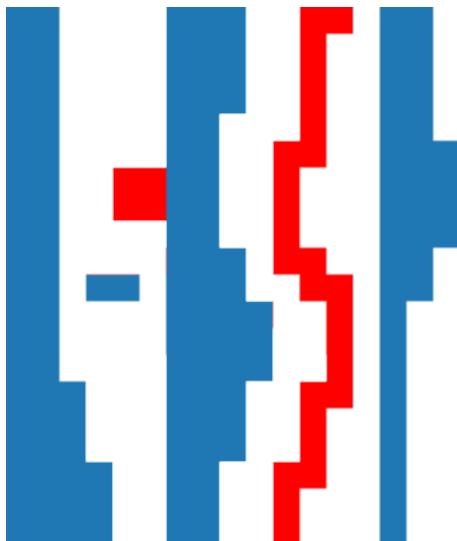
That can be useful, for example, to remove letters or symbols in the diagram.

What features are recognized can be defined in three ways:

1. Features that have a certain distance from the column start (see [Disconnected from the column start](#))
2. Features that have a certain distance from the previous feature in the column (see [Disconnected from the previous feature](#))
3. A combination of 1. and 2. (see [Disconnected from start and previous](#))

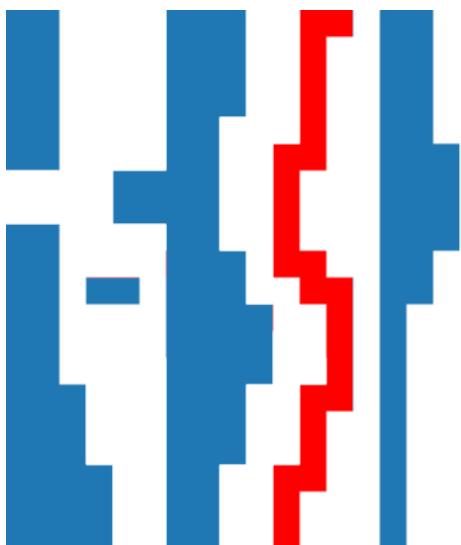
Disconnected from the column start

When only the checkbox *from column start* is checked, all features that have the given distance from the column start are selected.



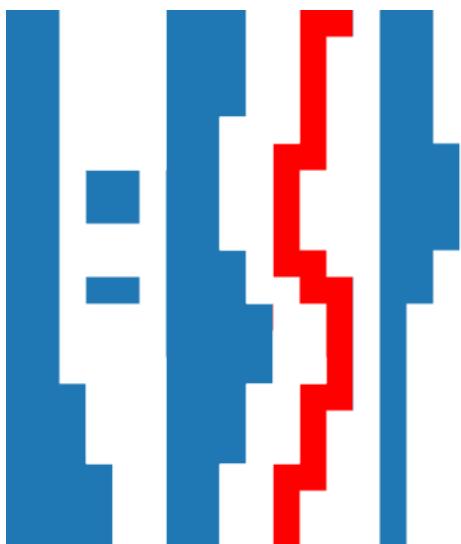
Disconnected from the previous feature

When only the checkbox *from previous feature* is selected, all features that have the given distance from the previous feature in the column (and everything to the right of it) is selected.



Disconnected from start and previous

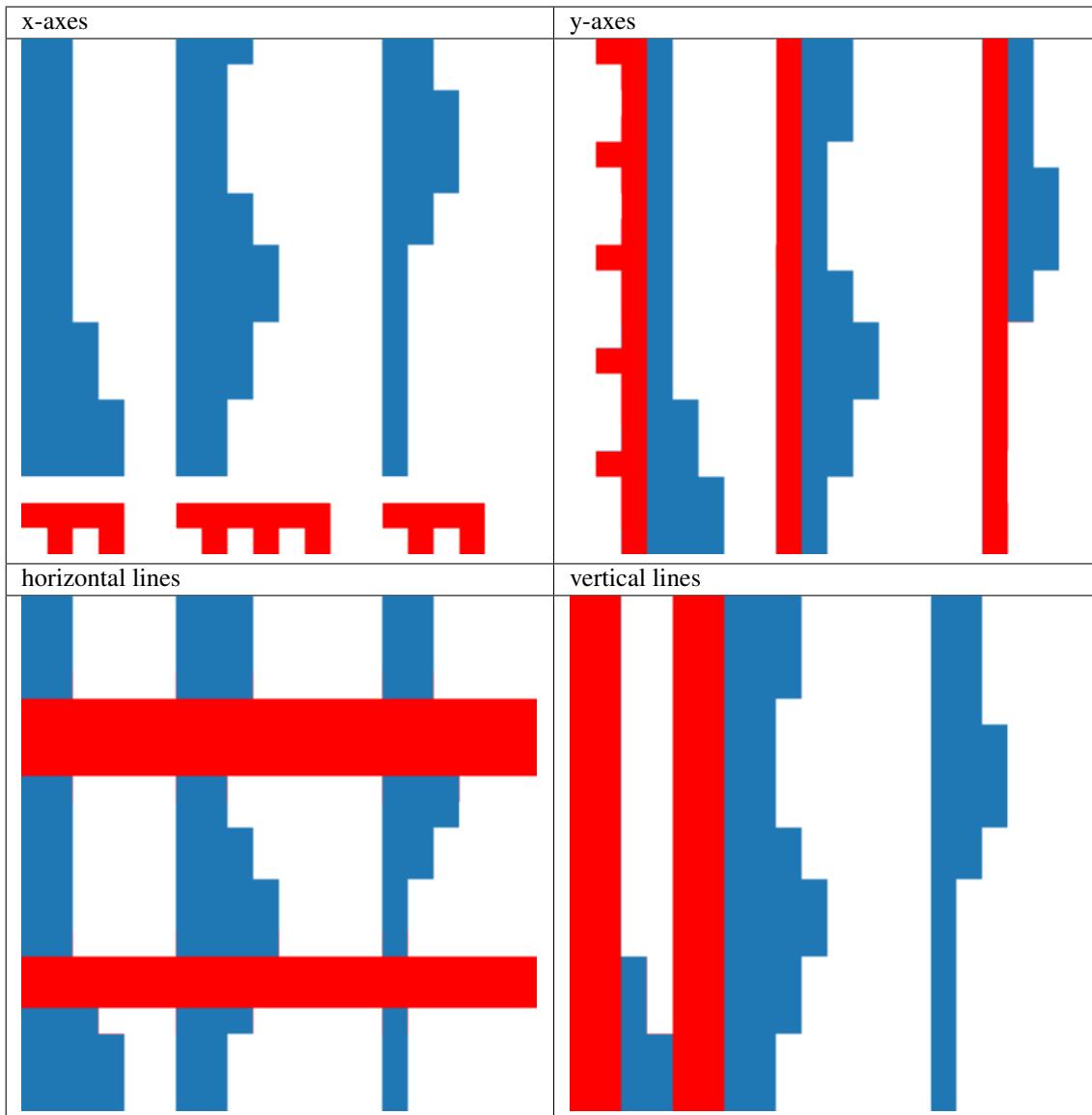
If both are selected, the selected feature must have the given distance from the column start and from the previous feature in the column to be selected.



Remove horizontal or vertical lines

The features described here automatically remove the y- and x-axes, or in general vertical or horizontal lines that span a certain fraction of the diagram part.

The methodology is simple: If a certain pixel column or row in the binary data image is covered, it is considered as a vertical or horizontal line, respectively.



You can modify the recognition using three options

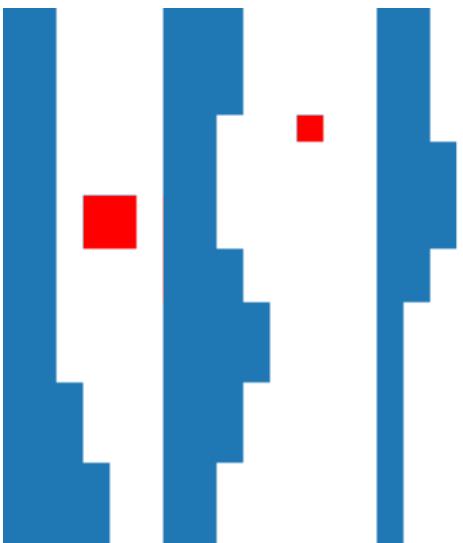
Minimum fraction: This is the minimum fraction of a pixel column (for or row that must be covered)

Minimum line width: Lines are only selected, if their line width is greater than or equal to the given minimum axis width.

Maximum line width: Lines are only selected up to the given maximum axis width

Remove small artifacts

This method recognizes small artifacts using the `skimage.morphology.remove_small_objects()` method.



The pixel size of the artifacts can be determined through the *Smaller than* field and you can always modify the selection through the tools in the selection toolbar.

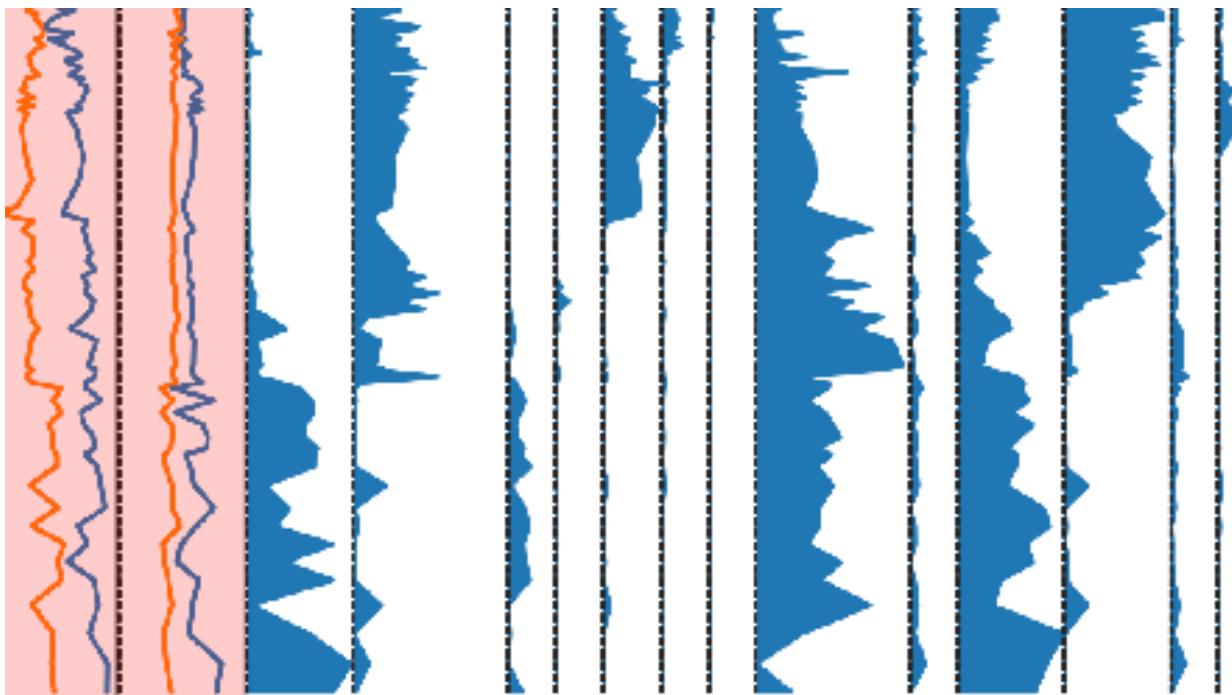
If you want a better visualization of the small artifacts that will be removed, use the *Highlight selection smaller than* button.

Column specific readers

Note: If the entire diagram is just based on one diagram type (see *Selecting the reader*), you can skip this step.

As noted in the *reader selection section*, you can choose different readers for different columns in order to account for multiple subdiagram styles within the stratigraphic diagram. The current reader, i.e. the one that is selected in the *Current reader* tab in the *Digitization control* is the one that can be accessed via the `stradi.data_reader` attribute in the console. It is also sometimes referred to as the *parent reader*.

Clicking the + button in the *Current reader* tab let's you select columns just by clicking on the data image.

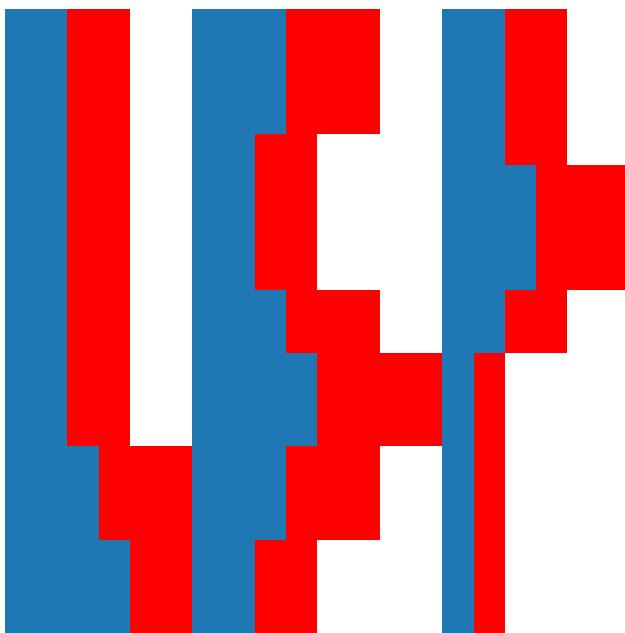


After you selected the necessary columns, hit the *Apply* button. In the appearing window, select the reader type you want (see [Selecting the reader](#)). You now initialized a new child reader for the selected columns that can be set as the parent reader using the dropdown menu in the *Current reader* tab.

Interpreting exaggerations

Visualizing an exaggerated version of the data is a common methodology in pollen diagrams, since even small pollen counts can say a lot about the environment.

Therefore it is common to not only plot the data, but also to exaggerate it by a certain factor, e.g. 10 or two like the red areas in the image below.



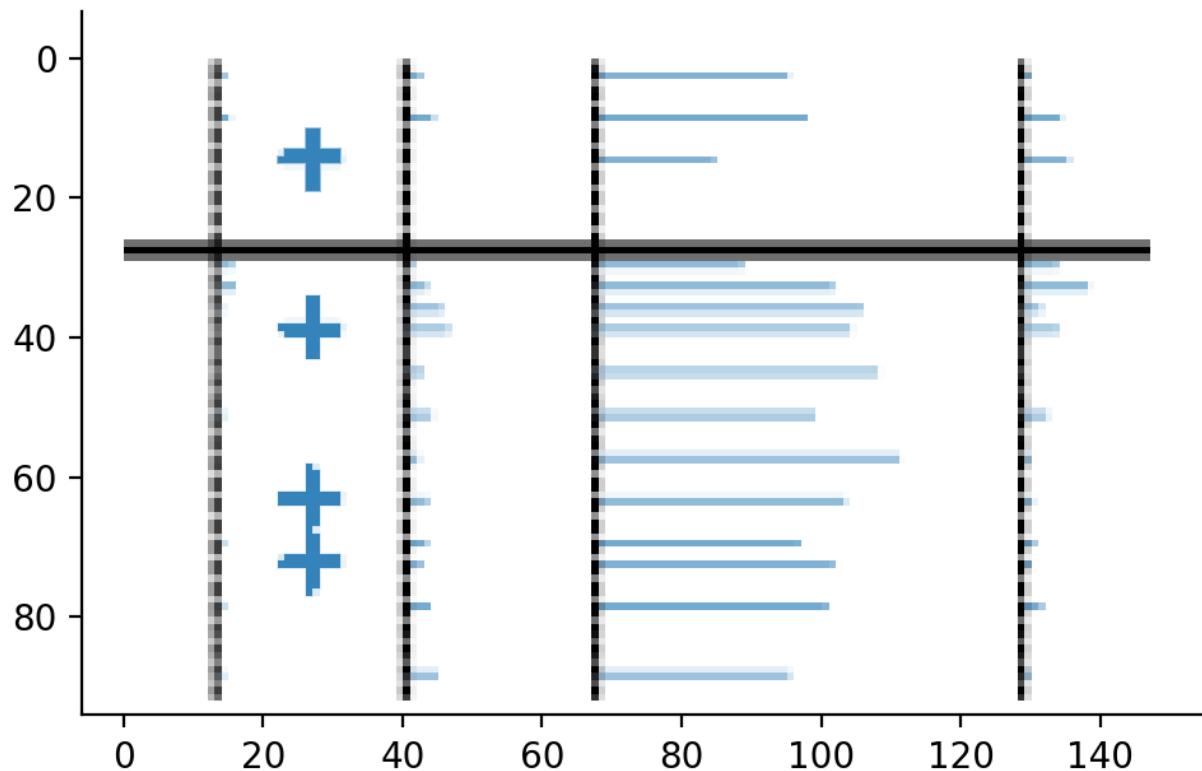
You have two choices: Either you remove these exaggerated areas (see [Removing features](#)), or you merge them into your results to improve the digitization result.

To include and interpret the exaggerations, straditiz can create a new reader dedicated for the exaggerations which you then have to select and specify when they should be used. This is all done in the *Exaggerations* tab of your digitization control.

1. Specify the *exaggeration factor*. This number is usually described in the caption of your diagram and must be greater than 1.
2. Select the data reader type for the exaggerations (see [Selecting the reader](#)). Most common this is one of *area* or *line*.
3. Initialize the reader for the exaggerations by clicking **+**. You will now find a new reader dedicated to the exaggerations in the dropdown menu of the *Current reader* tab. But for now, we stick with the original one.
4. Now, you then have to select the exaggerations. Click the *Select exaggerations* button and use the tools in the *selection toolbar* for selecting the features in the diagram that represent the exaggerations. When you're done with this, click the *Select* button. You can also repeat this step to select more and more features.
5. Specify when the exaggerations should be used. This can be either where the unexaggerated data is below a certain percentage of the image width and/or below a certain number of pixels.
6. When you now click the *Digitize exaggerations* button (but after you clicked the *Digitize* button for the original reader), the exaggerations will be digitized and merged into the digitization result of the non-exaggerated reader.

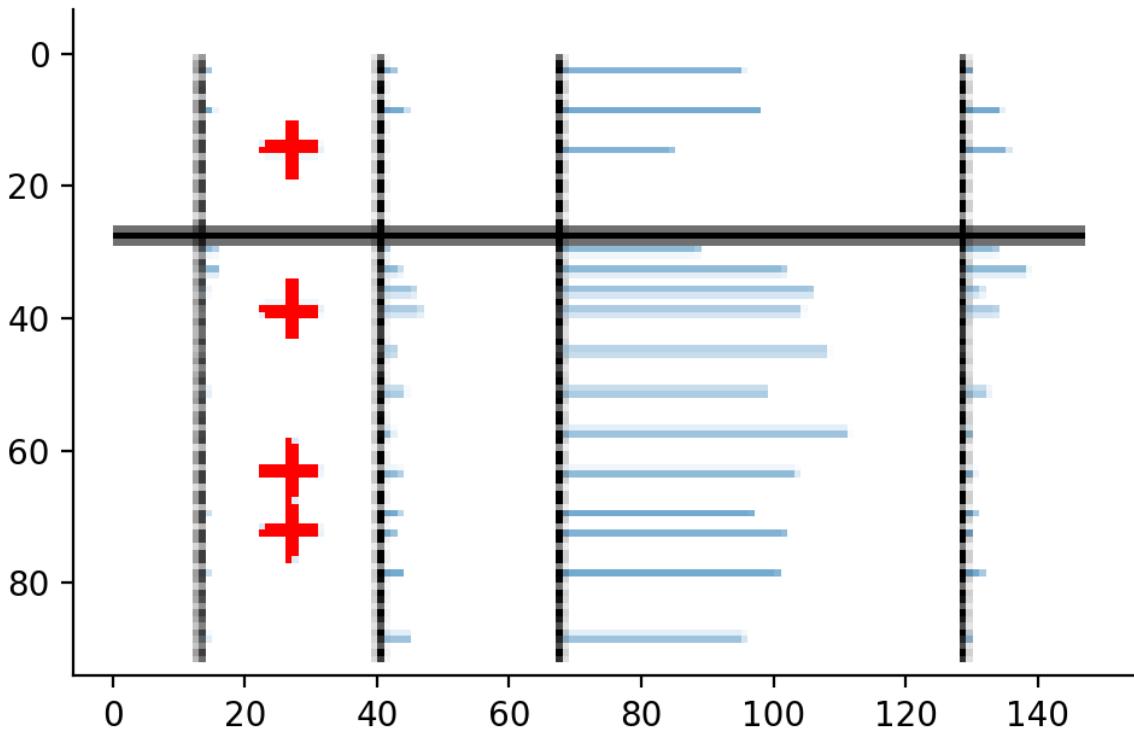
Select occurrences

Pollen diagrams often mark low taxon percentages to highlight the occurrence of a taxon, such as the **+** in the image below

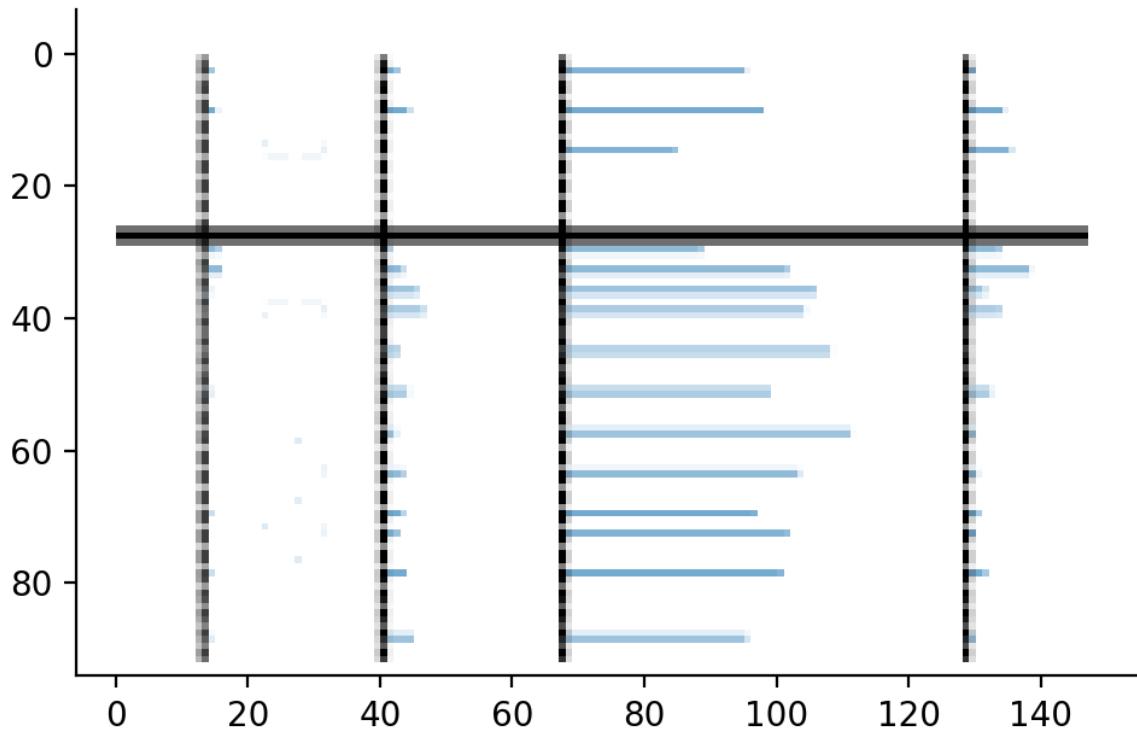


This is very useful information which can be considered by straditize.

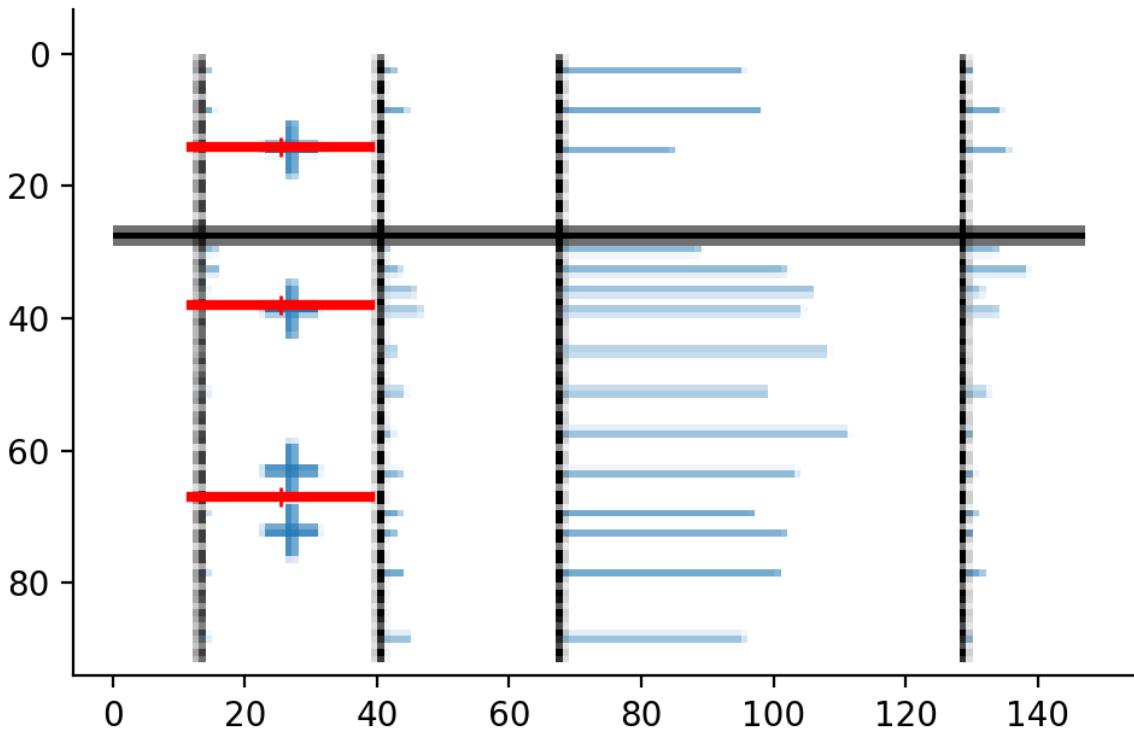
1. Expand the *Occurrences* tab in the *Digitization control* section
2. Click the *Select occurrences* button and select the occurrences using the tools from the *selection toolbar*, especially the  tool is useful



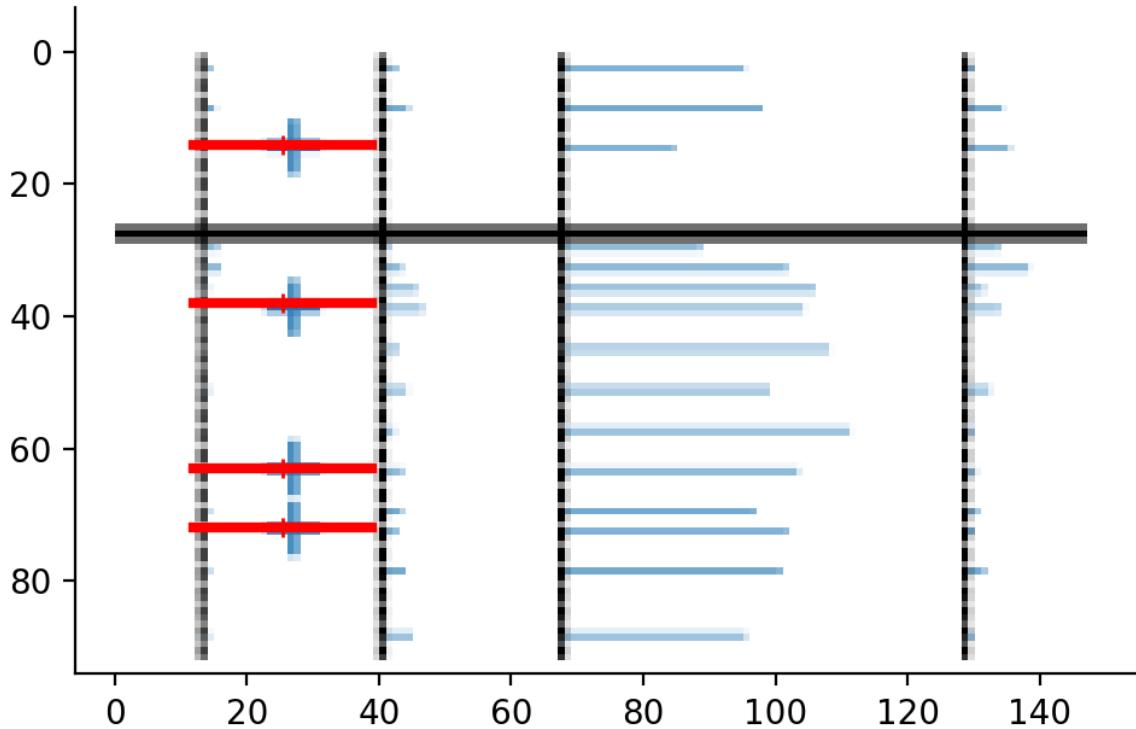
3. Click the *Apply button* and the selected features in the diagram will be registered as an *occurrence*. If you also checked the *Remove on apply* checkbox next the to *Select occurrences* button (which you normally should), the selected features in the diagram will be removed, too and you will end up with a plot like this



4. You can see and edit the occurrences using the *Edit occurrences* button. This will create horizontal markers at the positions of the original occurrences



In our demonstration, straditiz could not distinguish two of the occurrence markers because they were too close to each other. But you can just move the mark with Left-click and Shift+Leftclick on the plot to create a new marker



Click the *Apply* button when you are done.

5. The marked occurrences will appear later in the final data frame with the specified *Occurrence value*.

Digitizing the diagram

The digitization transforms your picture into a table. The outcome is a data frame with one value for each row and column in the data part of your image. After the digitization, you can access this raw data from the command line via:

```
stradi.data_reader.full_df
```

and you can plot it using the *Full digitized data* item in the *plot control*.

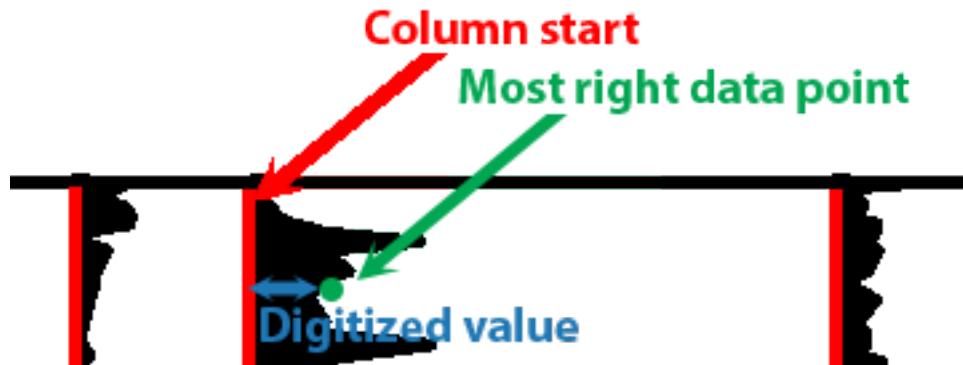


The digitization procedure depends on the reader type that you are using (see *Selecting the reader*).

Contents

- *Digitizing the diagram*
 - *Digitizing area and line readers*
 - *Digitizing bars*
 - * *Splitting bars*
 - *Digitizing stacked areas*

Digitizing area and line readers



This class of stratigraphic diagrams is the most popular within the pollen diagrams and it's digitization is fully automatic.

The algorithm simply uses the distance of the most right data (black) pixel to the column start.

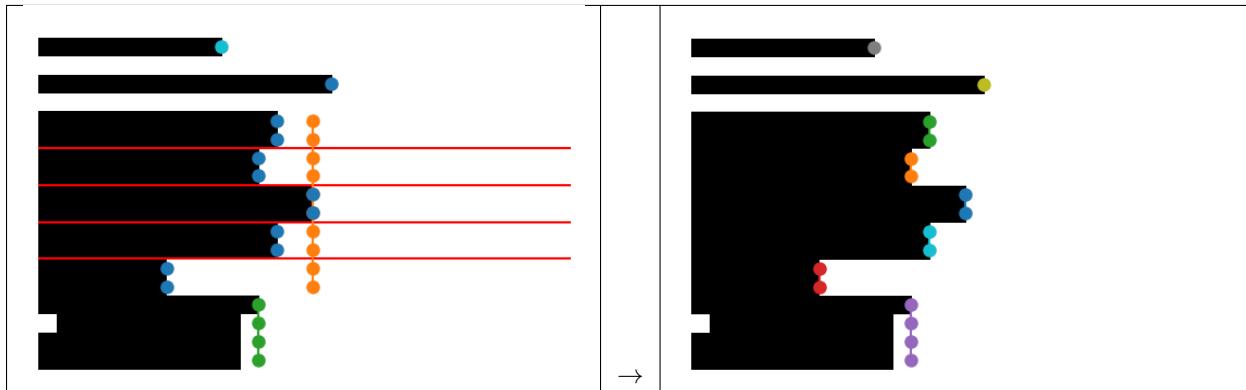
Digitizing bars



For digitizing bar diagrams, it is crucial that the software finds and distinguishes adjacent bars. This is mainly done through the *tolerance*. The algorithm is: if the pixel row, compared to the previous row, is higher or smaller by the given *tolerance*, the software assumes that those two rows belong to two distinct bars.

The image above illustrates this. The reader distinguishes the bars that are clearly separated or were one bar is significantly lower or higher than the other, but it cannot distinguish bars where there is only a small difference in height.

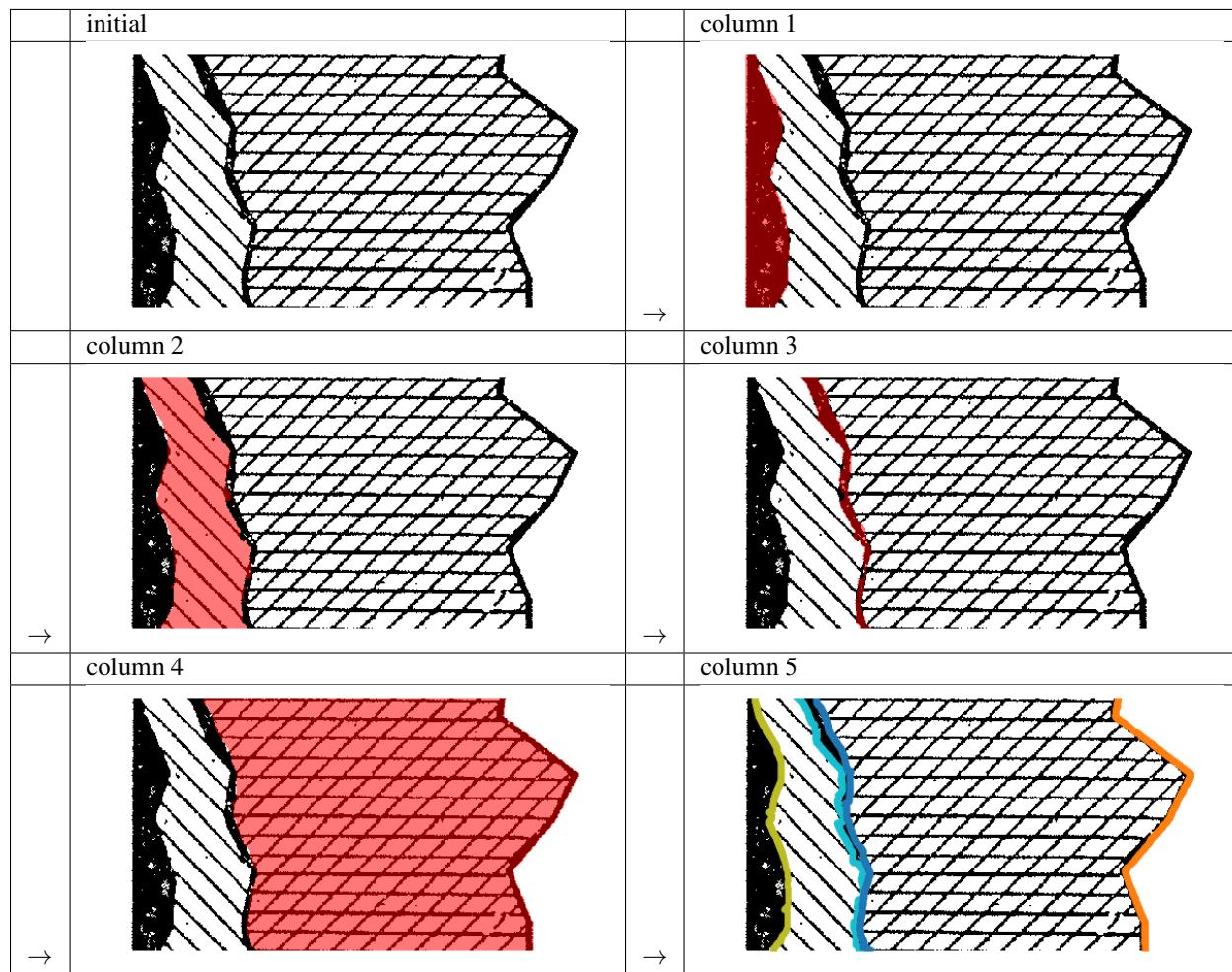
Splitting bars



To overcome this problem, straditize automatically finds the bars that are wider than the others and marks them for splitting. If the straditizer finds bars that are too long, they are shown in the *Split too long bars* tab of the digitization control. To split one bar:

1. Double-click an item in the table (the plot will zoom to the bar)
2. Left-click on the bar to indicate the vertical location for a split (right-click again on the same line to revert this). A red line will appear (left-click, see image above) or disappear (right-click)
3. Click the symbol to continue with the next bar that is too long or double-click an item in the table. Again split the bar as described in 2.
4. When you are done, hit the *Apply* button and the bars will be splitted right were you marked it

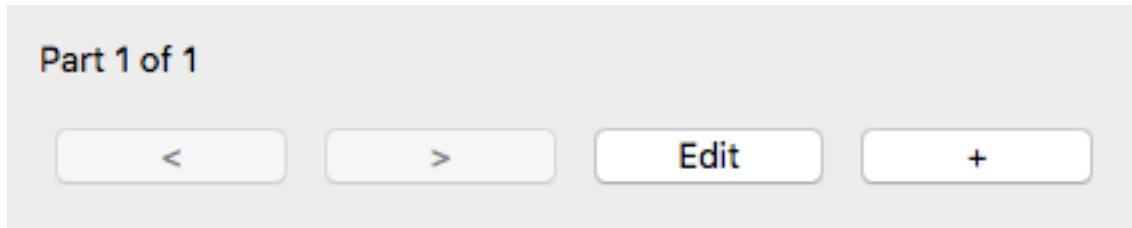
Digitizing stacked areas



Stacked diagrams are more difficult to digitize because of the variety of combinations. The columns might be distinguished by colors or patterns (hatches). Therefore, the user has to select the features corresponding to one column, one after the other.

The digitization here works as follows:

1. click the *Digitize* button, you will see a control for navigating within the columns



2. click the + button to setup a new column. Everything in the column will be selected and now you have to select only those features, that are part of the first column using the selection tools (see the *column 1* above).
3. When you did this, click the *Apply* button.
4. click the *Edit* button if you want to edit the column that you just selected

5. Click the > button to go to the right column
6. click the + to add a new column and select the second column (see above)
7. repeat the steps 3-6 until you digitized all columns

If you want to display your digitization result, use the *Full digitized data* in the *plot control*.

Hint: you can change the transparency of the selection with the transparency slider in the *selection toolbar*. That helps to still see the selected features

Samples

samples are the source data that has been used to create the diagram (see *Basics and Terminology*). The crucial part after the digitization of a diagram is to find out, where these samples exactly are.

Straditiz assists you with this through automatic sample finding algorithms, or you can load the sample locations from an external files or you just add and edit the samples manually.

Automatic samples identification

straditiz has reader specific sample identification algorithms implemented. The overall algorithm is based on two steps:

1. For each column: Identify the intervals that contain exactly one samples (the rough locations)
2. Align the overlapping intervals between the columns to estimate the exact location

Where step 2 is the same for all diagram types, step one is diagram specific. For bar diagrams, we use the bars that have been identified during the digitization (see *Digitizing bars*).

For area and line diagrams on the other hand, we identify the local extrema (minima and maxima) for each column. This strategy works well for pollen diagrams since they are supposed to sum up to 100%. However, this does not generally work for all stratigraphic diagrams.

Load samples locations from external file

samples can be loaded from an external CSV (comma-separated) file. The first column in this CSV file is expected to represent the y-locations of the samples. If the y-axis scale has already been specified (see *Translating the shared vertical axis*), this data should be on the same scale. Otherwise, we assume that the data is in pixel coordinates.

Warning: Make sure, that you first entered the correct *y-axis scale*. Otherwise the samples might not be interpreted correctly.

Editing samples

To edit samples, click the *Edit samples* button and a new figure will be created with one mark for each sample and a table is shown to you with the samples data. To edit the samples, you can edit the table or the marks in the plots

The sample table

Measurements editor

Zoom to selection Selection only Fit selected cells to selected data Plot reconstruction

	y	Column 0	Column 1	Column 2	Column 3	Column 4	Column 5	Column 6
1	3	1	36	3	4	14	12	6
2	12	0	20	2	4	9	12	4
3	21	0	30	3	3	18	12	3
4	31	0	39	2	4	22	5	2
5	35	0	45	3	4	18	4	3
6	40	0	38	3	4	20	4	2
7	48	0	37	2	3	26	3	2
8	52	0	34	2	3	32	5	2
9	57	0	28	2	3	35	4	2
10	63	0	34	2	3	32	3	2
11	80	0	26	2	3	22	4	2
12	87	0	27	2	3	23	4	2
13	97	0	25	2	3	25	4	2
14	102	0	19	2	3	22	4	2
15	110	0	19	3	4	3	3	2
16	121	0	32	2	4	3	2	2
--	--	--	--	--	--	--	--	--

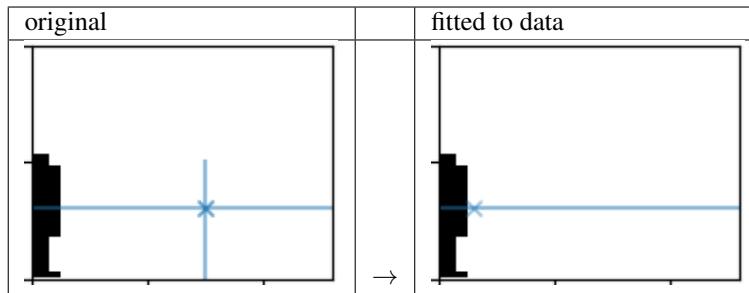
%0.6g

The sample table shows you the vertical coordinates of the samples and the values for each column obtained through the digitization.

Each row in this table represents one sample in the diagram.

The first column is the vertical location (y-axis) of the sample, the remaining columns are the x-values of the samples. The numbers shown in the table are in pixel coordinates of the diagram part. To interact with the samples, you can

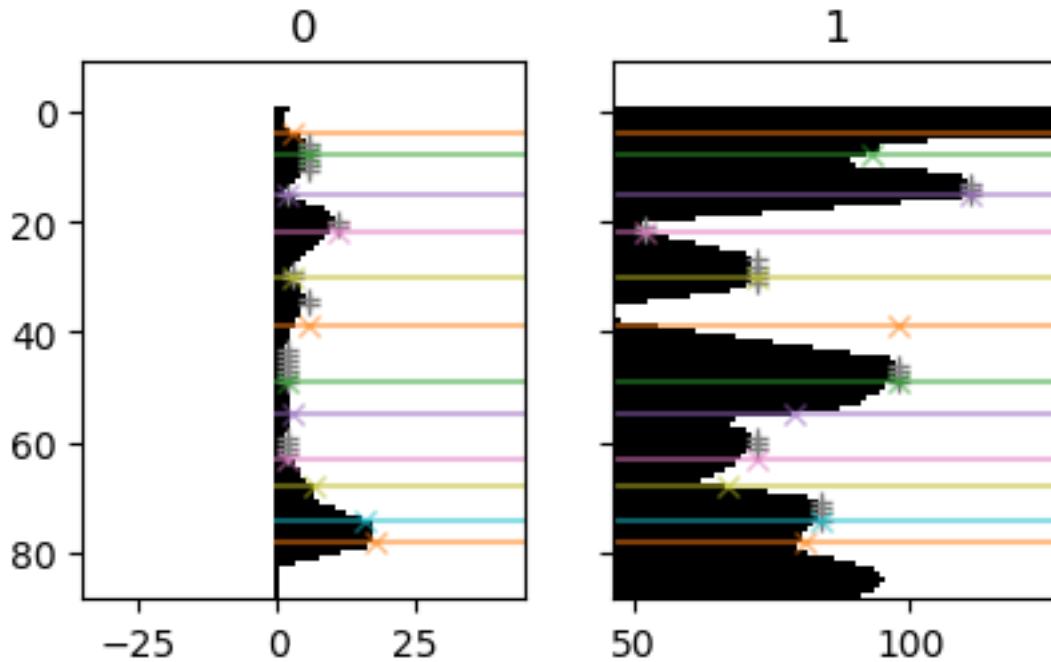
1. Right click the table to add new or delete old samples/rows below or above the selected row
2. Edit the numbers in the cells to change the values for the samples.
3. Fit the x-values of the samples to the data.
 - i. Either by right-click and *Fit selected cells to the data*, which will adjust the column values of the mark such, that it fits with the digitized value



- ii. Or by using the *Fit selected cells to selected data* checkbox. If this is checked:
1. select cells in the table
 2. click on the plot
 3. the cells will be updated with the x-values from the digitization at the click-position

You can also zoom the selection or hide everything else but the selection.

Editing the marks



Additional to the table, you have a visual representation of the samples in the figure. Here, you have one plot per column each mark represents the vertical location of a sample. The marker x in the line shows you the value of the sample (i.e. the location on the x-axes).

1. move a sample by
 - i. left-click a mark and hold the mouse button (the mark will change its color)
 - ii. while still holding the mouse button, drag the mark to a different location
2. delete a sample by right-clicking the mark

3. add a new sample by holding down the Shift button and left-click on one of the plots.

Translating from pixel to data coordinates

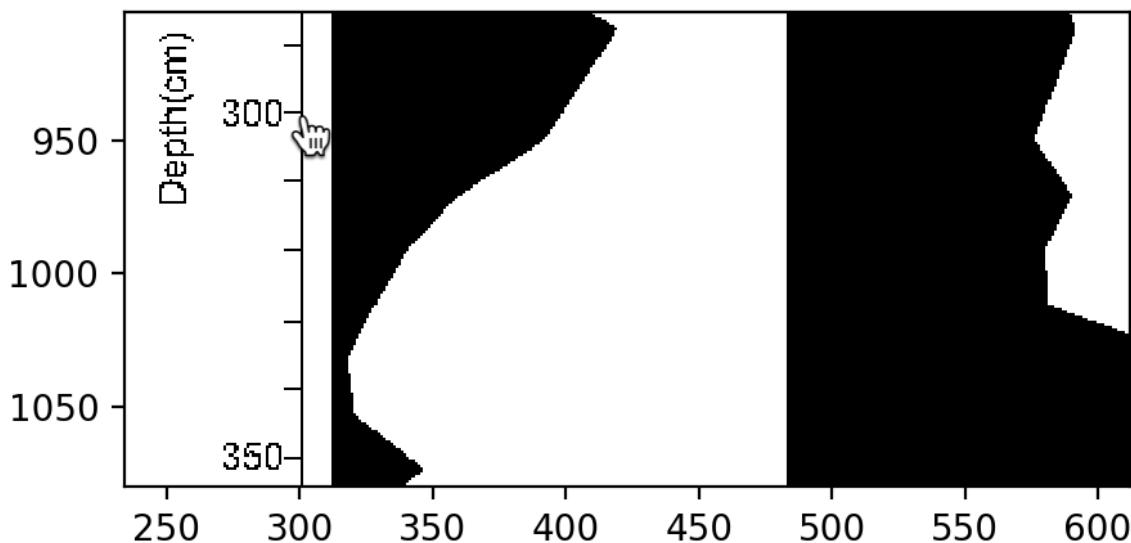
All the previous steps in the digitization were in pixel coordinates. However, for the final output, we need to translate this into the real units of the diagram.

Translating the shared vertical axis

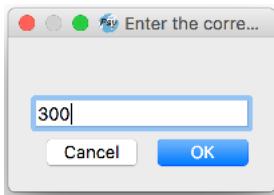
For your final product, straditize needs to know about how the data in the diagram is scaled, such that we can transform the final sample information from the pixel coordinates into the units of the diagram (e.g. years or meters in case of time or depth).

To translate the y-axis information,

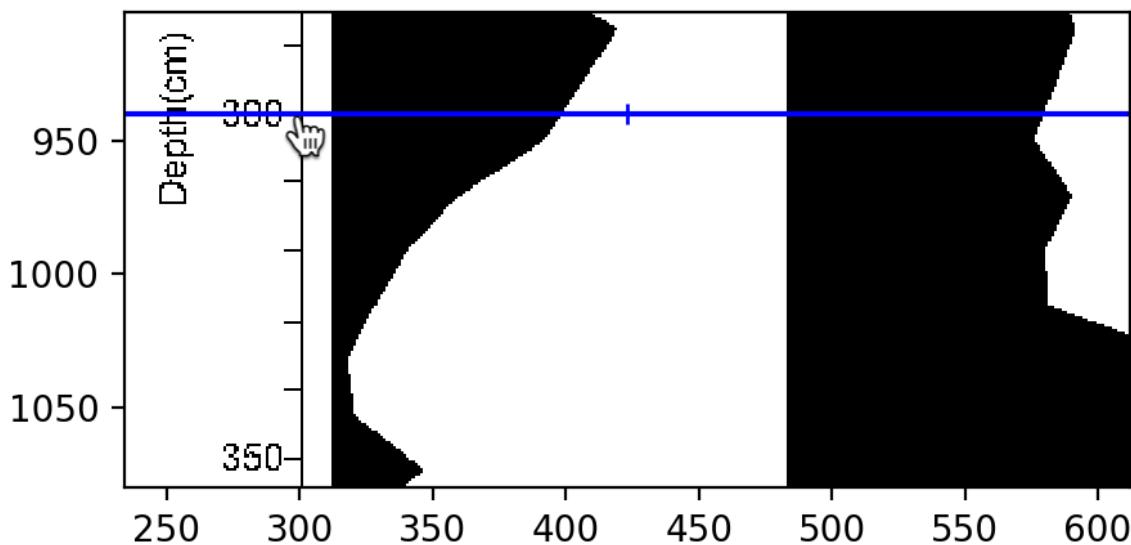
1. Expand the *Axes translations* tab in the digitization control
2. Click the *Insert Y-axis values* button in the *Axes translations* section of the straditzer control (if not already done)
3. Shift-leftclick on the plot to enter the corresponding y-value.



4. A small dialog will appear where you should enter the y-value to use (in this case, 300)

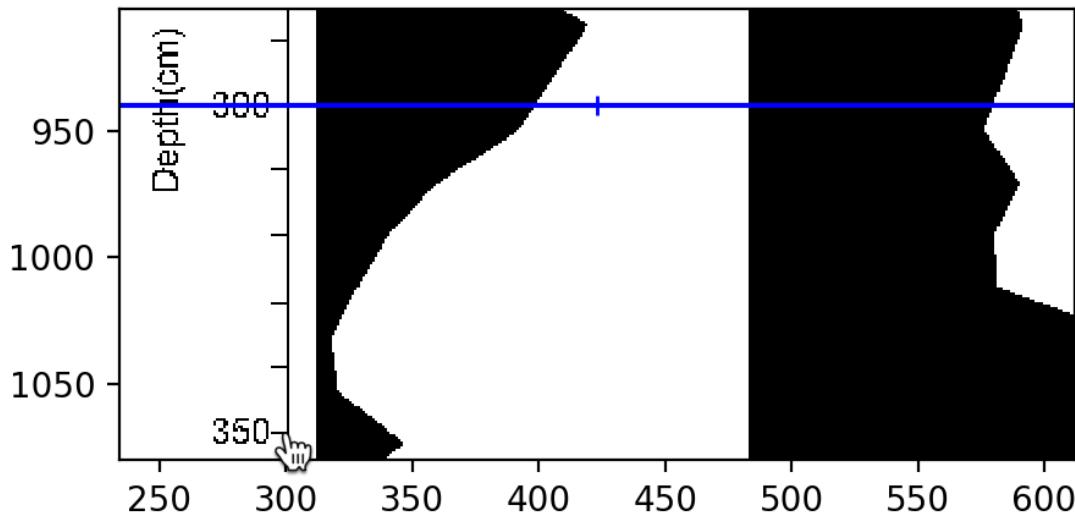


5. After hitting the *Ok* button, you will see a mark on the plot (blue line). You can select the mark via leftclick and drag it to a different location or you can delete it via rightclick.

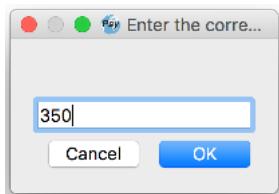


6. now repeat steps 2-4 on a second point on the y-axis

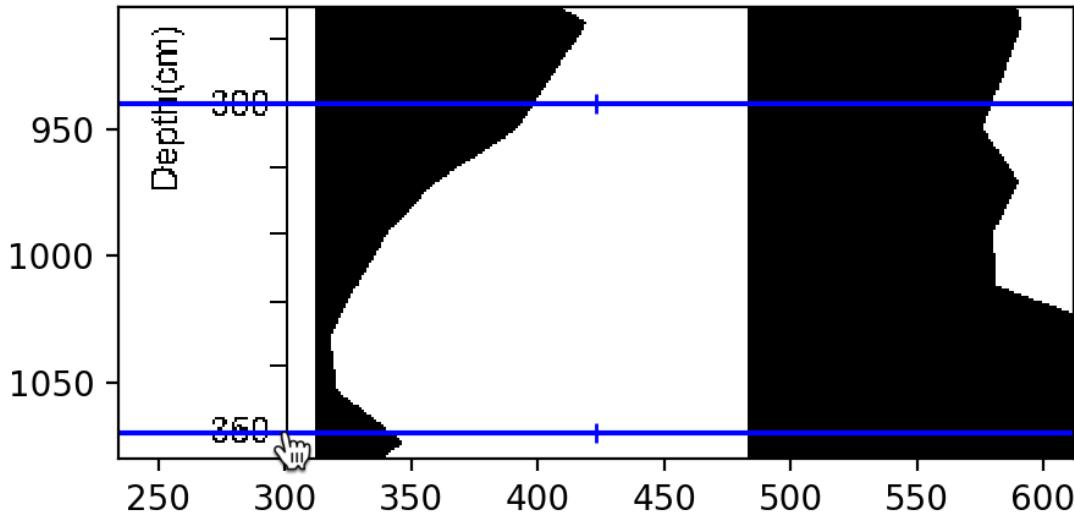
- Select another point



- Enter the corresponding value (here 350)



- A new mark is created that you can modify



7. Click the *Apply* button at the bottom of the straditizer control when you are done.

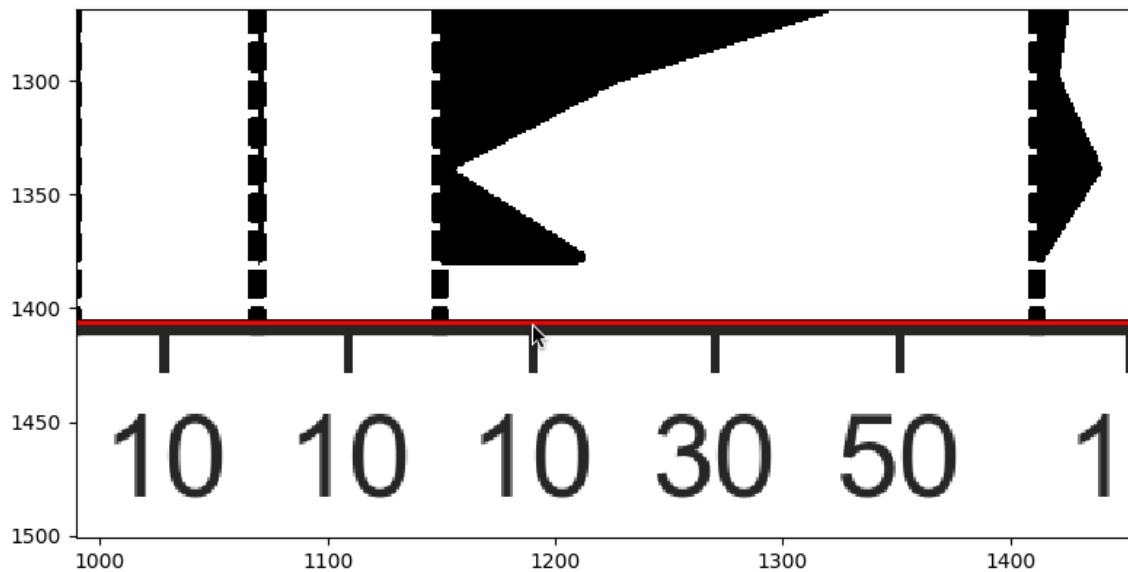
Note: If you drag a mark and hold the Shift button while releasing the mouse button, the dialog in point 3 from above will not pop up.

Translating the horizontal axes

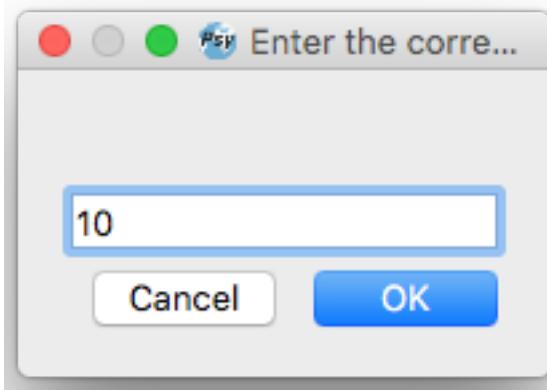
For your final product, straditize needs to know about how the data in the diagram is scaled, such that we can transform the final sample information from the pixel coordinates into the units of the diagram (e.g. percent for pollen data or Kelvin for temperature, etc.).

To translate the x-axis information,

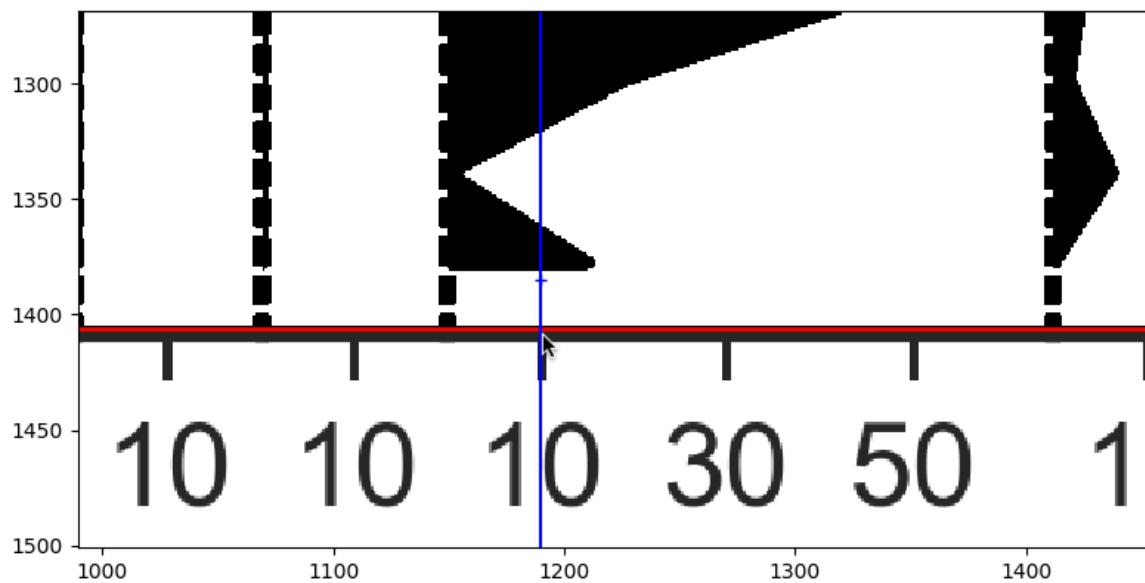
1. Expand the *Axes translations* tab in the digitization control
2. Click the *Insert X-axis values* button in the *Axes translations* section of the straditizer control (if not already done)
3. Shift-leftclick on the plot in one of the columns to enter the corresponding x-value.



4. A small dialog will appear where you should enter the x-value to use (in this case, 10)

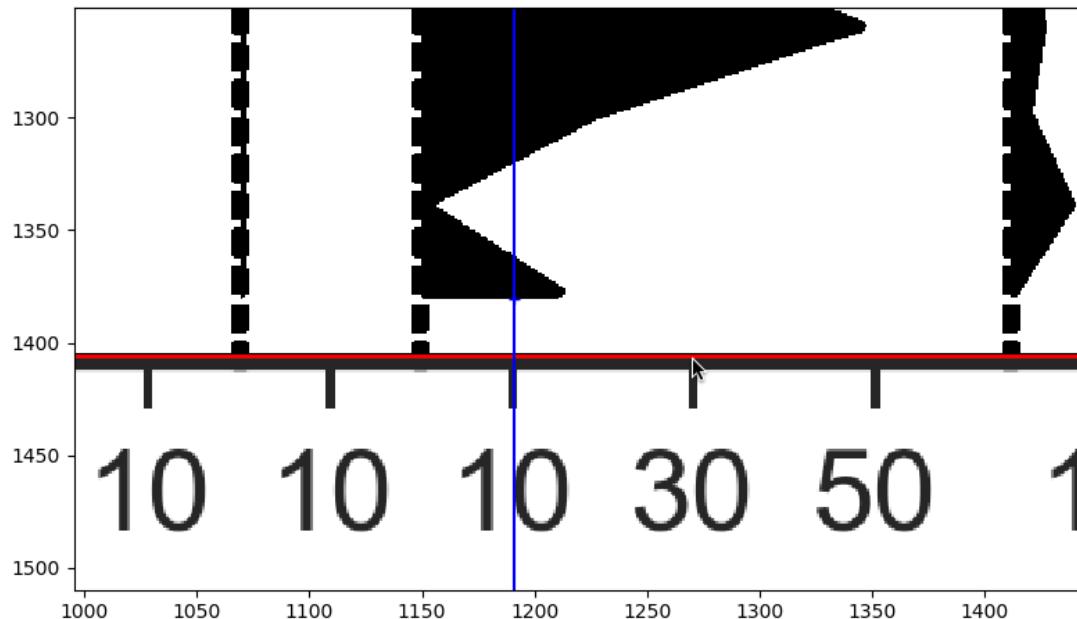


5. After hitting the *Ok* button, you will see a mark on the plot (blue line). You can select the mark via leftclick and drag it to a different location or you can delete it via rightclick.

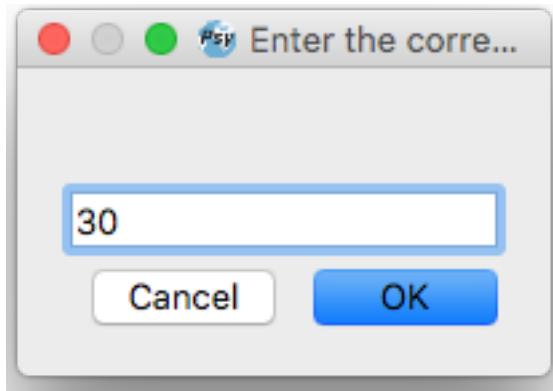


6. now repeat steps 3-5 on a second point in the same column

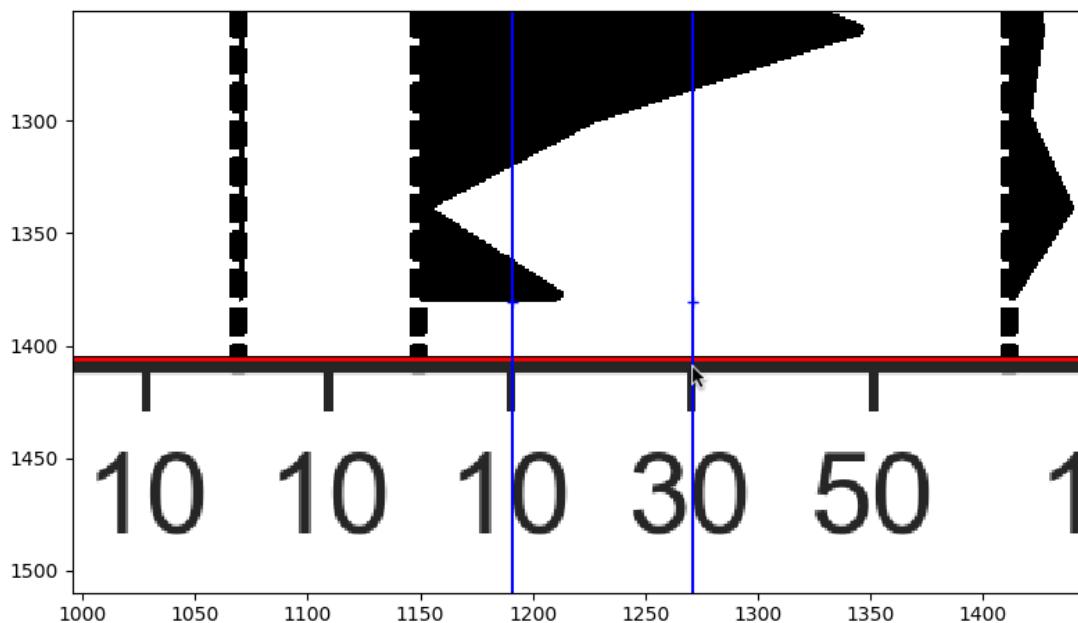
- Select another point



- Enter the corresponding value (here 30)



- A new mark is created that you can modify



7. Click the *Apply* button at the bottom of the straditzer control when you are done.

Note: If you have different units or different scalings in the diagram, create *column specific readers* and translate the x-axis separately for each reader/column.

Note: If you drag a mark and hold the *Shift* button while releasing the mouse button, the dialog in point 3 from above will not pop up.

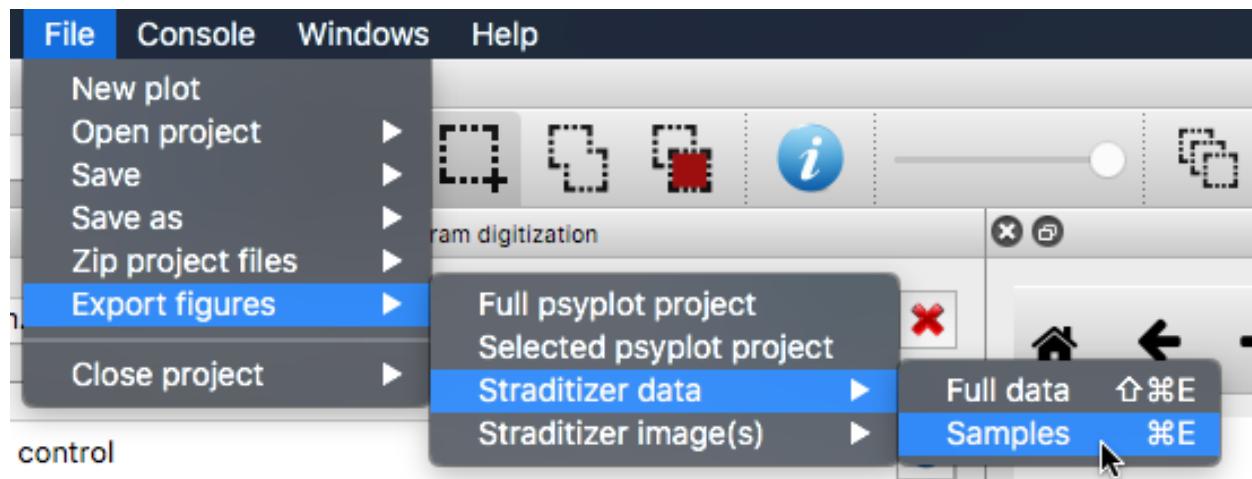
Export your results

Once you found the samples and added the y-axis and x-axis transformations, you can export them to a comma-separated (CSV) or an Excel file.

Just use the export menu via

File → Export figures → Straditzer data → Samples

or hit Control-E (Command-E on MacOS)



If you want to export the full digitized data, i.e. not only the samples, use

File → Export figures → Straditizer data → Full data

or hit Shift-Control-E (Shift-Command-E on MacOS)

1.6.3 Straditize helpers

Straditize has several generic tools to help you with various aspects.

The selection toolbar

The tools in this toolbar provides you the possibility to select features in the diagram. By default, the selected features will then be removed when you click the *Apply* or : guilabel:*Remove* button at the bottom of the straditizer window. The processing of the selection however varies depending on what task you are doing at the moment (for exaggerations, the selected features are for example marked as exaggerations).

The selection toolbar can *select from different sources*, provides *different mouse tools*, can be used in *different modes* provides *additional automatic tools for the selection*

Choosing the selection source

Using the combo box, you can also choose, where the features should be selected.

Straditizer If you choose the *Straditizer*, features will only be selected in the full image and the diagram part is untouched.

Reader If you choose the *Reader*, features will only be selected in the diagram image and the original image will be kept untouched

Reader - Greyscale The same as *Reader*, but whether two parts are connected or not is based on their color, not only on the binary image (see the tool below).

Selection tools

Too facilitate the selection for you, we implemented several tools to select features in the image:

select a rectangle or point With this tool you can select everything within a rectangle or a single point

select a polygon This tool can be accessed through the context menu of the tool (click and hold this button to open the menu). If activated and you click on the image, hold the mouse button and drag it around the features you want to select. Everything that is in the shape you draw while pressing the mouse button will be selected.

select based on connectivity or functionality This tool selects entire features in the image. The exact behaviour depends on what you are doing at the moment. If you hold and drag the mouse, you can select all features within a rectangle

In most cases, the tool just selects a feature based on the connectivity. If the selection source is the *Straditizer* or *Reader - Greyscale*, this connectivity is based on the greyscale image. For the *Reader* selection source, this is based on the binary image.

select based on color This tool can be accessed through the context menu of the tool (click and hold this button to open the menu). If activated and you click on the image, all connected cells that have the same color are selected. You can also relax this a bit, such that all colors that are close to the selected color will be selected. Furthermore, you can choose the select the colors in the *whole plot*, i.e. all pixels that have the same (or similar) color as the selected cell will be selected.

select the entire pixel row This tool also can be accessed through the context menu of the tool (click and hold this button to open the menu). If you click on the image, all selectable features on this horizontal level (i.e. on this pixel row in the image) are selected.

select the entire pixel column The same as the tool but for the vertical columns.

Selection mode

What exactly happens when you use the *selection tools described above* depends on the selection mode that you are in. This can be one of

new selection When you select something, it will create a new selection

add to selection Any new selection will be added to the current selection

remove from selection Any new selection will be removed from the current selection

Automatic tools

The selection toolbar also provides you some automatic tools to facilitate the selection for you

select all Everything that can be selected will be selected

expand selection Every selected pixel will be expanded based on its connectivity (see the tool above)

invert selection Everything that is selected will be unselected and everything that was not previously selected will be selected

clear selection Everything will be unselected

select everything to the right For each selected pixel in a column of the diagram part, we also select everything that is to the right of this pixel

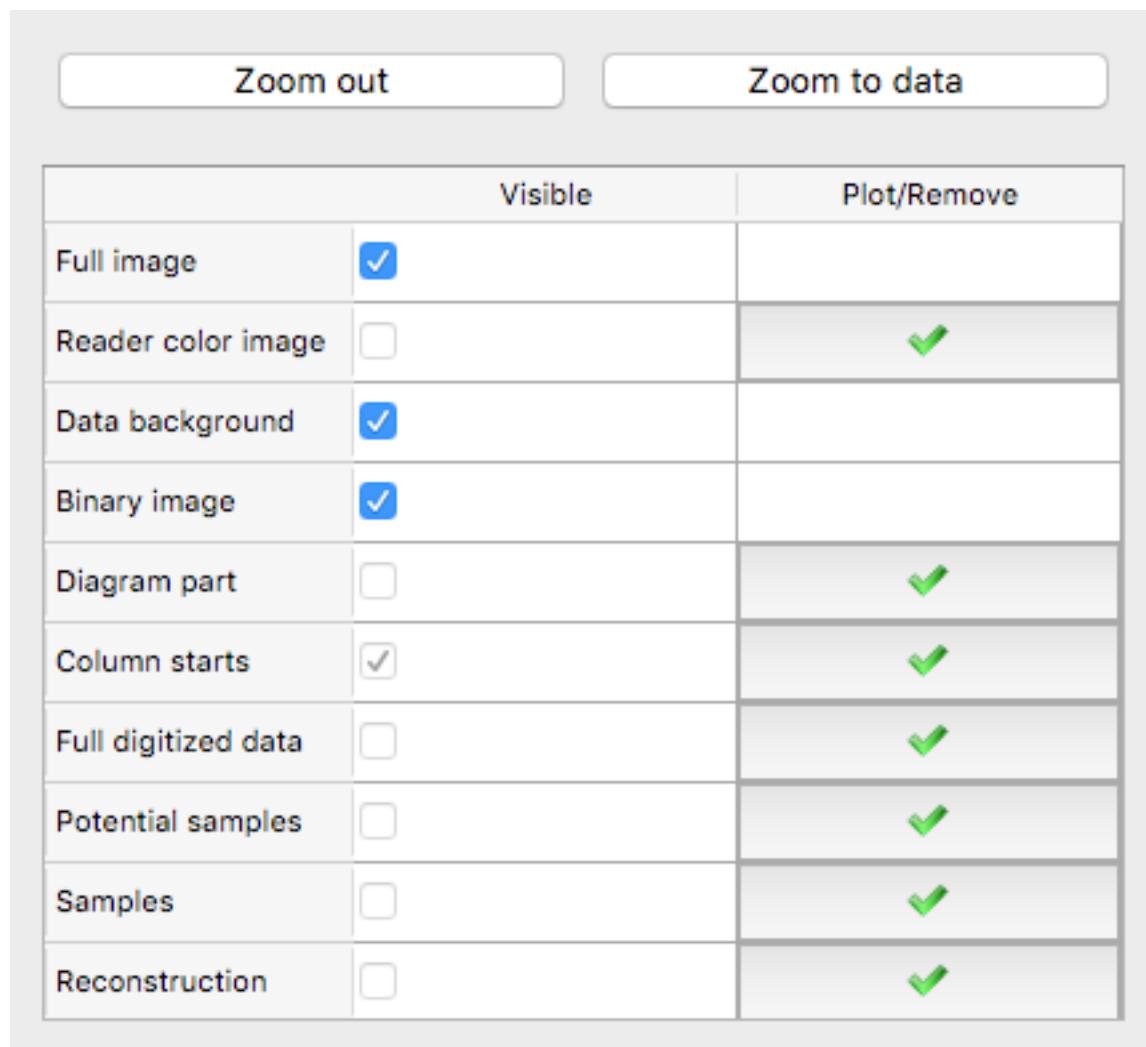
select based on a template  This method uses the `skimage.feature.match_template()` function to find a template within the current selection. This will then be selected or removed from the current selection based on the [current mode](#).

Visualize your progress with the plot control

To explore and validate your digitization, straditize implements multiple ways to visualize the results. You can show and remove these static visualizations using the items in the plot control.

Here you can hide the several plot objects using the *Visible* checkbox or you can plot the objects by clicking the  button or remove it by clicking the  button.

Note: Please note, that these are static plots. So if you change your data, e.g. through modifying the column starts, you have to remove () and plot () the *Column starts* item to visualize the changes.



	Visible	Plot/Remove
Full image	<input checked="" type="checkbox"/>	
Reader color image	<input type="checkbox"/>	
Data background	<input checked="" type="checkbox"/>	
Binary image	<input checked="" type="checkbox"/>	
Diagram part	<input type="checkbox"/>	
Column starts	<input checked="" type="checkbox"/>	
Full digitized data	<input type="checkbox"/>	
Potential samples	<input type="checkbox"/>	
Samples	<input type="checkbox"/>	
Reconstruction	<input type="checkbox"/>	

Some of the plotting features only visualize the currently selected columns. So if you have [column-specific readers](#) or [exaggerations](#), these options hide and show the plots for the currently selected columns only.

Full image: The full image is the diagram, that you are digitizing

Reader color image: This is the diagram part of the currently selected columns in color.

Data background: This is just a white layer that lies over the *Full image* in the diagram part. If you want to see the diagram part in the *Full image*, you have to hide this layer

Binary image: This is the image that is used to digitize the diagram part. As with the *Reader color image*, this option shows/hides the binary image for the currently selected columns.

Diagram part: This draws a red rectangle around the *diagram part* (see the *Basics and Terminology*)

Column starts: Red lines showing you, where each column starts of out of the currently selected columns starts

Full digitized data: After having hit the *Digitize* button, you can visualize the full digitized data for the currently selected columns.

Potential samples: For each column, we identify rough locations where samples may lie (see *Automatic samples identification*). This plot option lets you visualize these.

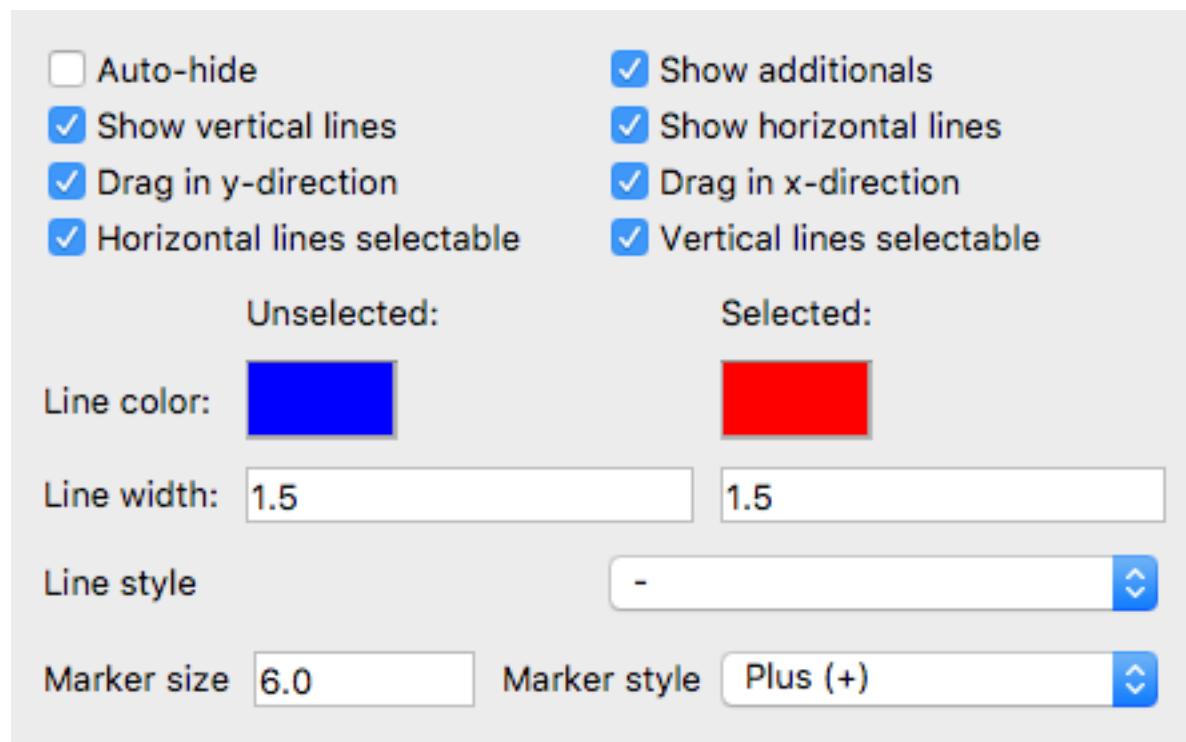
Samples: This option draws horizontal lines to show, the sample locations

Reconstruction: This plot option draws one line for each column, as the *Full digitized data* does, but it uses only the x- and y-values at the sample locations.

Configuring the appearance of markers

In the digitization, you will work with markers to select parts of your image. These can be the marks to *select the diagram part*, to *separate the columns* or to *edit your samples*.

You can modify the appearance of the marks using the *Marker control* section in the straditization control. This helps you to better visualize and select different parts in the diagram.



The functionalities are

Auto-hide Hide the lines of the marks, when they are not selected

Show additionals When *editing the samples*, for example, you will also see the rough locations of the samples for each column (little grey crosses). With the *Show additional*s setting, you can hide or show them

Show vertical lines and Show horizontal lines If your marks are crosses, e.g. when you are *selecting the diagram part*, you can hide and show the vertical or horizontal lines with these settings

Drag in y-direction and Drag in x-direction If unchecked, a mark cannot be moved along the corresponding axes.

You can use this option, if you want to change the y-value of a mark without changing the x-value (or the other way around)

Horizontal lines selectable and Vertical lines selectable If you uncheck one of these, you can prevent selecting a mark from the horizontal or vertical position. By default, you can select and move/drag a mark, by clicking on either of the lines (horizontal or vertical). But, for example, when editing the samples, many marks may lie on the same vertical position. Therefore it can be hard to move the mark of interest if you're selecting the vertical line of the mark.

Line color This changes the color of all marks.

Line width This changes the line width of the marks. Sometimes it is better, if the linewidth increases when you are selecting a line

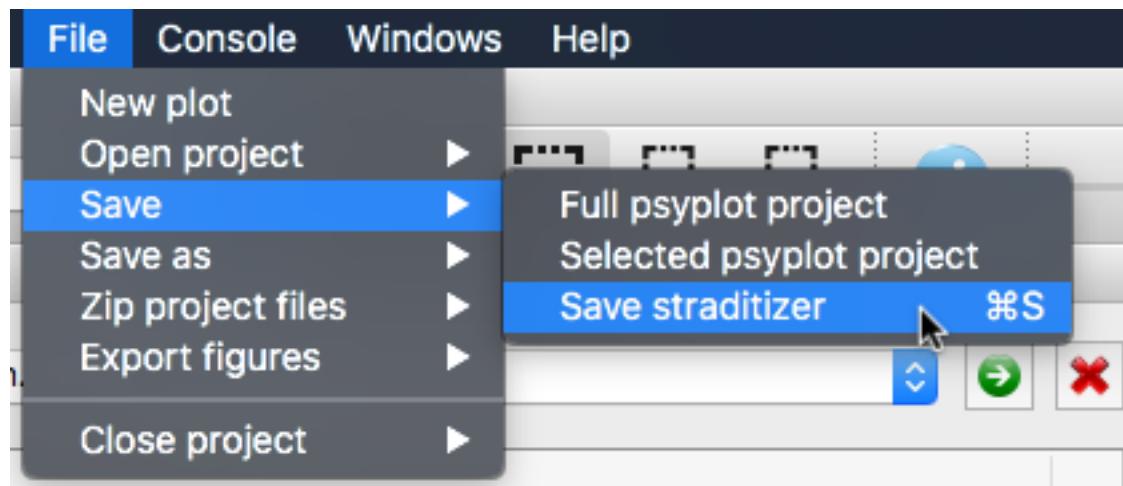
Line style You can change the style of the lines, to a dotted line, dashed line, etc.

Marker size and Marker style The location of the marks, i.e. where vertical and horizontal lines intersect, e.g. on corner of the diagram when *selecting the diagram part*, has a marker associated with it. It shows you, where the corner of your diagram is, where the sample value is, etc.. To make this more visible, you can change the size and the appearance of the mark.

Saving and loading your project

Saving a project

You can always save your current state to your hard-disc and continue working at it at a later point. To save a project, click *File* → *Save* → *Save straditzer*



or hit Control-S (Command-S on MacOS).

This will save your project, or, if you never saved it before or used the *Save As* option, it will open a file dialog to let you select the target location.

The default and recommended way is to save it as a [netCDF](#) file (ending with .nc or .nc4). This is a common self-describing, machine-independent data format that supports the creation, access, and sharing of array-oriented scientific

data. We recommend this format, because it works independent of the python libraries that you are using and let's you easily share the project with others or copy it to another machine.

The other possibility is to save it as a pickle (.pkl) file which uses the built-in `pickle` module to save the straditizer to a file. The reason, why we generally do not recommend this, is because the saved file might not work anymore if you upgrade your libraries (e.g. numpy, pandas or pillow).

Loading a project

Reloading the project works the same way as you open a new image file (see [Load a diagram](#)). Just go to *File* → *Open project* → *Open straditizer* → *Project or image* and choose the project file to load (or use the  button at the top of the straditizer control). If the chosen file ends with .nc, .nc4 or .pkl, straditize will recreate the given project.

1.7 API Reference

Digitizing stratigraphic diagrams with straditize

The core of straditize is the `Straditizer` class and the reader for the diagram part, the `DataReader`. Both classes work completely independent of the graphical user interface.

The graphical user interface is implemented as a plugin into the Graphical User Interface for the psyplot package package and is implemented in the `straditize.widgets` subpackage.

Authors

The code and GUI of straditize was developed by Philipp S. Sommer at the Institute of Earth System Dynamics (IDYST) at the University of Lausanne as part of the SNF funded HORNET Project (200021_169598).

The other contributors are Basil A. S. Davis, Manuel Chevalier and Dilan Rech who made significant contributions to the layout, workflow, beta tests and and reviewing of the software.

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1.7.1 Subpackages

`straditize.widgets` package

The straditizer widgets

This module contains widgets to digitize the straditizer diagrams through a GUI

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Classes

<i>EnableButton</i>	A <i>QPushButton</i> that emits a signal when enabled
<i>InfoButton</i> (parent[, fname, rst, name])	A button to display help informations in the help explorer
<i>StraditizerControlBase</i>	A base class for the straditizer widget
<i>StraditizerWidgets</i> (*args, **kwargs)	A widget that contains widgets to control the straditization in a GUI

Functions

<i>get_doc_file</i> (fname)	Return the path to a documentation file
<i>get_icon</i> (fname)	Return the path of an icon
<i>get_straditizer_widgets</i> ([mainwindow])	Get the <i>StraditizerWidgets</i> from the psyplot GUI mainwindow
<i>read_doc_file</i> (fname)	Return the content of a rst documentation file

class straditize.widgets.EnableButton

Bases: PyQt5.QtWidgets.QPushButton

A *QPushButton* that emits a signal when enabled **Attributes**

<i>enabled</i> (*args, **kwargs)	A signal that is emitted with a boolean whether if the button is
----------------------------------	--

Methods

<i>setEnabled</i> (b)	Reimplemented to emit the <i>enabled</i> signal
-----------------------	---

enabled(*args, **kwargs)

A signal that is emitted with a boolean whether if the button is enabled or disabled

setEnabled(b)

Reimplemented to emit the *enabled* signal

class straditize.widgets.InfoButton(parent, fname=None, rst=None, name=None)

Bases: PyQt5.QtWidgets.QToolButton

A button to display help informations in the help explorer

Parameters

- **parent** (*QWidget*) – The parent widget
- **fname** (*str*) – The name of the rst file. If None, specify the *rst* directly

- **rst** (*str*) – The restructured text to render when this button is clicked. If None, the *fname* has to be provided
- **name** (*str*) – The name to use for the document in the help_explorer

Methods

<code>show_docs()</code>	Show the docs
<code>show_docs()</code>	Show the docs
	Shows the docs in the in the <code>help_explorer</code>
class straditize.widgets. StraditizerControlBase	
Bases: <code>object</code>	
A base class for the straditizer widget	Methods
<code>add_info_button(child[, fname, rst, name, connections=[]])</code>	Add an info button to the given QTreeWidgetItem
<code>connect2apply(*funcs)</code>	Connect the functions to the <code>apply_button</code>
<code>connect2cancel(*funcs)</code>	Connect the functions to the <code>cancel_button</code>
<code>enable_or_disable_widgets(b)</code>	Enable or disable the widgets in this control
<code>init_straditizercontrol(straditizer_widgets)</code>	Initialize the straditizer control widget
<code>refresh()</code>	Refresh from the straditizer
<code>setup_children(item)</code>	Setup the children for this control
<code>should_be_enabled(w)</code>	Check if a widget should be enabled

Attributes

<code>apply_button</code>	The apply button of the straditizer_widgets
<code>cancel_button</code>	The cancel button of the straditizer_widgets
<code>help_explorer</code>	The <code>psyplot_gui.help_explorer.HelpExplorer</code> of the
<code>straditizer</code>	The current straditizer from the straditizer_widgets
<code>straditizer_widgets</code>	The StraditzerWidgets control
<code>widgets2disable</code>	A list of widgets to disable or enable if the apply_button is enabled

add_info_button (*child*, *fname=None*, *rst=None*, *name=None*, *connections=[]*)
Add an info button to the given QTreeWidgetItem

Parameters `child` (*QTreeWidgetItem*) – The item to which to add the infobutton

property apply_button
The apply button of the straditizer_widgets

property cancel_button
The cancel button of the straditizer_widgets

connect2apply (**funcs*)
Connect the functions to the `apply_button`

Parameters **funcs* – The callables that should be connected to the `apply_button`

```
connect2cancel (*funcs)
    Connect the functions to the cancel_button

    Parameters *funcs – The callables that should be connected to the cancel_button

enable_or_disable_widgets (b)
    Enable or disable the widgets in this control

    This method enables or disables the widgets2disable if the should_be_enabled() method
    evaluates to True

    Parameters b (bool) – If True, enable the widgets, if False, disable them

property help_explorer
    The psyplot_gui.help_explorer.HelpExplorer of the psyplot_gui.main.
    mainwindow

init_straditzercontrol (straditzer_widgets, item=None)
    Initialize the straditzer control widget

    This method should be called by every subclass when initializing. It sets the straditzer_widgets,
    connects the enable_or_disable_widgets() method and adds a new item to the
    StraditzerWidgets.tree

    Parameters
        • straditzer_widgets (StraditzerWidgets) – The main widget for the stra-
            ditizer GUI
        • item (QTreeWidgetItem) – The parent item in the StraditzerWidgets.
            tree. If given, the setup_children() is called with this item

refresh()
    Refresh from the straditzer

setup_children (item)
    Setup the children for this control

    This method is called to setup the children in the StraditzerWidgets.tree. By default, it just
    creates a child QTreeWidgetItem and sets this control as its widget

    Parameters item (QTreeWidgetItem) – The top level item in the
    StraditzerWidgets.tree

should_be_enabled (w)
    Check if a widget should be enabled

    This function checks if a given widget w from the widgets2disable attribute should be enabled or
    not

    Parameters w (QWidget) – The widget to check
    Returns True, if the widget should be enabled
    Return type bool

property straditzer
    The current straditzer from the straditzer_widgets

straditzer_widgets = None
    The StraditzerWidgets control

widgets2disable = []
    A list of widgets to disable or enable if the apply_button is enabled
```

```
class straditize.widgets.StraditizerWidgets(*args, **kwargs)
Bases: PyQt5.QtWidgets.QWidget, psyplot_gui.common.DockMixin
```

A widget that contains widgets to control the straditization in a GUI

This widget is the basis of the straditize GUI and implemented as a plugin into the psyplot gui. The open straditizers are handled in the `_straditizer` attribute.

The central parts of this widget are

- The combobox to manage the open straditizers
- The QTreeWidget in the `tree` attribute that contains all the controls to interface the straditizer
- the tutorial area
- the *Apply* and *Cancel* button

Methods

<code>add_info_button(child[, fname, rst, name, ...])</code>	Add an infobutton to the <code>tree</code> widget
<code>add_straditizer(stradi)</code>	Add a straditizer to the list of open straditzers
<code>autosave()</code>	Autosave the current straditzer
<code>close_all_straditzers()</code>	Close all straditzers
<code>close_straditzer()</code>	Close the current straditzer
<code>create_straditzer_from_args(fnames[, ...])</code>	Create a straditzer from the given file name
<code>disable_apply_button()</code>	Method that is called when the <code>cancel_button</code> is clicked
<code>edit_attrs()</code>	Edit the attributes of the current straditzer
<code>get_attr(stradi, attr)</code>	
<code>raise_figures()</code>	Raise the figures of the current straditzer in the GUI
<code>refresh()</code>	Refresh from the straditzer
<code>reload_autosaved()</code>	Reload the autosaved straditzer and close the old one
<code>reset_control()</code>	Reset the GUI of straditize
<code>set_current_stradi(i)</code>	Set the i-th straditzer to the current one
<code>show_or_hide_toolbar()</code>	Show or hide the toolbar depending on the visibility of this widget
<code>start_tutorial(state[, tutorial_cls])</code>	Start or stop the tutorial
<code>switch_to_straditzer_layout()</code>	Switch to the straditzer layout
<code>to_dock(main, *args, **kwargs)</code>	

Attributes

<code>always_yes</code>	Boolean that is True if all dialogs should be answered with <i>Yes</i>
<code>apply_button</code>	The apply button
<code>attrs_button</code>	The button to edit the straditzer attributes
<code>autosaved</code>	Auto-saved straditzers
<code>axes_translations</code>	The <code>straditize.widgets.axes_translations.AxesTranslations</code> to
<code>btn_close_stradi</code>	A button to close the current straditzer

Continued on next page

Table 9 – continued from previous page

<code>btn_open_stradi</code>	A button to open a new straditizer
<code>btn_reload_autosaved</code>	A button to reload the last autosaved state
<code>cancel_button</code>	The cancel button
<code>colnames_manager</code>	The <code>straditize.widgets.colnames.ColumnNamesManager</code> to interface
<code>digitizer</code>	The <code>straditize.widgets.data.DigitizingControl</code> to interface
<code>hidden</code>	<code>bool(x) -> bool</code>
<code>image_rescaler</code>	The <code>straditize.widgets.image_correction.ImageRescaler</code> class to
<code>image_rotator</code>	The <code>straditize.widgets.image_correction.ImageRotator</code> class to
<code>info_button</code>	An <code>InfoButton</code> to display the docs
<code>marker_control</code>	The <code>straditize.widgets.marker_control.MarkerControl</code> to modify
<code>open_external(*args, **kwargs)</code>	
<code>plot_control</code>	The <code>straditize.widgets.plots.PlotControl</code> to display additional
<code>progress_widget</code>	The <code>straditize.widgets.progress_widget.ProgressWidget</code> to
<code>selection_toolbar</code>	The <code>straditize.widgets.selection_toolbar.SelectionToolbar</code> to
<code>stradi_combo</code>	A QComboBox to select the current straditizer
<code>straditizer</code>	The <code>straditize.straditizer.Straditizer</code> instance
<code>title</code>	<code>str(object=”) -> str</code>
<code>tree</code>	The QTreeWidget that contains the different widgets for the digitization
<code>tutorial</code>	The <code>straditize.widgets.tutorial.Tutorial</code> class
<code>tutorial_button</code>	The button to start a tutorial

`add_info_button` (`child, fname=None, rst=None, name=None, connections=[]`)

Add an infobutton to the `tree` widget

Parameters

- **`child`** (`QTreeWidgetItem`) – The item to which to add the infobutton
- **`connections`** (`list of QPushButtons`) – Buttons that should be clicked when the info button is clicked

`add_straditizer` (`stradi`)

Add a straditizer to the list of open straditzers

`always_yes = False`

Boolean that is True if all dialogs should be answered with Yes

`apply_button = None`

The apply button

attrs_button = None
The button to edit the straditizer attributes

autosave()
Autosave the current straditizer

autosaved = []
Auto-saved straditzers

axes_translations = None
The `straditize.widgets.axes_translations.AxesTranslations` to handle the y- and x-axis conversions

btn_close_stradi = None
A button to close the current straditizer

btn_open_stradi = None
A button to open a new straditizer

btn_reload_autosaved = None
A button to reload the last autosaved state

cancel_button = None
The cancel button

close_all_straditzers()
Close all straditzers

close_straditizer()
Close the current straditizer

colnames_manager = None
The `straditize.widgets.colnames.ColumnNamesManager` to interface the `:straditize.straditizer.Straditizer.colnames_reader`

create_straditizer_from_args(fnames, project=None, xlim=None, ylim=None, full=False, reader_type='area')
Create a straditizer from the given file name

This method is called when the `psyplot_gui.mainmainwindow` receives a ‘straditize’ callback

digitizer = None
The `straditize.widgets.data.DigitizingControl` to interface the `:straditize.straditizer.Straditizer.data_reader`

disable_apply_button()
Method that is called when the `cancel_button` is clicked

dock_position = 1

edit_attrs()
Edit the attributes of the current straditizer

This creates a new dataframe editor to edit the `straditize.straditizer.Straditizer.attrs` meta informations

get_attr(stradi, attr)

hidden = True

image_rescaler = None
The `straditize.widgets.image_correction.ImageRescaler` class to rescale the image

image_rotator = None
The `straditize.widgets.image_correction.ImageRotator` class to rotate the image

```
info_button = None
    An InfoButton to display the docs

marker_control = None
    The straditize.widgets.marker_control.MarkerControl to modify the appearance of
    the marks of the current straditizer

open_external(*args, **kwargs)

plot_control = None
    The straditize.widgets.plots.PlotControl to display additional information on the dia-
    gram

progress_widget = None
    The straditize.widgets.progress_widget.ProgressWidget to display the progress of
    the straditization

raise_figures()
    Raise the figures of the current straditizer in the GUI

refresh()
    Refresh from the straditizer

reload_autosaved()
    Reload the autosaved straditizer and close the old one

reset_control()
    Reset the GUI of straditize

selection_toolbar = None
    The straditize.widgets.selection_toolbar.SelectionToolbar to select features in
    the stratigraphic diagram

set_current_stradi(i)
    Set the i-th straditizer to the current one

show_or_hide_toolbar()
    Show or hide the toolbar depending on the visibility of this widget

start_tutorial(state, tutorial_cls=None)
    Start or stop the tutorial

    Parameters
        • state (bool) – If False, the tutorial is stopped. Otherwise it is started
        • tutorial_cls (straditize.widgets.tutorial.beginner.Tutorial) –
            The tutorial class to use. If None, it will be asked in a QInputDialog

stradi_combo = None
    A QComboBox to select the current straditizer

straditizer = None
    The straditize.straditizer.Straditizer instance

switch_to_straditizer_layout()
    Switch to the straditizer layout

    This method makes this widget visible and stacks it with the psyplot content widget

title = 'Stratigraphic diagram digitization'

to_dock(main, *args, **kwargs)
```

```
tree = None
The QTreeWidget that contains the different widgets for the digitization

tutorial = None
The straditize.widgets.tutorial.Tutorial class

tutorial_button = None
The button to start a tutorial

window_layout_action = None

straditize.widgets.get_doc_file(fname)
Return the path to a documentation file

straditize.widgets.get_icon(fname)
Return the path of an icon

straditize.widgets.get_straditizer_widgets(mainwindow=None)
Get the StraditzerWidgets from the psyplot GUI mainwindow

    Parameters psyplot_gui.main.MainWindow – The mainwindow to use. If None, the
    psyplot_gui.main.mainwindow is used.

    Returns The straditizer widgets of the given mainwindow

    Return type StraditzerWidgets

straditize.widgets.read_doc_file(fname)
Return the content of a rst documentation file
```

Subpackages

straditize.widgets.tutorial package

Tutorials for straditize

The *beginner* tutorial serves as a first look into the software. The *hoya_del_castillo* tutorial on the other hand is a true pollen diagram with more than 20 taxa. Both tutorials can be started from the GUI through the *Tutorial* button.

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Submodules

straditize.widgets.tutorial.beginner module

The tutorial of straditize

This module contains a guided tour to get started with straditize

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Classes

<i>CleanImagePage</i> (filename, tutorial)	Tutorial page to clean the diagram part
<i>ColumnNames</i> (filename, tutorial)	The page for recognizing column names
<i>ControlIntro</i> (filename, tutorial)	Tutorial page for the control
<i>CreateReader</i> (filename, tutorial)	The page for creating the reader
<i>DigitizePage</i> (filename, tutorial)	The page for digitizing the diagram
<i>FinishPage</i> (filename, tutorial)	The last page of the tutorial
<i>LoadImage</i> (filename, tutorial)	TutorialPage for loading the straditzer image
<i>SamplesPage</i> (filename, tutorial)	The page for finding and editing the samples
<i>SelectDataPart</i> (filename, tutorial)	TutorialPage for selecting the data part
<i>SeparateColumns</i> (filename, tutorial)	The page for separating the columns
<i>TranslateXAxis</i> (filename, tutorial)	The tutorial page for translating the y-axis
<i>TranslateYAxis</i> (filename, tutorial)	The tutorial page for translating the y-axis
<i>Tutorial</i> (straditizer_widgets)	A tutorial for digitizing an area diagram
<i>TutorialDocs</i> (*args, **kwargs)	A documentation viewer for the tutorial docs
<i>TutorialNavigation</i> (nsteps, validate, *args, ...)	A widget for navigating through the tutorial.
<i>TutorialPage</i> (filename, tutorial)	A base class for the tutorial pages

class straditize.widgets.tutorial.beginner.**CleanImagePage** (filename, tutorial)

Bases: straditize.widgets.tutorial.beginner.TutorialPage

Tutorial page to clean the diagram part

Parameters

- **filename** (*str*) – The basename (without ending) of the RST file corresponding to this tutorial page
- **tutorial** (*Tutorial*) – The tutorial instance

Methods

<i>activate()</i>	Method that is called, when the page is activated
<i>clicked_btn_remove_xaxes()</i>	
<i>clicked_btn_remove_yaxes()</i>	

Continued on next page

Table 11 – continued from previous page

<code>deactivate()</code>	Method that is called, when the page is deactivated
<code>hint()</code>	A method that should display a hint to the user
<code>icon_to_bytes(icon)</code>	
<code>skip()</code>	Skip the steps in this page

Attributes

<code>btn_remove_xaxes_clicked</code>	<code>bool(x) -> bool</code>
<code>btn_remove_yaxes_clicked</code>	<code>bool(x) -> bool</code>
<code>is_finished</code>	Boolean that is True, if the steps are all finished

`activate()`

Method that is called, when the page is activated

`btn_remove_xaxes_clicked = False`

`btn_remove_yaxes_clicked = False`

`clicked_btn_remove_xaxes()`

`clicked_btn_remove_yaxes()`

`deactivate()`

Method that is called, when the page is deactivated

`hint()`

A method that should display a hint to the user

`icon_to_bytes(icon)`

`property is_finished`

Boolean that is True, if the steps are all finished

`skip()`

Skip the steps in this page

class straditize.widgets.tutorial.beginner.`ColumnNames` (*filename, tutorial*)

Bases: `straditize.widgets.tutorial.TutorialPage`

The page for recognizing column names

Parameters

- `filename (str)` – The basename (without ending) of the RST file corresponding to this tutorial page
- `tutorial (Tutorial)` – The tutorial instance

Methods

<code>activate()</code>	Method that is called, when the page is activated
<code>clicked_select_names_button()</code>	
<code>deactivate()</code>	Method that is called, when the page is deactivated
<code>hint()</code>	A method that should display a hint to the user
<code>hint_for_start_editing()</code>	
<code>hint_for_wrong_name(col, curr, ref)</code>	Display a hint if a name is not correctly set
<code>skip()</code>	Skip the steps in this page

Attributes

<code>column_names</code>	The column names in the diagram
<code>is_finished</code>	Boolean that is True, if the steps are all finished
<code>select_names_button_clicked</code>	<code>bool(x) -> bool</code>

`activate()`

Method that is called, when the page is activated

`clicked_select_names_button()`

```
column_names = ['Pinus', 'Juniperus', 'Quercus ilex-type', 'Chenopodiaceae']
The column names in the diagram
```

`deactivate()`

Method that is called, when the page is deactivated

`hint()`

A method that should display a hint to the user

`hint_for_start_editing()`

```
hint_for_wrong_name(col, curr, ref)
Display a hint if a name is not correctly set
```

`property is_finished`

Boolean that is True, if the steps are all finished

`select_names_button_clicked = False`

`skip()`

Skip the steps in this page

class straditize.widgets.tutorial.beginner.**ControlIntro** (*filename, tutorial*)

Bases: *straditize.widgets.tutorial.beginner.TutorialPage*

Tutorial page for the control

Parameters

- **filename** (*str*) – The basename (without ending) of the RST file corresponding to this tutorial page
- **tutorial** (*Tutorial*) – The tutorial instance

Methods

<code>activate()</code>	Method that is called, when the page is activated
-------------------------	---

`activate()`

Method that is called, when the page is activated

class straditize.widgets.tutorial.beginner.**CreateReader** (*filename, tutorial*)

Bases: *straditize.widgets.tutorial.beginner.TutorialPage*

The page for creating the reader

Parameters

- **filename** (*str*) – The basename (without ending) of the RST file corresponding to this tutorial page
- **tutorial** (*Tutorial*) – The tutorial instance

Methods

<code>hint()</code>	A method that should display a hint to the user
<code>skip()</code>	Skip the steps in this page

Attributes

<code>is_finished</code>	Boolean that is True, if the steps are all finished
--------------------------	---

hint()

A method that should display a hint to the user

property is_finished

Boolean that is True, if the steps are all finished

skip()

Skip the steps in this page

class straditize.widgets.tutorial.beginner.**DigitizePage** (*filename, tutorial*)
Bases: *straditize.widgets.tutorial.TutorialPage*

The page for digitizing the diagram

Parameters

- **filename** (*str*) – The basename (without ending) of the RST file corresponding to this tutorial page
- **tutorial** (*Tutorial*) – The tutorial instance

Methods

<code>hint()</code>	A method that should display a hint to the user
<code>skip()</code>	Skip the steps in this page

Attributes

<code>is_finished</code>	Boolean that is True, if the steps are all finished
--------------------------	---

hint()

A method that should display a hint to the user

property is_finished

Boolean that is True, if the steps are all finished

skip()

Skip the steps in this page

class straditize.widgets.tutorial.beginner.**FinishPage** (*filename, tutorial*)
Bases: *straditize.widgets.tutorial.TutorialPage*

The last page of the tutorial

Parameters

- **filename** (*str*) – The basename (without ending) of the RST file corresponding to this tutorial page
- **tutorial** (*Tutorial*) – The tutorial instance

Methods

<code>show()</code>	Reimplemented to release the help explorer lock
---------------------	---

show()
Reimplemented to release the help explorer lock

class straditize.widgets.tutorial.beginner.**LoadImage** (*filename, tutorial*)

Bases: *straditize.widgets.tutorial.beginner.TutorialPage*

TutorialPage for loading the straditizer image

Parameters

- **filename** (*str*) – The basename (without ending) of the RST file corresponding to this tutorial page
- **tutorial** (*Tutorial*) – The tutorial instance

Methods

<code>activate()</code>	Method that is called, when the page is activated
<code>deactivate()</code>	Method that is called, when the page is deactivated
<code>hint()</code>	A method that should display a hint to the user
<code>skip()</code>	Skip the steps in this page

Attributes

<code>is_finished</code>	Boolean that is True, if the steps are all finished
--------------------------	---

activate()
Method that is called, when the page is activated

deactivate()
Method that is called, when the page is deactivated

hint()
A method that should display a hint to the user

property is_finished
Boolean that is True, if the steps are all finished

skip()
Skip the steps in this page

class straditize.widgets.tutorial.beginner.**SamplesPage** (*filename, tutorial*)

Bases: *straditize.widgets.tutorial.beginner.TutorialPage*

The page for finding and editing the samples

Parameters

- **filename** (*str*) – The basename (without ending) of the RST file corresponding to this tutorial page
- **tutorial** (*Tutorial*) – The tutorial instance

Methods

<code>activate()</code>	Method that is called, when the page is activated
	Continued on next page

Table 23 – continued from previous page

<code>clicked_correct_button()</code>	
<code>deactivate()</code>	Method that is called, when the page is deactivated
<code>hint()</code>	A method that should display a hint to the user
<code>skip()</code>	Skip the steps in this page

Attributes

<code>correct_button_clicked</code>	bool(x) -> bool
<code>is_finished</code>	Boolean that is True, if the steps are all finished

activate()

Method that is called, when the page is activated

clicked_correct_button()**correct_button_clicked = False****deactivate()**

Method that is called, when the page is deactivated

hint()

A method that should display a hint to the user

property is_finished

Boolean that is True, if the steps are all finished

skip()

Skip the steps in this page

class straditize.widgets.tutorial.beginner.SelectDataPart (filename, tutorial)

Bases: `straditize.widgets.tutorial.beginner.TutorialPage`

TutorialPage for selecting the data part

Parameters

- `filename (str)` – The basename (without ending) of the RST file corresponding to this tutorial page
- `tutorial (Tutorial)` – The tutorial instance

Methods

<code>activate()</code>	Method that is called, when the page is activated
<code>check_mark(mark)</code>	
<code>clicked_correct_button()</code>	
<code>deactivate()</code>	Method that is called, when the page is deactivated
<code>display_reference_marks()</code>	
<code>hint()</code>	A method that should display a hint to the user
<code>is_valid_x(x)</code>	
<code>is_valid_y(y)</code>	
<code>skip()</code>	Skip the steps in this page
<code>validate_corners()</code>	

Attributes

<code>correct_button_clicked</code>	bool(x) -> bool
<code>is_finished</code>	Boolean that is True, if the steps are all finished
<code>marks</code>	Built-in mutable sequence.
<code>ref_lims</code>	The reference x- and y- limits
<code>remove_data_box</code>	true if the data box should be removed at the end
<code>valid_xlims</code>	Valid ranges for xmin and xmax
<code>valid_ylims</code>	Valid ranges for ymin and ymax

activate()

Method that is called, when the page is activated

check_mark(mark)**clicked_correct_button()****correct_button_clicked = False****deactivate()**

Method that is called, when the page is deactivated

display_reference_marks()**hint()**

A method that should display a hint to the user

property is_finished

Boolean that is True, if the steps are all finished

is_valid_x(x)**is_valid_y(y)****marks = []****ref_lims = array([[258, 1803], [375, 1666]])**

The reference x- and y- limits

remove_data_box = True

true if the data box should be removed at the end

skip()

Skip the steps in this page

valid_xlims = array([[221, 263], [1730, 1922]])

Valid ranges for xmin and xmax

valid_ylims = array([[346, 403], [1648, 1701]])

Valid ranges for ymin and ymax

validate_corners()

class straditize.widgets.tutorial.beginner.**SeparateColumns** (*filename, tutorial*)

Bases: straditize.widgets.tutorial.TutorialPage

The page for separating the columns

Parameters

- **filename** (*str*) – The basename (without ending) of the RST file corresponding to this tutorial page
- **tutorial** (*Tutorial*) – The tutorial instance

Methods

<code>activate()</code>	Method that is called, when the page is activated
<code>clicked_correct_button()</code>	
<code>deactivate()</code>	Method that is called, when the page is deactivated
<code>hint()</code>	A method that should display a hint to the user
<code>skip()</code>	Skip the steps in this page

Attributes

<code>correct_button_clicked</code>	<code>bool(x) -> bool</code>
<code>is_finished</code>	Boolean that is True, if the steps are all finished
<code>ncols</code>	<code>int([x]) -> integer</code>

`activate()`

Method that is called, when the page is activated

`clicked_correct_button()`

`correct_button_clicked = False`

`deactivate()`

Method that is called, when the page is deactivated

`hint()`

A method that should display a hint to the user

`property is_finished`

Boolean that is True, if the steps are all finished

`ncols = 4`

`skip()`

Skip the steps in this page

class straditize.widgets.tutorial.beginner.**TranslateXAxis** (*filename, tutorial*)
Bases: straditize.widgets.tutorial.beginner.TutorialPage

The tutorial page for translating the y-axis

Parameters

- **filename** (`str`) – The basename (without ending) of the RST file corresponding to this tutorial page
- **tutorial** (`Tutorial`) – The tutorial instance

Methods

<code>activate()</code>	Method that is called, when the page is activated
<code>clicked_correct_button()</code>	
<code>deactivate()</code>	Method that is called, when the page is deactivated
<code>hint()</code>	A method that should display a hint to the user
<code>skip()</code>	Skip the steps in this page

Attributes

<code>correct_button_clicked</code>	bool(x) -> bool
<code>is_finished</code>	Boolean that is True, if the steps are all finished

`activate()`

Method that is called, when the page is activated

`clicked_correct_button()`

`correct_button_clicked = False`

`deactivate()`

Method that is called, when the page is deactivated

`hint()`

A method that should display a hint to the user

`property is_finished`

Boolean that is True, if the steps are all finished

`skip()`

Skip the steps in this page

class straditize.widgets.tutorial.beginner.**TranslateYAxis** (*filename, tutorial*)

Bases: *straditize.widgets.tutorial.beginner.TutorialPage*

The tutorial page for translating the y-axis

Parameters

- **filename** (*str*) – The basename (without ending) of the RST file corresponding to this tutorial page
- **tutorial** (*Tutorial*) – The tutorial instance

Methods

<code>activate()</code>	Method that is called, when the page is activated
<code>clicked_correct_button()</code>	
<code>deactivate()</code>	Method that is called, when the page is deactivated
<code>hint()</code>	A method that should display a hint to the user
<code>skip()</code>	Skip the steps in this page

Attributes

<code>correct_button_clicked</code>	bool(x) -> bool
<code>is_finished</code>	Boolean that is True, if the steps are all finished

`activate()`

Method that is called, when the page is activated

`clicked_correct_button()`

`correct_button_clicked = False`

`deactivate()`

Method that is called, when the page is deactivated

`hint()`

A method that should display a hint to the user

```
property is_finished
    Boolean that is True, if the steps are all finished

skip()
    Skip the steps in this page

class straditize.widgets.tutorial.beginner.Tutorial(straditizer_widgets)
    Bases: straditize.widgets.StraditzerControlBase, straditize.widgets.
        tutorial.beginner.TutorialPage

A tutorial for digitizing an area diagram Methods
```

<code>close()</code>	Close the tutorial and remove the widgets
<code>display_hint(i)</code>	Display the hint for a tutorial page
<code>get_doc_files()</code>	Get the rst files for the tutorial
<code>goto_page(old, new)</code>	Go to another page
<code>refresh()</code>	Refresh from the straditizer
<code>setup_tutorial_pages()</code>	Setup the pages attribute and initialize the tutorial pages
<code>show()</code>	Show the documentation of the tutorial
<code>skip_page(i)</code>	Skip a tutorial page
<code>validate_page(i[, silent])</code>	Validate a tutorial page

Attributes

<code>current_page</code>	The current page of the tutorial (corresponding to the <code>TutorialNavigation.current_step</code>)
<code>load_image_step</code>	The number of the page that loads the diagram image (i.e.
<code>navigation</code>	A <code>TutorialNavigation</code> to navigate through the tutorial
<code>pages</code>	A list of the <code>TutorialPages</code> for this tutorial
<code>tutorial_docs</code>	A <code>TutorialDocs</code> to display the RST-files of the tutorial

`close()`

Close the tutorial and remove the widgets

`property current_page`

The current page of the tutorial (corresponding to the `TutorialNavigation.current_step`)

`display_hint(i)`

Display the hint for a tutorial page

Parameters `i` (`int`) – The index of the page in the `pages` attribute

See also:

`TutorialPage.hint()`

`get_doc_files()`

Get the rst files for the tutorial

Returns

- `str` – The path to the tutorial introduction file
- `list of str` – The paths of the remaining tutorial files

goto_page (*old, new*)
Go to another page

Parameters

- **old** (*int*) – The index of the old page in the *pages* attribute that is subject to be deactivated (see *TutorialPage.deactivate()*)
- **new** (*int*) – The index of the new page in the *pages* attribute that is subject to be activated (see *TutorialPage.activate()*)

See also:

TutorialPage.activate(), *TutorialPage.deactivate()*

property load_image_step

The number of the page that loads the diagram image (i.e. the index of the *LoadImage* instance in the *pages* attribute)

navigation = None

A *TutorialNavigation* to navigate through the tutorial

pages = []

A list of the *TutorialPages* for this tutorial

refresh()

Refresh from the straditizer

setup_tutorial_pages()

Setup the *pages* attribute and initialize the tutorial pages

show()

Show the documentation of the tutorial

skip_page (*i*)

Skip a tutorial page

Parameters **i** (*int*) – The index of the page in the *pages* attribute

See also:

TutorialPage.skip()

tutorial_docs = None

A *TutorialDocs* to display the RST-files of the tutorial

validate_page (*i, silent=False*)

Validate a tutorial page

Parameters

- **i** (*int*) – The index of the page in the *pages* attribute
- **silent** (*bool*) – If True, and the page is not yet finished (see *TutorialPage.is_finished*), the hint is displayed

Returns True, if the page *is_finished*

Return type bool

class straditize.widgets.tutorial.beginner.**TutorialDocs** (*args, **kwargs)
Bases: *psyplot_gui.help_explorer.UrlHelp*, *psyplot_gui.common.DockMixin*

A documentation viewer for the tutorial docs

This viewer is accessible through the `Tutorial.tutorial_docs` attribute and shows the `src_file` for the tutorial `pages` Attributes

<code>title</code>	<code>str(object="") -> str</code>
<code>dock_position = 2</code>	
<code>title = 'Straditize tutorial'</code>	
<code>class straditize.widgets.tutorial.beginner.TutorialNavigation(nsteps, validate, *args, **kwargs)</code>	Bases: PyQt5.QtWidgets.QWidget

A widget for navigating through the tutorial. It has a button to go to the previous step and to go to the next step. Furthermore it has a progressbar implemented a `hint` button

Parameters

- `nsteps (int)` – The total number of steps in the `Tutorial`
- `validate (callable)` – A callable that takes the `current_step` as an argument and returns a `bool` whether the current step is valid and finished, or not

Attributes

<code>current_step</code>	The current step in the tutorial
<code>hint_requested(*args, **kwargs)</code>	Signal that is emitted if the hint of the current step is requested
<code>skipped(*args, **kwargs)</code>	Signal that is emitted, if the step is skipped
<code>step_changed(*args, **kwargs)</code>	Signal that is emitted, when the step changes.

Methods

<code>display_hint()</code>	Trigger the <code>hint_requested</code> signal with the current step
<code>goto_next_step()</code>	Increase the <code>current_step</code> by one
<code>goto_prev_step()</code>	Decrease the <code>current_step</code> by one
<code>maybe_enable_widgets()</code>	Enable the buttons based on the <code>current_step</code>
<code>setEnabled(enable)</code>	Enable or disable the navigation buttons
<code>set_current_step(i)</code>	Change the <code>current_step</code>
<code>show_info()</code>	Trigger the <code>step_changed</code> signal with the current step
<code>skip()</code>	Skip the <code>current_step</code> and emit the <code>skipped</code> signal

`current_step = 0`
The current step in the tutorial

`display_hint ()`
Trigger the `hint_requested` signal with the current step

`goto_next_step ()`
Increase the `current_step` by one

`goto_prev_step ()`
Decrease the `current_step` by one

`hint_requested (*args, **kwargs)`

Signal that is emitted if the hint of the current step is requested

maybe_enable_widgets()

Enable the buttons based on the *current_step*

setEnabled(enable)

Enable or disable the navigation buttons

Parameters **enable** (*bool*) – Whether to enable or disable the buttons

setCurrent_step(i)

Change the *current_step*

Parameters **i** (*int*) – The *current_step* to switch to

show_info()

Trigger the *step_changed* signal with the current step

skip()

Skip the *current_step* and emit the *skipped* signal

skipped(*args, **kwargs)

Signal that is emitted, if the step is skipped

step_changed(*args, **kwargs)

Signal that is emitted, when the step changes. The first integer is the old step, the second one the current step

class straditize.widgets.tutorial.beginner.TutorialPage (filename, tutorial)

Bases: *object*

A base class for the tutorial pages

Subclasses show implement the *show_hint()* method and the *is_finished()* property

Parameters

- **filename** (*str*) – The basename (without ending) of the RST file corresponding to this tutorial page
- **tutorial** (*Tutorial*) – The tutorial instance

Methods

<i>activate()</i>	Method that is called, when the page is activated
<i>deactivate()</i>	Method that is called, when the page is deactivated
<i>hint()</i>	A method that should display a hint to the user
<i>lock_viewer(lock)</i>	Set or unset the url lock of the HTML viewer
<i>refresh()</i>	
<i>show()</i>	Show the page and browse the <i>filename</i> in the tutorial docs
<i>show_hint()</i>	Show a hint to the user
<i>show_tooltip_at_widget</i> (tooltip, widget[, ...])	Show a tooltip close to a widget
<i>show_tooltip_in_plot</i> (tooltip, x, y[, ...])	Show a tooltip in the matplotlib figure at the given coordinates
<i>skip()</i>	Skip the steps in this page

Attributes

<code>is_finished</code>	Boolean that is True, if the steps are all finished
<code>is_selecting</code>	True if the user clicked the <code>btn_select_data</code> button
<code>src_base</code>	The basename of the stratigraphic diagram image for this tutorial
<code>src_dir</code>	The source directory for the docs
<code>src_file</code>	The complete path to the of the stratigraphic diagram image for this

activate()

Method that is called, when the page is activated

deactivate()

Method that is called, when the page is deactivated

hint()

A method that should display a hint to the user

property is_finished

Boolean that is True, if the steps are all finished

property is_selecting

True if the user clicked the `btn_select_data` button

lock_viewer(lock)

Set or unset the url lock of the HTML viewer

refresh()**show()**

Show the page and browse the `filename` in the tutorial docs

show_hint()

Show a hint to the user

show_tooltip_at_widget(tooltip, widget, timeout=20000)

Show a tooltip close to a widget

Parameters

- **tooltip** (`str`) – The tooltip to display
- **widget** (`QWidget`) – The widget that should be close to the tooltip
- **timeout** (`int`) – The time that the tool tip shall be displayed, in milliseconds

show_tooltip_in_plot(tooltip, x, y, timeout=20000, transform=None)

Show a tooltip in the matplotlib figure at the given coordinates

Parameters

- **tooltip** (`str`) – The tooltip to display
- **x** (`float`) – The x-coordinate of where to display the tooltip
- **y** (`float`) – The y-coordinate of where to display the tooltip
- **timeout** (`int`) – The time that the tool tip shall be displayed, in milliseconds
- **transform** (`matplotlib transformation`) – The matplotlib transformation to use. If None, the `self.stradi.ax.transData` transformation is used and x and y are expected to be in data coordinates

```
skip()
Skip the steps in this page

src_base = 'beginner-tutorial.png'
The basename of the stratigraphic diagram image for this tutorial

src_dir = '/home/docs/checkouts/readthedocs.org/user_builds/straditize/checkouts/latest'
The source directory for the docs

src_file = '/home/docs/checkouts/readthedocs.org/user_builds/straditize/checkouts/latest'
The complete path to the of the stratigraphic diagram image for this tutorial
```

straditize.widgets.tutorial.hoya_del_castillo module

The tutorial of straditize

This module contains an advanced guided tour through straditize

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Classes

<i>ColumnNames</i> (filename, tutorial)	The page for recognizing column names
<i>ColumnNamesOCR</i> (filename, tutorial)	Tutorial page for column names with tesserocr support
<i>EditMeta</i> (filename, tutorial)	Tutorial page for editing the meta attributes
<i>HoyaDelCastilloTutorial</i> (straditzer_widgets)	A tutorial for digitizing an area diagram
<i>LoadImage</i> (filename, tutorial)	param filename The basename (without ending) of the RST file corresponding to this
<i>RemoveLines</i> (filename, tutorial)	Tutorial page for removing horizontal and vertical lines
<i>SelectDataPart</i> (filename, tutorial)	TutorialPage for selecting the data part
<i>SeparateColumns</i> (filename, tutorial)	The page for separating the columns
<i>TranslateXAxis</i> (filename, tutorial)	The tutorial page for translating the x-axes
<i>TranslateYAxis</i> (filename, tutorial)	The tutorial page for translating the y-axis
<i>TutorialPage</i> (filename, tutorial)	param filename The basename (without ending) of the RST file corresponding to this

```
class straditize.widgets.tutorial.hoya_del_castillo.ColumnNames (filename, tutorial)
```

Bases: `straditize.widgets.tutorial.beginner.ColumnNames`

The page for recognizing column names

Parameters

- **filename** (`str`) – The basename (without ending) of the RST file corresponding to this tutorial page
- **tutorial** (`Tutorial`) – The tutorial instance

Attributes

<code>column_names</code>	Built-in mutable sequence.
<code>select_names_button_clicked</code>	<code>bool(x) -> bool</code>

```
column_names = ['Charcoal', 'Pinus', 'Juniperus', 'Quercus ilex-type', 'Quercus suber-'
    select_names_button_clicked = False
```

class `straditize.widgets.tutorial.hoya_del_castillo.ColumnNamesOCR` (`filename,`
 `tutorial`)

Bases: `straditize.widgets.tutorial.hoya_del_castillo.ColumnNames`

Tutorial page for column names with tesserocr support

Parameters

- **filename** (`str`) – The basename (without ending) of the RST file corresponding to this tutorial page
- **tutorial** (`Tutorial`) – The tutorial instance

Attributes

<code>colpic_extents</code>	Built-in mutable sequence.
<code>colpic_sizes</code>	Built-in mutable sequence.

Methods

<code>hint_for_start_editing()</code>	
<code>hint_for_wrong_name(col, curr, ref)</code>	Display a hint if a name is not correctly set

```
colpic_extents = [(1051, 1127, 573, 588), (1176, 1225, 704, 719), (1338, 1422, 866, 881),
    colpic_sizes = [(270, 88), (189, 88), (307, 109), (494, 109), (550, 109), (159, 85), (181, 85),
        hint_for_start_editing()
        hint_for_wrong_name(col, curr, ref)
```

Display a hint if a name is not correctly set

class `straditize.widgets.tutorial.hoya_del_castillo.EditMeta` (`filename, tutorial`)
 Bases: `straditize.widgets.tutorial.hoya_del_castillo.TutorialPage`

Tutorial page for editing the meta attributes

Parameters

- **filename** (`str`) – The basename (without ending) of the RST file corresponding to this tutorial page

- **tutorial** (`Tutorial`) – The tutorial instance

Methods

<code>hint()</code>	A method that should display a hint to the user
<code>skip()</code>	Skip the steps in this page

Attributes

<code>is_finished</code>	Boolean that is True, if the steps are all finished
--------------------------	---

`hint()`

A method that should display a hint to the user

`property is_finished`

Boolean that is True, if the steps are all finished

`skip()`

Skip the steps in this page

class straditize.widgets.tutorial.hoya_del_castillo.**HoyaDelCastilloTutorial** (`straditizer_widgets`)
Bases: `straditize.widgets.tutorial.beginner.Tutorial`

A tutorial for digitizing an area diagram **Methods**

<code>setup_tutorial_pages()</code>	Setup the pages attribute and initialize the tutorial pages
<code>show()</code>	Show the documentation of the tutorial

Attributes

<code>src_base</code>	<code>str(object=’’)</code> -> <code>str</code>
<code>src_dir</code>	<code>str(object=’’)</code> -> <code>str</code>
<code>src_file</code>	<code>str(object=’’)</code> -> <code>str</code>

`setup_tutorial_pages()`

Setup the pages attribute and initialize the tutorial pages

`show()`

Show the documentation of the tutorial

`src_base = ‘‘hoya-del-castillo.png’‘`

`src_dir = ‘‘/home/docs/checkouts/readthedocs.org/user_builds/straditize/checkouts/latest’‘`

`src_file = ‘‘/home/docs/checkouts/readthedocs.org/user_builds/straditize/checkouts/latest’‘`

class straditize.widgets.tutorial.hoya_del_castillo.**LoadImage** (`filename`, `tutorial`)
Bases: `straditize.widgets.tutorial.beginner.LoadImage`, `straditize.widgets.tutorial.hoya_del_castillo.TutorialPage`

Parameters

- **filename** (`str`) – The basename (without ending) of the RST file corresponding to this tutorial page
- **tutorial** (`Tutorial`) – The tutorial instance

class straditize.widgets.tutorial.hoya_del_castillo.**RemoveLines** (*filename*, *tutorial*)
Bases: *straditize.widgets.tutorial.hoya_del_castillo.TutorialPage*

Tutorial page for removing horizontal and vertical lines

Parameters

- **filename** (*str*) – The basename (without ending) of the RST file corresponding to this tutorial page
- **tutorial** (*Tutorial*) – The tutorial instance

Methods

<i>activate()</i>	Method that is called, when the page is activated
<i>clicked_hlines_button()</i>	
<i>clicked_vlines_button()</i>	
<i>deactivate()</i>	Method that is called, when the page is deactivated
<i>hint()</i>	A method that should display a hint to the user
<i>skip()</i>	Skip the steps in this page

Attributes

<i>hlines_button_clicked</i>	bool(x) -> bool
<i>is_finished</i>	Boolean that is True, if the steps are all finished
<i>vlines_button_clicked</i>	bool(x) -> bool

activate()

Method that is called, when the page is activated

clicked_hlines_button()

clicked_vlines_button()

deactivate()

Method that is called, when the page is deactivated

hint()

A method that should display a hint to the user

hlines_button_clicked = False

property is_finished

Boolean that is True, if the steps are all finished

skip()

Skip the steps in this page

vlines_button_clicked = False

class straditize.widgets.tutorial.hoya_del_castillo.**SelectDataPart** (*filename*, *tutorial*)
Bases: *straditize.widgets.tutorial.beginner.SelectDataPart*

TutorialPage for selecting the data part

Parameters

- **filename** (*str*) – The basename (without ending) of the RST file corresponding to this tutorial page

- **tutorial** (`Tutorial`) – The tutorial instance

Attributes

<code>ref_lims</code>	The reference x- and y- limits
<code>remove_data_box</code>	<code>bool(x) -> bool</code>
<code>valid_xlims</code>	Valid ranges for xmin and xmax
<code>valid_ylims</code>	Valid ranges for ymin and ymax

```
ref_lims = array([[ 315, 1946], [ 511, 1311]])  
The reference x- and y- limits  
remove_data_box = False  
valid_xlims = array([[ 310, 319], [1928, 1960]])  
Valid ranges for xmin and xmax  
valid_ylims = array([[ 508, 513], [1307, 1312]])  
Valid ranges for ymin and ymax
```

class straditize.widgets.tutorial.hoya_del_castillo.**SeparateColumns** (`filename, tutorial`)
Bases: *straditize.widgets.tutorial.beginner.SeparateColumns*

The page for separating the columns

Parameters

- **filename** (`str`) – The basename (without ending) of the RST file corresponding to this tutorial page
- **tutorial** (`Tutorial`) – The tutorial instance

Attributes

<code>ncols</code>	<code>int([x]) -> integer</code>
--------------------	-------------------------------------

`ncols = 28`

class straditize.widgets.tutorial.hoya_del_castillo.**TranslateXAxis** (`filename, tutorial`)
Bases: *straditize.widgets.tutorial.hoya_del_castillo.TutorialPage*

The tutorial page for translating the x-axes

Parameters

- **filename** (`str`) – The basename (without ending) of the RST file corresponding to this tutorial page
- **tutorial** (`Tutorial`) – The tutorial instance

Methods

<code>activate()</code>	Method that is called, when the page is activated
<code>clicked_add_reader_button()</code>	
<code>clicked_translations_button()</code>	
<code>deactivate()</code>	Method that is called, when the page is deactivated
<code>hint()</code>	A method that should display a hint to the user
<code>hint_for_col(col)</code>	

Continued on next page

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<code>refresh()</code>	
<code>skip()</code>	Skip the steps in this page

Attributes

<code>add_reader_button_clicked</code>	Built-in mutable sequence.
<code>is_finished</code>	Boolean that is True, if the steps are all finished
<code>xaxis_translations_button_clicked</code>	<code>bool(x) -> bool</code>

`activate()`

Method that is called, when the page is activated

`add_reader_button_clicked = []`

`clicked_add_reader_button()`

`clicked_translations_button()`

`deactivate()`

Method that is called, when the page is deactivated

`hint()`

A method that should display a hint to the user

`hint_for_col(col)`

`property is_finished`

Boolean that is True, if the steps are all finished

`refresh()`

`skip()`

Skip the steps in this page

`xaxis_translations_button_clicked = False`

class straditize.widgets.tutorial.hoya_del_castillo.TranslateYAxis(`filename, tutorial`)

Bases: `straditize.widgets.tutorial.hoya_del_castillo.TutorialPage`

The tutorial page for translating the y-axis

Parameters

- `filename (str)` – The basename (without ending) of the RST file corresponding to this tutorial page
- `tutorial (Tutorial)` – The tutorial instance

Methods

<code>activate()</code>	Method that is called, when the page is activated
<code>clicked_correct_button()</code>	
<code>deactivate()</code>	Method that is called, when the page is deactivated
<code>hint()</code>	A method that should display a hint to the user
<code>skip()</code>	Skip the steps in this page

Attributes

<code>correct_button_clicked</code>	bool(x) -> bool
<code>is_finished</code>	Boolean that is True, if the steps are all finished

activate()

Method that is called, when the page is activated

clicked_correct_button()**correct_button_clicked = False****deactivate()**

Method that is called, when the page is deactivated

hint()

A method that should display a hint to the user

property is_finished

Boolean that is True, if the steps are all finished

skip()

Skip the steps in this page

class straditize.widgets.tutorial.hoya_del_castillo.TutorialPage (*filename*, *tutorial*)

Bases: straditize.widgets.tutorial.beginner.TutorialPage

Parameters

- **filename** (*str*) – The basename (without ending) of the RST file corresponding to this tutorial page
- **tutorial** (*Tutorial*) – The tutorial instance

Attributes

<code>src_base</code>	str(object="") -> str
<code>src_dir</code>	str(object="") -> str
<code>src_file</code>	str(object="") -> str

`src_base = 'hoya-del-castillo.png'`

`src_dir = '/home/docs/checkouts/readthedocs.org/user_builds/straditize/checkouts/latest'`

`src_file = '/home/docs/checkouts/readthedocs.org/user_builds/straditize/checkouts/latest'`

Submodules**straditize.widgets.axes_translations module**

Module for translating the x- and y-axes from pixel into data coordinates

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Classes

<code>AxesTranslations(straditizer_widgets, item)</code>	The control for translating x- and y-axes
--	---

class straditize.widgets.axes_translations.**AxesTranslations** (*straditizer_widgets, item*)

Bases: *straditize.widgets.StraditizerControlBase*

The control for translating x- and y-axes

This object creates two buttons for translating x- and y-axes from pixel to data coordinates

Parameters

- **straditizer_widgets** (*StraditizerWidgets*) – The main widget for the straditzer GUI
- **item** (*QTreeWidgetItem*) – The parent item in the *StraditizerWidgets.tree*. If given, the *setup_children()* is called with this item

Methods

<code>marks_for_x([at_col_start])</code>	Create (or enable) the marks for the x-axis translation
<code>marks_for_y()</code>	Create (or enable) the marks for the y-axis translation
<code>setup_children(item)</code>	Setup the children for this control
<code>should_be_enabled(w)</code>	Check if a widget should be enabled

Attributes

`tree`

marks_for_x (*at_col_start=True*)

Create (or enable) the marks for the x-axis translation

See also:

straditze.straditizer.Straditizer.marks_for_x_values(), straditze.straditizer.Straditizer.update_xvalues()

marks_for_y ()

Create (or enable) the marks for the y-axis translation

See also:

straditze.straditizer.Straditizer.marks_for_y_values(), straditze.straditizer.Straditizer.update_yvalues()

setup_children (*item*)

Setup the children for this control

This method is called to setup the children in the *StraditizerWidgets.tree*. By default, it just creates a child *QTreeWidgetItem* and sets this control as its widget

Parameters `item` (`QTreeWidgetItem`) – The top level item in the `StraditizerWidgets.tree`

should_be_enabled(`w`)
Check if a widget should be enabled
This function checks if a given widget `w` from the `widgets2disable` attribute should be enabled or not

Parameters `w` (`QWidget`) – The widget to check

Returns True, if the widget should be enabled

Return type `bool`

property tree

straditize.widgets.colnames module

Widget for handling column names

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Classes

<code>ColumnNamesManager(straditizer_widgets[, item])</code>	Manage the column names of the reader
<code>DummyNavigationToolbar2(canvas)</code>	Reimplemented <code>NavigationToolbar2</code> just to add an <code>_init_toolbar</code> method

class straditize.widgets.colnames.**ColumnNamesManager** (`straditizer_widgets,`
`item=None, *args, **kwargs)`
Bases: `straditize.widgets.StraditizerControlBase, psyplot_gui.common.DockMixin, PyQt5.QtWidgets.QSplitter`

Manage the column names of the reader

Parameters

- **straditizer_widgets** (`StraditizerWidgets`) – The main widget for the straditzer GUI
- **item** (`QTreeWidgetItem`) – The parent item in the `StraditizerWidgets.tree`. If given, the `setup_children()` is called with this item

Attributes

<code>NAVIGATION_LABEL</code>	<code>str(object="") -> str</code>
<code>SELECT_LABEL</code>	<code>str(object="") -> str</code>
<code>btn_find</code>	A QPushButton to find column names in the visible part of the
<code>btn_load_image</code>	A QPushButton to load the highres image
<code>btn_recognize</code>	A QPushButton to recognize text in the colpic
<code>btn_select_colpic</code>	A checkable QPushButton to initialize a selector to select the
<code>btn_select_names</code>	The QPushButton in the <code>straditize.widgets.StraditzerWidgets</code>
<code>cb_find_all_cols</code>	A QCheckBox to find the column names (see <code>btn_find</code>) for all
<code>cb_fliph</code>	A QCheckBox to set the <code>straditize.colnames.ColNamesReader.mirror</code>
<code>cb_flipv</code>	A QCheckBox to set the <code>straditize.colnames.ColNamesReader.flip</code>
<code>cb_ignore_data_part</code>	A QCheckBox to ignore the part within the
<code>colnames_reader</code>	The <code>straditize.straditzer.Straditzer.colnames_reader</code>
<code>colnames_table</code>	A QTableWidget to display the column names
<code>colpic</code>	The <code>PIL.Image.Image</code> of the column name (see also)
<code>colpic_ax</code>	The <code>matplotlib.axes.Axes</code> to display the <code>colpic_im</code>
<code>colpic_canvas</code>	The canvas to display the <code>colpic_im</code>
<code>colpic_extents</code>	The extents of the <code>colpic</code> in the <code>im_rotated</code>
<code>colpic_im</code>	The matplotlib image of the <code>colpic</code>
<code>current_col</code>	The currently selected column
<code>fig_h</code>	The original height of the <code>main_canvas</code>
<code>fig_w</code>	The original width of the <code>main_canvas</code>
<code>im_rotated</code>	The matplotlib image of the
<code>main_ax</code>	The <code>matplotlib.axes.Axes</code> to display the <code>im_rotated</code>
<code>main_canvas</code>	The canvas to display the <code>im_rotated</code>
<code>rect</code>	The rectangle to highlight a column (see <code>highlight_selected_col()</code>)
<code>refreshing</code>	True if the widget is refreshing
<code>selector</code>	The <code>matplotlib.widgets.RectangleSelector</code> to select the
<code>txt_rotate</code>	A QLineEdit to set the <code>straditize.colnames.ColNamesReader.rotate</code>

Methods

<code>adjust_lims()</code>	Adjust the limits of the <code>main_ax</code> to fill the entire figure
<code>adjust_lims_after_resize(event)</code>	Adjust the limits of the <code>main_ax</code> after resize of the figure
<code>cancel_colpic_selection()</code>	Stop the colpic selection in the <code>im_rotated</code>
<code>change_ignore_data_part(checked)</code>	Change <code>straditize.colnames.ColNamesReader.ignore_data_part</code>

Continued on next page

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<code>colname_changed(row, column)</code>	Update the column name in the <code>colnames_reader</code>
<code>create_selector()</code>	Create the selector to enable <code>colpic</code> selection
<code>enable_or_disable_btn_find(*args, **kwargs)</code>	
<code>find_colnames([warn, full_image, all_cols])</code>	Find the column names automatically
<code>flip(checked)</code>	TFlip the image
<code>highlight_selected_col()</code>	Highlight the column selected in the <code>colnames_tables</code>
<code>load_image()</code>	Load a high resolution image
<code>maybe_tabify()</code>	
<code>mirror(checked)</code>	Mirror the image
<code>plot_colpic()</code>	Plot the <code>colpic</code> in the <code>colpic_ax</code>
<code>read_colpic()</code>	Recognize the text in the <code>colpic</code>
<code>refresh()</code>	Refresh from the straditizer
<code>remove_images()</code>	Remove the <code>im_rotated</code> and the <code>colpic_im</code>
<code>remove_selector()</code>	Remove and disconnect the selector
<code>replot_figure()</code>	Remove and replot the <code>im_rotated</code>
<code>reset_control()</code>	Reset the dialog
<code>rotate(val)</code>	Rotate the image
<code>set_xc_yc()</code>	Set the x- and y-center before rotating or flipping
<code>setup_children(item)</code>	Setup the children for this control
<code>should_be_enabled(w)</code>	Check if a widget should be enabled
<code>to_dock(main[, title, position])</code>	
<code>toggle_colpic_selection()</code>	Enable or disable the <code>colpic</code> selection
<code>toggle_dialog()</code>	Close the dialog when the <code>btn_select_names</code> button is clicked
<code>update_image(*args, **kwargs)</code>	Update the <code>colpic</code> with the extents of the selector

```

NAVIGATION_LABEL = 'Use left-click of your mouse to move the image below and right-click to rotate'
SELECT_LABEL = 'Left-click and hold on the image to select the column name'
adjust_lims()
    Adjust the limits of the main_ax to fill the entire figure
adjust_lims_after_resize(event)
    Adjust the limits of the main_ax after resize of the figure
btn_find = None
    A QPushButton to find column names in the visible part of the im_rotated
btn_load_image = None
    A QPushButton to load the highres image
btn_recognize = None
    A QPushButton to recognize text in the colpic
btn_select_colpic = None
    A checkable QPushButton to initialize a selector to select the colpic
btn_select_names = None
    The QPushButton in the straditize.widgets.StraditzerWidgets to toggle the column names dialog

```

```

cancel_colpic_selection()
    Stop the colpic selection in the im_rotated

cb_find_all_cols = None
    A QCheckBox to find the column names (see btn_find) for all columns and not just the one selected in
    the colnames_table

cb_fliph = None
    A QCheckBox to set the straditize.colnames.ColNamesReader.mirror

cb_flipv = None
    A QCheckBox to set the straditize.colnames.ColNamesReader.flip

cb_ignore_data_part = None
    A QCheckBox to ignore the part within the straditize.colnames.ColNamesReader.data_ylim

change_ignore_data_part(checked)
    Change straditize.colnames.ColNamesReader.ignore_data_part

colname_changed(row, column)
    Update the column name in the colnames_reader

    This method is called when a cell in the colnames_table has been changed and updates the corre-
    sponding name in the colnames_reader

Parameters

- row (int) – The row of the cell in the colnames_table that changed
- column (int) – The column of the cell in the colnames_table that changed

property colnames_reader
    The straditize.straditizer.Straditizer.colnames_reader of the current straditizer

colnames_table = None
    A QTableWidget to display the column names

colpic = None
    The PIL.Image.Image of the column name (see also straditize.colnames.  

ColNamesReader.colpics)

colpic_ax = None
    The matplotlib.axes.Axes to display the colpic_im

colpic_canvas = None
    The canvas to display the colpic_im

colpic_extents = None
    The extents of the colpic in the im_rotated

colpic_im = None
    The matplotlib image of the colpic

create_selector()
    Create the selector to enable colpic selection

property current_col
    The currently selected column

enable_or_disable_btn_find(*args, **kwargs)

fig_h = None
    The original height of the main_canvas

```

fig_w = None

The original width of the `main_canvas`

find_colnames (warn=True, full_image=False, all_cols=None)

Find the column names automatically

See also:

`straditize.colnames.ColNamesReader.find_colnames()`

flip (checked)

TFlip the image

highlight_selected_col ()

Highlight the column selected in the `colnames_tables`

See also:

`straditize.colnames.ColNamesReader.highlight_column()`

im_rotated = None

The matplotlib image of the `straditize.colnames.ColNamesReader.rotated_image`

load_image ()

Load a high resolution image

main_ax = None

The `matplotlib.axes.Axes` to display the `im_rotated`

main_canvas = None

The canvas to display the `im_rotated`

maybe_tabify ()

mirror (checked)

Mirror the image

plot_colpic ()

Plot the `colpic` in the `colpic_ax`

read_colpic ()

Recognize the text in the `colpic`

See also:

`straditize.colnames.ColNamesReader.recognize_text()`

rect = None

The rectangle to highlight a column (see `highlight_selected_col ()`)

refresh ()

Refresh from the straditizer

property refreshing

True if the widget is refreshing

remove_images ()

Remove the `im_rotated` and the `colpic_im`

remove_selector ()

Remove and disconnect the `selector`

replot_figure ()

Remove and replot the `im_rotated`

reset_control()
Reset the dialog

rotate(val)
Rotate the image

Parameters `float` – The angle for the rotation

selector = None
The `matplotlib.widgets.RectangleSelector` to select the `colpic`

set_xc_yc()
Set the x- and y-center before rotating or flipping

setup_children(item)
Setup the children for this control

This method is called to setup the children in the `StraditzerWidgets.tree`. By default, it just creates a child `QTreeWidgetItem` and sets this control as its widget

Parameters `item (QTreeWidgetItem)` – The top level item in the `StraditzerWidgets.tree`

should_be_enabled(w)
Check if a widget should be enabled

This function checks if a given widget `w` from the `widgets2disable` attribute should be enabled or not

Parameters `w (QWidget)` – The widget to check

Returns True, if the widget should be enabled

Return type `bool`

to_dock(main, title=None, position=None, *args, **kwargs)

toggle_colpic_selection()
Enable or disable the colpic selection

toggle_dialog()
Close the dialog when the `btn_select_names` button is clicked

txt_rotate = None
A QLineEdit to set the `straditize.colnames.ColNamesReader.rotate`

update_image(*args, **kwargs)
Update the `colpic` with the extents of the `selector`
`*args` and `**kwargs` are ignored

class straditize.widgets.colnames.DummyNavigationToolbar2(canvas)
Bases: `matplotlib.backend_bases.NavigationToolbar2`

Reimplemented `NavigationToolbar2` just to add an `_init_toolbar` method **Methods**

<code>set_cursor(cursor)</code>	Set the current cursor to one of the <code>Cursors</code> enums values.
---------------------------------	---

set_cursor(cursor)
Set the current cursor to one of the `Cursors` enums values.
If required by the backend, this method should trigger an update in the backend event loop after the cursor is set, as this method may be called e.g. before a long-running task during which the GUI is not updated.

straditize.widgets.data module

The main control widget for handling the data

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Classes

<code>BarSplitter(straditizer_widgets, *args, **kwargs)</code>	A widget for splitting bars
<code>DigitizingControl(straditizer_widgets, item)</code>	An interface to <code>straditze.straditzer.Straditzer.data_reader</code>

Functions

<code>get_reader_name(reader)</code>	Get the reader key in the straditize.binary.readers dictionary
<code>int_list2str(numbers)</code>	Create a short string representation of an integer list

class straditze.widgets.data.**BarSplitter**(straditizer_widgets, *args, **kwargs)
Bases: PyQt5.QtWidgets.QTreeWidget, straditze.widgets.StraditzerControlBase

A widget for splitting bars **Methods**

<code>add_toolbar_widgets()</code>	Add an action to switch between the bars to the matplotlib toolbar
<code>disconnect()</code>	Disconnect the events to split an item
<code>enable_or_disable_widgets(b)</code>	Enable or disable the widgets in this control
<code>fill_table([source])</code>	Fill the table with the bars that should be splitted
<code>get_overlapping_bars()</code>	Get the bars the overlap with multiple bars in another column
<code>go_to_next_bar()</code>	Go to the next_item
<code>go_to_prev_bar()</code>	Go to the previous_item
<code>new_split(y, y0[, draw_figure])</code>	Mark the current item to be splitted at y
<code>prepare_for_split(event)</code>	Select or deselect a split location
<code>refresh()</code>	Reimplemented to use the <code>fill_table()</code> method
<code>remove_actions()</code>	Remove the actions added by <code>add_toolbar_widgets()</code>
<code>remove_lines()</code>	Remove the plotted lines
<code>remove_split_children()</code>	Remove all the child items that mark a split
<code>revert_split(y, y0)</code>	Revert the split

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<code>set_suggestions_fig_titles(ax)</code>	Set the title in the suggestion figure with the displayed column
<code>split_bars()</code>	Split the bars after they have been separated manually
<code>start_splitting(item, *args, **kwargs)</code>	Enable the splitting for the selected item
<code>suggest_splits()</code>	Find overlaps for the current selected bar in other columns

Attributes

<code>next_item</code>	The QTreeWidgetItem for the next bar to split
<code>prev_action</code>	The action in the matplotlib toolbar to go to the previous bar to split
<code>previous_item</code>	The QTreeWidgetItem for the previous bar to split
<code>selected_child</code>	The QTreeWidgetItem that is currently shown in the plot
<code>suggestions_fig</code>	A figure to show the other columns

`add_toolbar_widgets()`

Add an action to switch between the bars to the matplotlib toolbar

`disconnect()`

Disconnect the events to split an item

`enable_or_disable_widgets(b)`

Enable or disable the widgets in this control

This method enables or disables the `widgets2disable` if the `should_be_enabled()` method evaluates to True

Parameters `b` (`bool`) – If True, enable the widgets, if False, disable them

`fill_table(source='too-long')`

Fill the table with the bars that should be splitted

Parameters `source` ({ 'too-long' / 'overlaps' / 'all' }) – The source with what to fill the table.

too-long Only display the bars that are considered as *too long*

overlap Only display the bars that overlap with multiple bars in another column (see `get_overlapping_bars()`)

all Display all bars

`get_overlapping_bars()`

Get the bars the overlap with multiple bars in another column

`go_to_next_bar()`

Go to the `next_item`

`go_to_prev_bar()`

Go to the `previous_item`

`new_split(y, y0, draw_figure=True)`

Mark the current item to be splitted at `y`

This method draws a horizontal line at `y` and adds a new child QTreeWidgetItem to the `selected_child` to mark the split

Parameters

- **y** (*int*) – The vertical position of the split in the data image coordinate system
- **y0** (*int*) – The vertical start of the data image (see *straditiz.binary.Reader.extent*)
- **draw_figure** (*bool*) – If True, draw the figure

property next_item

The QTreeWidgetItem for the next bar to split

prepare_for_split (*event*)

Select or deselect a split location

LeftButton selects the given location for a new split (see *new_split()*), RightButton deselects it (see *revert_split()*)

prev_action = None

The action in the matplotlib toolbar to go to the previous bar to split

property previous_item

The QTreeWidgetItem for the previous bar to split

refresh()

Reimplemented to use the *fill_table()* method

remove_actions()

Remove the actions added by *add_toolbar_widgets()*

remove_lines()

Remove the plotted lines

remove_split_children()

Remove all the child items that mark a split

revert_split (*y, y0*)

Revert the split

Parameters

- **y** (*int*) – The vertical position of the split in the data image coordinate system
- **y0** (*int*) – The vertical start of the data image (see *straditiz.binary.Reader.extent*)

selected_child = None

The QTreeWidgetItem that is currently shown in the plot

set_suggestions_fig_titles (*ax*)

Set the title in the suggestion figure with the displayed column

split_bars()

Split the bars after they have been separated manually

start_splitting (*item, *args, **kwargs*)

Enable the splitting for the selected item

suggest_splits()

Find overlaps for the current selected bar in other columns

suggestions_fig = None

A figure to show the other columns

```
class straditize.widgets.data.DigitizingControl(straditizer_widgets, item)
```

Bases: straditize.widgets.StraditzerControlBase

An interface to straditize.straditzer.Straditzer.data_reader

This widgets contains the functionalities to interface with the data readers for the stratigraphic diagram

Parameters

- **straditizer_widgets** (StraditzerWidgets) – The main widget for the straditzer GUI
- **item** (QTreeWidgetItem) – The parent item in the StraditzerWidgets.tree. If given, the `setup_children()` is called with this item

Methods

<code>align_vertical()</code>	Create marks for vertical alignment of the columns
<code>change_reader(txt)</code>	Change the current parent reader
<code>digitize()</code>	Digitize the data
<code>digitize_exaggerations()</code>	Digitize the data
<code>edit_occurrences()</code>	Enable the editing of occurrences
<code>edit_samples()</code>	Enable the sample editing
<code>enable_col_selection_for_new_reader()</code>	Start the selection process to get a new reader for specific cols
<code>enable_occurrences_selection()</code>	Enable the selection of occurrences
<code>enable_or_disable_btn_highlight_small()</code>	Enable the btn_highlight_small_selection during a selection
<code>enable_or_disable_widgets(*args, **kwargs)</code>	Enable or disable the widgets in this control
<code>fill_cb_readers()</code>	Fill the cb_readers combo based on the current reader
<code>find_samples()</code>	
<code>finish_exaggerated_features()</code>	Save the exaggerations in the exaggerations reader
<code>init_exaggerated_reader()</code>	Initialize the reader for exaggeration features
<code>init_reader()</code>	Initialize the reader
<code>load_samples([fname])</code>	Load the samples of a text file
<code>maybe_show_btn_reset_columns()</code>	Show the btn_reset_columns if the column starts are set
<code>maybe_show_btn_reset_samples()</code>	Show the btn_reset_samples if the samples are set
<code>modify_column_ends()</code>	Modify the column ends
<code>new_reader_for_selection([cls])</code>	Create a new child reader for the selected columns
<code>refresh()</code>	Refresh from the straditzer
<code>remove_hlines()</code>	Remove horizontal lines
<code>remove_vlines()</code>	Remove vertical lines
<code>remove_xaxes()</code>	Remove x-axes in the plot
<code>remove_yaxes()</code>	Remove y-axes in the plot
<code>reset_column_starts()</code>	Reset the column starts
<code>reset_samples()</code>	Reset the samples
<code>select_column_starts()</code>	Estimate the column starts and draw marks
<code>select_data_part([guess_lims])</code>	Enable the selection of the diagram part
<code>select_exaggerated_features()</code>	Enable the selection of exaggerated features
<code>select_occurrences()</code>	Save (and potentially remove) the selected occurrences

Continued on next page

Table 68 – continued from previous page

<code>set_occurrences_value(value)</code>	Set the <code>occurrences_value</code>
<code>setup_children(item)</code>	Set up the child items for a topLevelItem in the control tree
<code>should_be_enabled(w)</code>	Check if a widget should be enabled
<code>show_cross_column_features()</code>	Remove cross column features
<code>show_disconnected_parts()</code>	Remove disconnected parts
<code>show_parts_at_column_ends()</code>	Remove parts that touch the column ends
<code>show_small_parts()</code>	Remove parts that touch the column ends
<code>toggle_bar_split_source(i)</code>	Fill the <code>tree_bar_split</code> based on the <code>cb_split_source</code>
<code>toggle_btn_highlight_small_selection()</code>	Enable or disable the <code>btn_highlight_small_selection</code>
<code>toggle_sp_max_lw(state)</code>	Toggle <code>sp_max_lw</code> based on <code>cb_max_lw</code>
<code>toggle_txt_edit_rows(state)</code>	Toggle <code>txt_edit_rows</code> based on <code>cb_edit_separate</code>
<code>toggle_txt_from0(state)</code>	Toggle <code>txt_from0</code> based on <code>cb_from0</code>
<code>toggle_txt_fromlast(state)</code>	Toggle <code>txt_fromlast</code> based on <code>cb_fromlast</code>
<code>toggle_txt_tolerance(s)</code>	Set the visibility of the <code>txt_tolerance</code> based on the reader
<code>update_tolerance(s)</code>	Set the readers tolerance

Attributes

<code>btn_column_ends</code>	Button for selecting and modifying column ends
<code>btn_column_starts</code>	Button for selecting and modifying column starts, see the <code>straditize.straditizer.Straditizer.marks_for_column_starts()</code> method.
<code>btn_digitize</code>	Button for digitizing the diagram
<code>btn_digitize_exag</code>	Button to digitize the exaggerations
<code>btn_edit_occurrences</code>	A button to edit the occurrences with the
<code>btn_edit_samples</code>	A button to edit the samples (see the
<code>btn_find_samples</code>	A button to find the samples with the
<code>btn_highlight_small_selection</code>	A button to highlight small selections using the
<code>btn_init_reader</code>	Button for initializing the reader
<code>btn_load_samples</code>	A button to load samples from a file
<code>btn_new_child_reader</code>	Button to add a new column-specific child reader
<code>btn_new_exaggeration</code>	Button to add an exaggerations reader
<code>btn_remove_hlines</code>	button for removing horizontal lines
<code>btn_remove_vlines</code>	Button for removing vertical lines
<code>btn_remove_xaxes</code>	button for removing x-axes
<code>btn_remove_yaxes</code>	Button for removing y-axes
<code>btn_reset_columns</code>	Button to reset the column starts and ends
<code>btn_reset_samples</code>	A button to reset the samples
<code>btn_select_data</code>	Button for selecting the data box, see the <code>straditize.straditizer.Straditizer.marks_for_data_selection()</code> method.
<code>btn_select_exaggerations</code>	Button to select the exaggerations
<code>btn_select_occurrences</code>	A button to select occurrences in the data part (see the

Continued on next page

Table 69 – continued from previous page

<code>btn_show_cross_column</code>	A button to
<code>btn_show_disconnected_parts</code>	Button for removing disconnected parts in the plot.
<code>btn_show_parts_at_column_ends</code>	A button to show the parts that touch the column end
<code>btn_show_small_parts</code>	A button to <code>show_small_parts()</code>
<code>cb_edit_separate</code>	A QCheckBox to edit the samples in a separate figure and not inside the
<code>cb_exag_reader_type</code>	A QComboBox to select the reader type for exaggerations
<code>cb_from0</code>	A QCheckBox to enable and disable the <code>from0</code> keyword in the
<code>cb_fromlast</code>	A QCheckBox to enable and disable the <code>fromlast</code> keyword in the
<code>cb_max_lw</code>	A QCheckBox to enable and disable the maximum linewidth as a criterion
<code>cb_reader_type</code>	Combobox for selecting the reader type
<code>cb_remove_occurrences</code>	A QCheckBox to remove the occurrences in the plot after selection
<code>cb_split_source</code>	A QComboBox to select whether to fill the <code>tree_bar_split</code> with
<code>reader</code>	The <code>straditize.straditizer.Straditizer.data_reader</code>
<code>selection_toolbar</code>	
<code>sp_max_lw</code>	A QSpinBox to select the maximum linewidth
<code>sp_min_lw</code>	A QSpinBox to select the minimum linewidth
<code>sp_pixel_tol</code>	A QSpinBox to set the minimum distance between to samples in the
<code>tree</code>	The <code>straditize.widgets.StraditizerWidgets.tree</code>
<code>tree_bar_split</code>	A <code>BarSplitter</code> to split too long bars
<code>txt_column_thresh</code>	A QLineEdit to set the threshold for the column starts detection
<code>txt_cross_column_px</code>	A QLineEdit to select the minimum pixels (<code>min_px</code>) for a cross column
<code>txt_edit_rows</code>	A QLineEdit to specify the number of rows in a plot for editing the
<code>txt_exag_factor</code>	A QLineEdit for the exaggeration factor
<code>txt_from0</code>	A QLineEdit to set the <code>from0</code> keyword for the
<code>txt_fromlast</code>	A QLineEdit to set the <code>fromlast</code> keyword for the
<code>txt_line_fraction</code>	LineEdit for specifying the fraction of vertical and horizontal lines
<code>txt_max_len</code>	A QLineEdit to specify the maximum length of a potential sample to be
<code>txt_max_small_size</code>	A QLineEdit to set the size for small parts
<code>txt_min_highlight</code>	A QLineEdit to set the maximal size for
<code>txt_min_len</code>	A QLineEdit to specify the minimum length of a potential sample to be
<code>txt_occurrences_value</code>	A QLineEdit to set the value for occurrences in the final data
<code>txt_tolerance</code>	Line edit for setting the tolerance for bars

`align_vertical()`

Create marks for vertical alignment of the columns

See also:

`straditize.straditizer.Straditizer.marks_for_vertical_alignment()`,
`straditize.straditizer.Straditizer.align_columns()`

btn_column_ends = None

Button for selecting and modifying column ends

btn_column_starts = None

Button for selecting and modifying column starts, see the `straditize.straditizer.Straditizer.marks_for_column_starts()` method.

btn_digitize = None

Button for digitizing the diagram

btn_digitize_exag = None

Button to digitize the exaggerations

btn_edit_occurrences = None

A button to edit the occurrences with the `straditize.straditizer.Straditizer.marks_for_occurrences()` method

btn_edit_samples = None

A button to edit the samples (see the `straditize.straditizer.Straditizer.marks_for_samples()` and `straditize.straditizer.Straditizer.marks_for_samples_sep()`)

btn_find_samples = None

A button to find the samples with the `straditize.binary.DataReader.find_samples()` method

btn_highlight_small_selection = None

A button to highlight small selections using the `straditize.label_selection.LabelSelection.highlight_small_selections()` method

btn_init_reader = None

Button for initializing the reader

btn_load_samples = None

A button to load samples from a file

btn_new_child_reader = None

Button to add a new column-specific child reader

btn_new_exaggeration = None

Button to add an exaggerations reader

btn_remove_hlines = None

button for removing horizontal lines

btn_remove_vlines = None

Button for removing vertical lines

btn_remove_xaxes = None

button for removing x-axes

btn_remove_yaxes = None

Button for removing y-axes

btn_reset_columns = None

Button to reset the column starts and ends

btn_reset_samples = None
A button to reset the samples

btn_select_data = None
Button for selecting the data box, see the `straditize.straditizer.Straditizer.marks_for_data_selection()` method.

btn_select_exaggerations = None
Button to select the exaggerations

btn_select_occurrences = None
A button to select occurrences in the data part (see the `enable_occurrences_selection()` method)

btn_show_cross_column = None
A button to `show_cross_column_features()`

btn_show_disconnected_parts = None
Button for removing disconnected parts in the plot. See the `straditize.binary.DataReader.show_disconnected_parts()` method

btn_show_parts_at_column_ends = None
A button to show the parts that touch the column end

btn_show_small_parts = None
A button to `show_small_parts()`

cb_edit_separate = None
A QCheckBox to edit the samples in a separate figure and not inside the original diagram

cb_exag_reader_type = None
A QComboBox to select the reader type for exaggerations

cb_from0 = None
A QCheckBox to enable and disable the `from0` keyword in the `straditize.binary.DataReader.show_disconnected_parts()` method

cb_fromlast = None
A QCheckBox to enable and disable the `fromlast` keyword in the `straditize.binary.DataReader.show_disconnected_parts()` method

cb_max_lw = None
QCheckBox to enable and disable the maximum linewidth as a criterion

cb_reader_type = None
Combobox for selecting the reader type

cb_remove_occurrences = None
A QCheckBox to remove the occurrences in the plot after selection

cb_split_source = None
A QComboBox to select whether to fill the `tree_bar_split` with too long, overlapping, or all bars

change_reader(txt)
Change the current parent reader

This changes the `straditize.straditizer.Straditizer.data_reader` using the `straditize.binary.DataReader.set_as_parent()` method

Parameters s (str) – A string matching 'Columns (\d.*)', where the numbers are the columns of the reader to use

digitize()
Digitize the data

This method uses the `straditize.binary.DataReader.digitize()` method to digitize the data of the current reader

digitize_exaggerations()

Digitize the data

This method uses the `straditize.binary.DataReader.digitize_exaggerated()` method to digitize the exaggerated data of the current reader and merge it into the data obtained by the `digitize()` method.

digitize_item = None**edit_occurences()**

Enable the editing of occurences

This enables the editing of occurences using the `straditize.straditizer.Straditizer.marks_for_occurences()` method for the occurences selected by the `select_occurences()` method

edit_samples()

Enable the sample editing

This method opens a `straditize.widgets.samples_table.MultiCrossMarksEditor` or a `straditize.widgets.samples_table.SingleCrossMarksEditor` to edit the samples in the GUI. Depending on whether the `cb_edit_separate` is checked or not, we use the `straditize.straditizer.Straditizer.marks_for_samples_sep()` or `straditize.straditizer.Straditizer.marks_for_samples()` method.

enable_col_selection_for_new_reader()

Start the selection process to get a new reader for specific cols

enable_occurences_selection()

Enable the selection of occurences

This method starts the selection of features in the data image and connects the `select_occurences()` to the `apply_button`.

enable_or_disable_btn_highlight_small_selection()

Enable the `btn_highlight_small_selection` during a selection

enable_or_disable_widgets(*args, **kwargs)

Enable or disable the widgets in this control

This method enables or disables the `widgets2disable` if the `should_be_enabled()` method evaluates to True

Parameters `b` (`bool`) – If True, enable the widgets, if False, disable them

fill_cb_readers()

Fill the `cb_readers` combo based on the current reader

find_samples()**finish_exaggerated_features()**

Save the exaggerations in the exaggerations reader

This method finalizes the operation initialized by the `select_exaggerated_features()` by calling the `straditize.binary.DataReader.mark_as_exaggerations()` method

init_exaggerated_reader()

Initialize the reader for exaggeration features

init_reader()

Initialize the reader

Initialize the data reader with the `straditize.straditizer.Straditizer.init_reader()` method

load_samples (`fname=None`)
Load the samples of a text file

This method asks for a filename to update the samples. The first column in this file is taken as the sample locations. If the y-axis translation is already done, the new data is assumed to be in this transformed unit.

Parameters `fname` (`str`) – The path to the file to use. If None, a QDialog is opened and we ask for a name

maybe_show_btn_reset_columns ()
Show the `btn_reset_columns` if the column starts are set

maybe_show_btn_reset_samples ()
Show the `btn_reset_samples` if the samples are set

modify_column_ends ()
Modify the column ends

After having selected the `column starts`, this method enables the modification of the column ends

See also:

<code>select_column_starts()</code> ,	<code>straditize.straditizer.Straditizer.</code>
<code>marks_for_column_ends()</code> ,	<code>straditize.straditizer.Straditizer.</code>
<code>update_column_ends()</code>	

new_reader_for_selection (`cls=None`)
Create a new child reader for the selected columns

This method finishes the process started by `enable_col_selection_for_new_reader()`

Parameters `cls` (`type`) – The subclass of the `straditize.binary.DataReader` class to use for the new reader. If None, a QDialog is opened and we ask for a reader

property reader
The `straditize.straditizer.Straditizer.data_reader`

refresh ()
Refresh from the straditizer

remove_hlines ()
Remove horizontal lines

This method uses the `straditize.binary.DataReader.recognize_hlines()` method to identify horizontal lines in the plot

remove_vlines ()
Remove vertical lines

This method uses the `straditize.binary.DataReader.recognize_vlines()` method to identify vertical lines in the plot

remove_xaxes ()
Remove x-axes in the plot

This method uses the `straditize.binary.DataReader.recognize_xaxes()` method to identify x-axes in the plot

remove_yaxes ()
Remove y-axes in the plot

This method uses the `straditiz.binary.DataReader.recognize_yaxes()` method to identify y-axes in the plot

reset_column_starts()

Reset the column starts

Reset the column starts by calling the `straditiz.binary.DataReader.reset_column_starts()` method

reset_samples()

Reset the samples

Reset the samples by calling the `straditiz.binary.DataReader.reset_samples()` method

select_column_starts()

Estimate the column starts and draw marks

This methods estimates the column starts (if they are not yet set) based on the threshold in the `txt_column_thresh` and draws `straditiz.cross_marks.DraggableVLine` marks on the plot.

See also:

`straditiz.straditizer.Straditizer.marks_for_column_starts()`,
`straditiz.straditizer.Straditizer.update_column_starts()`

select_data_part(`guess_lims=True`)

Enable the selection of the diagram part

This method uses the `straditiz.straditizer.Straditizer.marks_for_data_selection()` method to draw cross marks on the image for the diagram part

select_exaggerated_features()

Enable the selection of exaggerated features

select_occurrences()

Save (and potentially remove) the selected occurrences

Save the occurrences with the `straditiz.binary.DataReader.get_occurrences()` method and remove them if the `cb_remove_occurrences` is checked

property selection_toolbar**set_occurrences_value(`value`)**

Set the `occurrences_value`

Set the `straditiz.binary.DataReader.occurrence_value` of the `data_reader` with the given value

Parameters `value` (`float`) – The value to use for occurrences

setup_children(`item`)

Set up the child items for a `topLevelItem` in the control tree

should_be_enabled(`w`)

Check if a widget should be enabled

This function checks if a given widget `w` from the `widgets2disable` attribute should be enabled or not

Parameters `w` (`QWidget`) – The widget to check

Returns True, if the widget should be enabled

Return type `bool`

show_cross_column_features()
Remove cross column features
This method highlights features that span multiple columns using the `straditize.binary.DataReader.show_cross_column_features()` method. The algorithm can be modified with the `txt_cross_column_px` line editor

show_disconnected_parts()
Remove disconnected parts
This method uses the `straditize.binary.DataReader.show_disconnected_parts()` to highlight and remove disconnected features in the diagram part. The algorithm can be modified by the `txt_fromlast` and `txt_from0` text editors

show_parts_at_column_ends()
Remove parts that touch the column ends
This method highlights features that touch the column ends using the `straditize.binary.DataReader.show_parts_at_column_ends()` method.

show_small_parts()
Remove parts that touch the column ends
This method highlights small features in the data image using the `straditize.binary.DataReader.show_small_parts()` method. The maximal size of the small features can be taken from the `txt_max_small_size` line editor

sp_max_lw = None
A QSpinBox to select the maximum linewidth

sp_min_lw = None
A QSpinBox to select the minimum linewidth

sp_pixel_tol = None
A QSpinBox to set the minimum distance between samples in the sample finding algorithm

toggle_bar_split_source(*i*)
Fill the `tree_bar_split` based on the `cb_split_source`
Parameters *i* (`int`) – The `BarSplitter.fill_table()` is called with either 'too-long' (if *i* is 0), 'overlaps' (if *i* is 1) or 'all'

toggle_btn_highlight_small_selection()
Enable or disable the `btn_highlight_small_selection`
This method enables the `btn_highlight_small_selection` button if we are selecting something at the moment

toggle_sp_max_lw(*state*)
Toggle `sp_max_lw` based on `cb_max_lw`

toggle_txt_edit_rows(*state*)
Toggle `txt_edit_rows` based on `cb_edit_separate`

toggle_txt_from0(*state*)
Toggle `txt_from0` based on `cb_from0`

toggle_txt_fromlast(*state*)
Toggle `txt_fromlast` based on `cb_fromlast`

toggle_txt_tolerance(*s*)
Set the visibility of the `txt_tolerance` based on the reader

Parameters `s` (`str`) – The reader name. If there is `bars` in `s`, then the `txt_tolerance` is displayed

property tree
The `straditize.widgets.StraditzerWidgets.tree`

tree_bar_split = None
A `BarSplitter` to split too long bars

txt_column_thresh = None
A QLineEdit to set the threshold for the column starts detection

txt_cross_column_px = None
A QLineEdit to select the minimum pixels (`min_px`) for a cross column feature

txt_edit_rows = None
A QLineEdit to specify the number or rows in a plot for editing the samples in a separate figure (see `btn_edit_samples` and `cb_edit_separate`)

txt_exag_factor = None
A QLineEdit for the exaggeration factor

txt_from0 = None
A QLineEdit to set the `from0` keyword for the `straditize.binary.DataReader.show_disconnected_parts()` method

txt_fromlast = None
A QLineEdit to set the `fromlast` keyword for the `straditize.binary.DataReader.show_disconnected_parts()` method

txt_line_fraction = None
LineEdit for specifying the fraction of vertical and horizontal lines

txt_max_len = None
A QLineEdit to specify the maximum length of a potential sample to be included in the sample finding algorithm (see `btn_find_samples`)

txt_max_small_size = None
A QLineEdit to set the size for small parts

txt_min_highlight = None
A QLineEdit to set the maximal size for `highlighting small features`

txt_min_len = None
A QLineEdit to specify the minimum length of a potential sample to be included in the sample finding algorithm (see `btn_find_samples`)

txt_occurrences_value = None
A QLineEdit to set the value for occurrences in the final data

txt_tolerance = None
Line edit for setting the tolerance for bars

update_tolerance(s)
Set the readers tolerance

Parameters `s` (`str` or `int`) – The tolerance for the `straditzer.binary.BarDataReader.tolerance` attribute

`straditize.widgets.data.get_reader_name(reader)`
Get the reader key in the `straditize.binary.readers` dictionary

Parameters `reader` (`straditize.binary.DataReader`) – The reader for which to get the key in the `straditize.binary.readers` dictionary

Returns The key in the `straditize.binary.readers` dictionary whose value corresponds to the class of the given `reader`

Return type `str`

`straditize.widgets.data.int_list2str(numbers)`

Create a short string representation of an integer list

Parameters `numbers` (*list of ints*) – Integer list

Returns The string representation

Return type `str`

Examples

`[1, 2, 3]` becomes '`1-3`', `[1, 2, 3, 5, 7, 8, 9]` becomes '`1-3, 5, 7-9`'

straditize.widgets.image_correction module

Image correction methods

Disclaimer

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Classes

<code>ImageRescaler(straditizer_widgets, item, ...)</code>	A button to rescale the straditze image
--	---

<code>ImageRotator(straditizer_widgets[, item])</code>	Widget to rotate the image
--	----------------------------

class `straditize.widgets.image_correction.ImageRescaler(straditizer_widgets, item, *args, **kwargs)`

Bases: `straditize.widgets.StraditzerControlBase`, `PyQt5.QtWidgets.QPushButton`

A button to rescale the straditze image **Methods**

<code>adjust_orig_limits(*args, **kwargs)</code>	Readjust <code>ax_orig</code> after changes in <code>ax_rescale</code>
--	--

<code>adjust_rescaled_limits(*args, **kwargs)</code>	Readjust <code>ax_rescale</code> after changes in <code>ax_orig</code>
--	--

<code>close_figs()</code>	Close the fig
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<code>draw_figure()</code>	
----------------------------	--

<code>equalize_axes([event])</code>	Set both axes to the same size
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<code>raise_figure()</code>	Raise the figure for rescaling
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<code>rescale([ask])</code>	Rescale and start a new straditizer
<code>rescale_plot(percentage)</code>	Replot <code>im_rescale</code> after adjustments of the slider
<code>resize_stradi_image(percentage)</code>	Resize the straditizer image
<code>should_be_enabled(w)</code>	Check if a widget should be enabled
<code>start_rescaling()</code>	Create the rescaling figure

Attributes

<code>ax_orig</code>	The matplotlib axes for the <code>im_orig</code>
<code>ax_rescale</code>	The matplotlib axes for the <code>im_rescale</code>
<code>fig</code>	The matplotlib figure for the rescaling
<code>im_orig</code>	The matplotlib image for the original diagram
<code>im_rescale</code>	The matplotlib image for the rescaled diagram
<code>rescaling</code>	Boolean that is true if one of the axes is rescaling
<code>slider</code>	A <code>matplotlib.widgets.Slider</code> for specifying the size of the

`adjust_orig_limits(*args, **kwargs)`

Readjust `ax_orig` after changes in `ax_rescale`

`adjust_rescaled_limits(*args, **kwargs)`

Readjust `ax_rescale` after changes in `ax_orig`

`ax_orig = None`

The matplotlib axes for the `im_orig`

`ax_rescale = None`

The matplotlib axes for the `im_rescale`

`close_figs()`

Close the `fig`

`draw_figure()`

`equalize_axes(event=None)`

Set both axes to the same size

`fig = None`

The matplotlib figure for the rescaling

`im_orig = None`

The matplotlib image for the original diagram

`im_rescale = None`

The matplotlib image for the rescaled diagram

`raise_figure()`

Raise the figure for rescaling

`rescale(ask=None)`

Rescale and start a new straditizer

Parameters `ask (bool)` – Whether to ask with a QMessageBox. If None, it defaults to the `straditize.widgets.StraditzerWidgers.always_yes`

`rescale_plot(percentage)`

Replot `im_rescale` after adjustments of the `slider`

property rescaling

Boolean that is true if one of the axes is rescaling

resize_stradi_image(percentage)

Resize the straditizer image

Parameters `percentage (float)` – A float between 0 and 100 specifying the target size of the straditize.straditzer.Straditzer.image

Returns The resized image of the current straditzer

Return type `PIL.Image.Image`

should_be_enabled(w)

Check if a widget should be enabled

This function checks if a given widget `w` from the `widgets2disable` attribute should be enabled or not

Parameters `w (QWidget)` – The widget to check

Returns True, if the widget should be enabled

Return type `bool`

slider = None

A `matplotlib.widgets.Slider` for specifying the size of the rescaled image

start_rescaling()

Create the rescaling figure

```
class straditize.widgets.image_correction.ImageRotator(straditzer_widgets,
                                                       item=None,           *args,
                                                       **kwargs)
```

Bases: `straditize.widgets.StraditzerControlBase`, `PyQt5.QtWidgets.QWidget`

Widget to rotate the image

This control mainly adds a `QLineEdit` `txt_rotate` to the `straditize.widgets.StraditzerWidgets` to rotate the image. It also enables the user to specify the rotation angle using two connected `CrossMarks`. Here the user can decide between a horizontal alignment (`btn_rotate_horizontal`) or a vertical alignment (`btn_rotate_vertical`)

Parameters

- **straditzer_widgets** (`StraditzerWidgets`) – The main widget for the straditzer GUI
- **item** (`QTreeWidgetItem`) – The parent item in the `StraditzerWidgets.tree`. If given, the `setup_children()` is called with this item

Attributes

<code>angle</code>	The rotation angle from <code>txt_rotate</code> as a float
<code>btn_rotate_horizontal</code>	A QPushButton for horizontal alignment
<code>btn_rotate_vertical</code>	A QPushButton for vertical alignment
<code>txt_rotate</code>	A QLineEdit to display the rotation angle

Methods**draw_figure()**

Continued on next page

Table 74 – continued from previous page

<code>enable_or_disable_widgets(b)</code>	Enable or disable the widgets in this control
<code>remove_marks()</code>	Remove the cross marks used for the rotation angle
<code>rotate_image()</code>	Rotate the image based on the specified <code>angle</code>
<code>should_be_enabled(w)</code>	Check if a widget should be enabled
<code>start_horizontal_alignment()</code>	Start the horizontal alignment
<code>start_rotation()</code>	Start the rotation (if not already started)
<code>start_vertical_alignment()</code>	Start the vertical alignment
<code>update_txt_rotate(*args[, marks])</code>	Update the <code>txt_rotate</code> from the displayed cross marks

property angle

The rotation angle from `txt_rotate` as a float

btn_rotate_horizontal = None

A QPushButton for horizontal alignment

btn_rotate_vertical = None

A QPushButton for vertical alignment

draw_figure()**enable_or_disable_widgets (b)**

Enable or disable the widgets in this control

This method enables or disables the `widgets2disable` if the `should_be_enabled()` method evaluates to True

Parameters `b` (`bool`) – If True, enable the widgets, if False, disable them

remove_marks ()

Remove the cross marks used for the rotation angle

rotate_image ()

Rotate the image based on the specified `angle`

should_be_enabled (w)

Check if a widget should be enabled

This function checks if a given widget `w` from the `widgets2disable` attribute should be enabled or not

Parameters `w` (`QWidget`) – The widget to check

Returns True, if the widget should be enabled

Return type `bool`

start_horizontal_alignment ()

Start the horizontal alignment

start_rotation ()

Start the rotation (if not already started)

start_vertical_alignment ()

Start the vertical alignment

txt_rotate = None

A QLineEdit to display the rotation angle

update_txt_rotate (*args, marks=[], **kwargs)

Update the `txt_rotate` from the displayed cross marks

straditize.widgets.marker_control module

A widget for controlling the appearance of markers

This module defines the `MarkerControl` to control the appearance and behaviour of `CrossMarks` instances in the `straditizer.straditizer.Straditizer.marks` attribute

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Classes

<code>ColorLabel([color])</code>	A QTableWidget with one cell and no headers to just display a color
<code>MarkerControl(straditizer_widgets, item, ...)</code>	Widget to control the appearance of the marks

class straditize.widgets.marker_control.`ColorLabel` (`color='w'`, `*args`, `**kwargs`)

Bases: PyQt5.QtWidgets.QTableWidget

A QTableWidget with one cell and no headers to just display a color

The color to display

Parameters `color (object)` – Either a QtGui.QColor object or a color that can be converted to RGBA using the `matplotlib.colors.to_rgba()` function

Methods

<code>adjust_height()</code>	Adjust the height to match the row height
<code>select_color(*args)</code>	Select a color using PyQt5.QtWidgets.QColorDialog.getColor()
<code>set_color(color)</code>	Set the color of the label
<code>sizeHint()</code>	Reimplemented to use the rowHeight as height

Attributes

<code>color</code>	QtCore.QColor.
<code>color_changed(*args, **kwargs)</code>	a signal that is emitted with an rgba color if the chosen color changes

`adjust_height()`

Adjust the height to match the row height

`color = None`

QtCore.QColor. The current color that is displayed

color_changed(*args, **kwargs)
a signal that is emitted with an rgba color if the chosen color changes

select_color(*args)
Select a color using PyQt5.QtWidgets.QColorDialog.getColor()

set_color(color)
Set the color of the label

This method sets the given *color* as background color for the cell and emits the `color_changed` signal

Parameters `color`(*object*) – Either a QtGui.QColor object or a color that can be converted to RGBA using the `matplotlib.colors.to_rgba()` function

sizeHint()

Reimplemented to use the rowHeight as height

class straditize.widgets.marker_control.MarkerControl(straditizer_widgets, item, *args, **kwargs)

Bases: `straditize.widgets.StraditzerControlBase`, PyQt5.QtWidgets.QWidget

Widget to control the appearance of the marks

This widget controls the appearance of the `straditize.cross_mark.CrossMarks` instances in the `marks` attribute of the `straditizer`

Parameters

- `straditizer_widgets`(StraditzerWidgets) – The main widget for the straditzer GUI
- `item`(QTreeWidgetItem) – The parent item in the StraditzerWidgets.tree. If given, the `setup_children()` is called with this item

Methods

<code>add_toolbar_widgets(mark)</code>	Add the navigation actions to the toolbar
<code>change_auto_hide(auto_hide)</code>	Toggle the <code>auto_hide</code>
<code>change_hline_draggable(state)</code>	Enable or disable the dragging of horizontal lines
<code>change_hline_selectable(state)</code>	Enable or disable the selection of horizontal lines
<code>change_line_style(i)</code>	Change the line style of the marks
<code>change_line_widths(lw)</code>	Change the linewidth of the marks
<code>change_marker_size(markersize)</code>	Change the size of the markers
<code>change_marker_style(i)</code>	Change the marker style of the marks
<code>change_select_colors(color)</code>	Change the selection color of the marks
<code>change_selection_line_widths(lw)</code>	Change the linewidth for selected marks
<code>change_show_connected_artists(show)</code>	Change the visibility of connected artists
<code>change_show_hlines(state)</code>	Enable or disable the visibility of horizontal lines
<code>change_show_vlines(state)</code>	Enable or disable the visibility of vertical lines
<code>change_unselect_colors(color)</code>	Change the straditize.cross_mark.CrossMark.unselect color
<code>change_vline_draggable(state)</code>	Enable or disable the dragging of vertical lines
<code>change_vline_selectable(state)</code>	Enable or disable the selection of vertical lines
<code>draw_figs()</code>	Draw the figures of the marks
<code>enable_or_disable_widgets(b)</code>	Renabled to use the <code>refresh()</code> method
<code>fill_after_adding(mark)</code>	
<code>fill_from_mark(mark)</code>	Set the widgets of this <code>MarkerControl</code> from a mark

Continued on next page

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<code>fill_linestyles()</code>	Fill the <code>combo_line_style</code> combobox
<code>fill_markerstyles()</code>	Fill the <code>combo_marker_style</code> combobox
<code>go_to_greater_x_mark(x)</code>	Move the plot to the next mark with a x-position greater than <code>x</code>
<code>go_to_greater_y_mark(y)</code>	Move the plot to the next mark with a y-position greater than <code>y</code>
<code>go_to_left_mark()</code>	Move the plot to the previous left cross mark
<code>go_to_lower_mark()</code>	Go to the next mark below the current y-limits
<code>go_to_right_mark()</code>	Move the plot to the next right cross mark
<code>go_to_smaller_x_mark(x)</code>	Move the plot to the next mark with a x-position smaller than <code>x</code>
<code>go_to_smaller_y_mark(y)</code>	Move the plot to the next mark with a y-position smaller than <code>x</code>
<code>go_to_upper_mark()</code>	Go to the next mark above the current y-limits
<code>refresh()</code>	Reimplemented to also set the properties of this widget
<code>remove_actions()</code>	Remove the navigation actions from the toolbar
<code>set_line_style_item(ls)</code>	Switch the <code>combo_line_style</code> to the given linestyle
<code>set_marker_item(marker)</code>	Switch the <code>combo_marker_style</code> to the given marker
<code>should_be_enabled(w)</code>	Check if a widget <code>w</code> should be enabled or disabled
<code>update_mark(mark)</code>	Update the properties of a mark to match the settings

Attributes

<code>line_props</code>	The properties of the lines as a <code>dict</code>
<code>marks</code>	The <code>CrossMarks</code> of the straditizer
<code>select_props</code>	The properties of selected marks as a <code>dict</code>

`add_toolbar_widgets(mark)`

Add the navigation actions to the toolbar

`change_auto_hide(auto_hide)`

Toggle the `auto_hide`

This method disables or enables the `auto_hide` of the marks

Parameters `auto_hide` (`bool` or `PyQt5.QtGui.Qt.Checked` or `PyQt5.QtGui.Qt.Unchecked`) – The value to use for the `auto_hide`. `PyQt5.QtGui.Qt.Checked` is equivalent to `True`

`change_hline_draggable(state)`

Enable or disable the dragging of horizontal lines

Parameters `state` (`Qt.Checked` or `Qt.Unchecked`) – If `Qt.Checked`, the horizontal lines can be dragged and dropped

`change_hline_selectable(state)`

Enable or disable the selection of horizontal lines

Parameters `state` (`Qt.Checked` or `Qt.Unchecked`) – If `Qt.Checked`, the horizontal lines can be selected

`change_line_style(i)`

Change the line style of the marks

Parameters `i` (`int`) – The index of the line style in the `line_styles` attribute to use

change_line_widths (`lw`)

Change the linewidth of the marks

Parameters `lw` (`float`) – The line width to use

change_marker_size (`markersize`)

Change the size of the markers

Parameters `markersize` (`float`) – The size of the marker to use

change_marker_style (`i`)

Change the marker style of the marks

Parameters `i` (`int`) – The index of the marker style in the `marker_styles` attribute to use

change_select_colors (`color`)

Change the selection color of the marks

Change the selection color of the marks to the given color.

Parameters `color` (`PyQt5.QtGui.QColor` or a `matplotlib` color) – The color to use

change_selection_line_widths (`lw`)

Change the linewidth for selected marks

Parameters `lw` (`float`) – The linewidth for selected marks

change_show_connected_artists (`show`)

Change the visibility of connected artists

Parameters `show` (`bool`) – The visibility for the `straditiz.cross_mark.CrossMarks.set_connected_artists_visible()` method

change_show_hlines (`state`)

Enable or disable the visibility of horizontal lines

Parameters `state` (`Qt.Checked` or `Qt.Unchecked`) – If Qt.Checked, all horizontal lines are hidden

change_show_vlines (`state`)

Enable or disable the visibility of vertical lines

Parameters `state` (`Qt.Checked` or `Qt.Unchecked`) – If Qt.Checked, all vertical lines are hidden

change_unselect_colors (`color`)

Change the `straditiz.cross_mark.CrossMark.cunselect` color

Change the unselection color of the marks to the given color.

Parameters `color` (`PyQt5.QtGui.QColor` or a `matplotlib` color) – The color to use

change_vline_draggable (`state`)

Enable or disable the dragging of vertical lines

Parameters `state` (`Qt.Checked` or `Qt.Unchecked`) – If Qt.Checked, the vertical lines can be dragged and dropped

change_vline_selectable (`state`)

Enable or disable the selection of vertical lines

Parameters `state` (`Qt.Checked` or `Qt.Unchecked`) – If Qt.Checked, the vertical lines can be selected

draw_figs()

Draw the figures of the `marks`

enable_or_disable_widgets(b)

Renabled to use the `refresh()` method

fill_after_adding(mark)**fill_from_mark(mark)**

Set the widgets of this `MarkerControl` from a mark

This method sets the color labels, combo boxes, check boxes and text edits to match the properties of the given `mark`

Parameters `mark` (`straditize.cross_mark.CrossMark`) – The mark to use the attributes from

fill_linestyles()

Fill the `combo_line_style` combobox

fill_markerstyles()

Fill the `combo_marker_style` combobox

go_to_greater_x_mark(x)

Move the plot to the next mark with a x-position greater than `x`

Parameters `x` (`float`) – The reference x-position that shall be smaller than the new centered mark

go_to_greater_y_mark(y)

Move the plot to the next mark with a y-position greater than `y`

Parameters `y` (`float`) – The reference y-position that shall be smaller than the new centered mark

go_to_left_mark()

Move the plot to the previous left cross mark

go_to_lower_mark()

Go to the next mark below the current y-limits

go_to_right_mark()

Move the plot to the next right cross mark

go_to_smaller_x_mark(x)

Move the plot to the next mark with a x-position smaller than `x`

Parameters `x` (`float`) – The reference x-position that shall be greater than the new centered mark

go_to_smaller_y_mark(y)

Move the plot to the next mark with a y-position smaller than `y`

Parameters `y` (`float`) – The reference y-position that shall be greater than the new centered mark

go_to_upper_mark()

Go to the next mark above the current y-limits

property line_props

The properties of the lines as a `dict`

property marks

The `CrossMarks` of the straditizer

refresh()

Reimplemented to also set the properties of this widget

remove_actions()

Remove the navigation actions from the toolbar

property select_props

The properties of selected marks as a `dict`

set_line_style_item(ls)

Switch the `combo_line_style` to the given linestyle

Parameters `ls (str)` – The matplotlib linestyle string

set_marker_item(marker)

Switch the `combo_marker_style` to the given `marker`

Parameters `marker (str)` – A matplotlib marker string

should_be_enabled(w)

Check if a widget `w` should be enabled or disabled

Parameters `w (PyQt5.QtWidgets.QWidget)` – The widget to (potentially) enable

Returns True if the straditizer of this instance has marks. Otherwise False

Return type `bool`

update_mark(mark)

Update the properties of a mark to match the settings

Parameters `mark (straditize.cross_mark.CrossMarks)` – The mark to update

straditize.widgets.menu_actions module

The main control widget for a straditizer

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Classes

`ExportDfDialog(df, straditizer[, fname])`

A QDialog to export a `pandas.DataFrame` to Excel or CSV

`StraditizerMenuActions(straditizer_widgets)`

An object to control the main functionality of a Straditzer

class straditize.widgets.menu_actions.**ExportDfDialog**(*df*, *straditizer*, *fname=None*,
 **args, **kwargs*)

Bases: PyQt5.QtWidgets.QDialog

A QDialog to export a `pandas.DataFrame` to Excel or CSV

Parameters

- **df** (`pandas.DataFrame`) – The DataFrame to be exported
- **straditizer** (`straditize.straditizer.Straditizer`) – The source straditzer
- **fname** (`str`) – The file name to export to

Methods

`accept(self)`

`cancel()`

`export_df(parent, df, straditizer[, fname, ...])` Open a dialog for exporting a DataFrame

`get_open_file_name()` Ask the user for a filename for saving the data frame

`accept (self)`

`cancel ()`

classmethod `export_df(parent, df, straditizer, fname=None, exec_=True)`

Open a dialog for exporting a DataFrame

Parameters `parent` (`QWidget`) – The parent widget

`get_open_file_name ()`

Ask the user for a filename for saving the data frame

class straditize.widgets.menu_actions.**StraditizerMenuActions**(*straditizer_widgets*)

Bases: `straditize.widgets.StraditizerControlBase`

An object to control the main functionality of a Straditizer

This object is creates menu actions to load the straditizer **Attributes**

`all_actions`

`save_actions`, `data_actions` and
`text_actions`

`data_actions`

The QActions to save the exported DataFrames and
data images

`save_actions`

The QActions to save the straditizer, straditizer im-
age, etc.

`text_actions`

The QActions to save the column names images

`window_layout_action`

The action to

Methods

`create_sliders(stradi)`

Create sliders to navigate in the given axes

`export_final([fname])`

Export the final results

`export_full([fname])`

Export the full digitized data

`finish_loading(stradi)`

Finish the opening of a straditizer

`from_clipboard()`

Open a straditizer from an Image in the clipboard

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<code>import_binary_image([fname])</code>	Import the binary data reader image from an external file
<code>import_data_image([fname])</code>	Import the data reader image from an external file
<code>import_full_image([fname])</code>	Import the straditizer image from an external file
<code>import_text_image(fname)</code>	Import the column names reader image from an external file
<code>open_straditizer([fname])</code>	Open a straditizer from an image or project file
<code>refresh()</code>	Refresh from the straditizer
<code>save_data_image([fname])</code>	Save the binary image of the data reader
<code>save_full_image([fname])</code>	Save the image of the straditizer
<code>save_straditizer()</code>	Save the straditizer to a file
<code>save_straditizer_as([fname])</code>	Save the straditizer to a file
<code>save_text_image([fname])</code>	Save the image of the colnames reader
<code>set_ax_xlim(ax_ref)</code>	Define a function to update xlim from a given centered value
<code>set_ax_ylim(ax_ref)</code>	Define a function to update ylim from a given centered value
<code>set_stradi_in_console()</code>	Set the straditizer in the console of the GUI main-window
<code>setup_children(item)</code>	Setup the children for this control
<code>setup_menu_actions(main)</code>	Create the actions for the file menu
<code>setup_shortcuts(main)</code>	Setup the shortcuts when switched to the straditizer layout
<code>stack_zoom_window()</code>	Stack the magnifier image above the help explorer
<code>update_x_navigation_sliders(ax)</code>	Update the horizontal navigation slider for the given ax'
<code>update_y_navigation_sliders(ax)</code>	Update the vertical navigation slider for the given ax'

property all_actions`save_actions, data_actions and text_actions`**create_sliders (stradi)**

Create sliders to navigate in the given axes

Parameters `stradi` (`straditize.straditizer.Straditizer`) – The straditizer that just has been opened

data_actions = []

The QActions to save the exported DataFrames and data images

export_final (fname=None)

Export the final results

This method exports the `straditize.straditizer.Straditizer.final_df` of the current `straditizer`

Parameters `fname` (`str or None`) – The path of the target filename where to save the `df` (see the `ExportDialog.export_df()`)

export_full (fname=None)

Export the full digitized data

This method exports the `straditize.straditizer.Straditizer.full_df` of the current `straditizer`

Parameters `fname` (`str or None`) – The path of the target filename where to save the `df` (see the `ExportDialog.export_df()`)

finish_loading(stradi)

Finish the opening of a straditizer

This method sets the `straditizer.widgets.StraditizerWidgets.straditizer`, shows the straditizer image, creates the navigation sliders and sets the straditizer in the console

Parameters `stradi` (`straditize.straditizer.Straditizer`) – The straditizer that just has been opened

from_clipboard()

Open a straditizer from an Image in the clipboard

This method uses the `PIL.ImageGrab.grabclipboard()` function to open a new straditizer from the clipboard.

import_binary_image(fname=None)

Import the binary data reader image from an external file

This method imports the `straditize.binary.DataReader.binary` from an external file and sets it to the current `data_reader`.

Parameters `fname` (`str, PIL.Image.Image` or `None`) – The path of the image file or the `PIL.Image.Image`. If `None`, a `QFileDialog` is opened to request the file from the user.

import_data_image(fname=None)

Import the data reader image from an external file

This method imports the `straditize.binary.DataReader.image` from an external file and sets it to the current `data_reader`.

Parameters `fname` (`str, PIL.Image.Image` or `None`) – The path of the image file or the `PIL.Image.Image`. If `None`, a `QFileDialog` is opened to request the file from the user.

See also:

[`import_binary_image\(\)`](#) To import the binary file

import_full_image(fname=None)

Import the straditizer image from an external file

This method imports the `straditize.straditizer.Straditizer.image` from an external file and sets it to the current straditizer.

Parameters `fname` (`str, PIL.Image.Image` or `None`) – The path of the image file or the `PIL.Image.Image`. If `None`, a `QFileDialog` is opened to request the file from the user.

import_text_image(fname)

Import the column names reader image from an external file

This method imports the `straditize.colnames.ColNamesReader.highres_image` from an external file and sets it to the current `colnames_reader`.

Parameters `fname` (`str, PIL.Image.Image` or `None`) – The path of the image file or the `PIL.Image.Image`. If `None`, a `QFileDialog` is opened to request the file from the user.

open_straditizer(fname=None, *args, **kwargs)

Open a straditizer from an image or project file

Parameters `fname` (`str, PIL.Image.Image` or `None`) – The path to the file to import. If `None`, a `QFileDialog` is opened and the user is asked for a file name. The action then depends on the ending of `fname`:

'`.nc`' or '`.nc4`' we expect a netCDF file and open it with `xarray.open_dataset()` and load the straditizer with the `straditize.straditizer.Straditizer.from_dataset()` constructor

'`.pk1`' We expect a pickle file and load the straditizer with `pickle.load()`

any other ending We expect an image file and use the `PIL.Image.open()` function

At the end, the loading is finished with the `finish_loading()` method

refresh()
Refresh from the straditizer

save_actions = []
The QActions to save the straditizer, straditizer image, etc.

save_data_image(fname=None)
Save the binary image of the data reader
Save the `straditize.binary.DataReader.binary` of the current `data_reader`
Parameters %(`StraditizerMenuActions._save_image.parameters.fname`) s

save_full_image(fname=None)
Save the image of the straditizer
Save the `straditize.straditizer.Straditizer.image` of the current `straditizer`
Parameters %(`StraditizerMenuActions._save_image.parameters.fname`) s

save_straditizer()
Save the straditizer to a file

save_straditizer_as(fname=None)
Save the straditizer to a file
Parameters `fname` (str or None) – If None, a QDialog is opened and the user has to provide a filename. The final action then depends on the ending of the chose file:
'`.pk1`' We save the straditizer with `pickle.dump()`
`else` We use the `straditize.straditizer.Straditizer.to_dataset()` method and save the resulting dataset using the `xarray.Dataset.to_netcdf()` method

save_text_image(fname=None)
Save the image of the colnames reader
Save the `straditize.colnames.ColNamesReader.image` of the current `colnames_reader`
Parameters `fname` (str or None) – The path of the target filename where to save the `image`. If None, A QDialog is opened and we ask the user for a filename

static set_ax_xlim(ax_ref)
Define a function to update xlim from a given centered value
Parameters `ax_ref` (`matplotlib.axes.Axes`) – The axes whose x-limits to update when the returned function is called
Returns The function that can be called with a `val` to set the x-center of the `ax_ref`
Return type callable

static set_ax_ylim(ax_ref)

Define a function to update ylim from a given centered value

Parameters `ax_ref (matplotlib.axes.Axes)` – The axes whose y-limits to update when the returned function is called

Returns The function that can be called with a *val* to set the y-center of the *ax_ref*

Return type callable

set_stradi_in_console()

Set the straditizer in the console of the GUI mainwindow

This sets the current `straditzer` in psyplots `console` as the `stradi` variable

setup_children(item)

Setup the children for this control

This method is called to setup the children in the `StraditizerWidgets.tree`. By default, it just creates a child `QTreeWidgetItem` and sets this control as its widget

Parameters `item (QTreeWidgetItem)` – The top level item in the `StraditizerWidgets.tree`

setup_menu_actions(main)

Create the actions for the file menu

Parameters `main (psyplot_gui.main.MainWindow)` – The mainwindow whose menubar shall be adapted

setup_shortcuts(main)

Setup the shortcuts when switched to the straditizer layout

Parameters `main (psyplot_gui.main.MainWindow)` – The psyplot mainwindow

stack_zoom_window()

Stack the magnifier image above the help explorer

text_actions = []

The QActions to save the column names images

static update_x_navigation_sliders(ax)

Update the horizontal navigation slider for the given *ax*‘

Set the horizontal slider of the straditizer figure depending on the x-limits of its corresponding axes

Parameters `ax (matplotlib.axes.Axes)` – The `straditize.straditizer.Straditizer.ax` attribute of a straditizer

static update_y_navigation_sliders(ax)

Update the vertical navigation slider for the given *ax*‘

Set the vertical slider of the straditizer figure depending on the y-limits of its corresponding axes

Parameters `ax (matplotlib.axes.Axes)` – The `straditize.straditizer.Straditizer.ax` attribute of a straditizer

window_layout_action = None

The action to `switch_to_straditizer_layout()`

straditize.widgets.pattern_selection module

A wdiget to select patterns in the image

The `PatternSelectionWidget` is used by the `straditize.widget.selection_toolbar.SelectionToolbar` to select patterns in the straditizer image

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Classes

<code>EmbededMplCanvas([parent])</code>	Ultimately, this is a QWidget (as well as a FigureCanvasAgg, etc.).
<code>PatternSelectionWidget(arr, data_obj[, ...])</code>	A widget to select patterns in the image

```
class straditize.widgets.pattern_selection.EmbededMplCanvas (parent=None, *args,
                                                               **kwargs)
Bases: matplotlib.backends.backend_qt5agg.FigureCanvasQTAgg
Ultimately, this is a QWidget (as well as a FigureCanvasAgg, etc.).

class straditize.widgets.pattern_selection.PatternSelectionWidget (arr,
                                                               data_obj,
                                                               re-
                                                               move_selection=False,
                                                               *args,
                                                               **kwargs)
Bases: PyQt5.QtWidgets.QWidget, psyplot_gui.common.DockMixin
A widget to select patterns in the image
```

This widget consist of an `EmbededMplCanvas` to display the template for the pattern and uses the `skimage.feature.match_template()` function to identify it in the `arr`

See also:

`straditize.widget.selection_toolbar.SelectionToolbar.start_pattern_selection`

Attributes

<code>selector</code>	The selector to select the template in the original image
<code>sl_thresh</code>	A QSlider to set the threshold for the template correlation
<code>template</code>	The template to look for in the <code>arr</code>
<code>template_extents</code>	The extents of the template in the original image
<code>template_fig</code>	The <code>EmbededMplCanvas</code> to display the template
<code>template_im</code>	The matplotlib artist of the template in the

Methods

<code>cancel()</code>	
<code>correlate_template(arr, template[, ...])</code>	Correlate a template with the <code>arr</code>
<code>maybe_tabify()</code>	
<code>modify_selection(i)</code>	Modify the selection based on the correlation threshold
<code>remove_plugin()</code>	Remove this plugin and close it
<code>start_correlation()</code>	Look for the correlations of template and source
<code>to_dock(main[, title, position, docktype])</code>	
<code>toggle_correlation_plot()</code>	Toggle the correlation plot between template and arr
<code>toggle_selection()</code>	Modify the selection (or not) based on the template correlation
<code>toggle_template_selection()</code>	Enable or disable the template selection
<code>update_image(*args, **kwargs)</code>	Update the template image based on the selector extents

Parameters

- `arr` (np.ndarray of shape (Ny, Nx)) – The labeled selection array
- `data_obj` (straditize.label_selection.LabelSelection) – The data object whose image shall be selected
- `remove_selection` (`bool`) – If True, remove the selection on apply

`axes = None`

`cancel()`

`correlate_template(arr, template, fraction=False, increment=1, report=True)`

Correlate a template with the `arr`

This method uses the `skimage.feature.match_template()` function to find the given `template` in the source array `arr`.

Parameters

- `arr` (np.ndarray of shape (Ny, Nx)) – The labeled selection array (see `arr`), the source of the given `template`
- `template` (np.ndarray of shape (nx, ny)) – The template from `arr` that shall be searched
- `fraction` (`float`) – If not null, we will look through the given fraction of the template to look for partial matches as well
- `increment` (`int`) – The increment of the loop with the `fraction`.
- `report` (`bool`) – If True and `fraction` is not null, a QProgressDialog is opened to inform the user about the progress

`key_press_cid = None`

`maybe_tabify()`

`modify_selection(i)`

Modify the selection based on the correlation threshold

Parameters `i` (`int`) – An integer between 0 and 100, the value of the `sl_thresh` slider

```
remove_plugin()
    Remove this plugin and close it

selector = None
    The selector to select the template in the original image

sl_thresh = None
    A QSlider to set the threshold for the template correlation

start_correlation()
    Look for the correlations of template and source

template = None
    The template to look for in the arr

template_extents = None
    The extents of the template in the original image

template_fig = None
    The EmbededMplCanvas to display the template

template_im = None
    The matplotlib artist of the template in the template_fig

to_dock(main, title=None, position=None, docktype='df', *args, **kwargs)
toggle_correlation_plot()
    Toggle the correlation plot between template and arr

toggle_selection()
    Modify the selection (or not) based on the template correlation

toggle_template_selection()
    Enable or disable the template selection

update_image(*args, **kwargs)
    Update the template image based on the selector extents
```

straditize.widgets.plots module

Plot control widgets for straditize

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Classes

<i>PlotControl</i> (straditzer_widgets, item, ...)	A widget for controlling the plot
<i>PlotControlTable</i> (straditzer_widgets, *args, ...)	A widget to control the plots
<i>ResultsPlot</i> (straditzer_widgets)	A widget for plotting the final results

```
class straditize.widgets.plots.PlotControl(straditizer_widgets, item, *args, **kwargs)
Bases: straditize.widgets.StraditzerControlBase, PyQt5.QtWidgets.QWidget
```

A widget for controlling the plot

This widgets holds a *PlotControlTable* to display visual diagnostics in the plot. Additionally it contains zoom buttons (*btn_view_global* and *btn_view_data*) and a widget to plot the results (*results_plot*)

Attributes

<i>btn_view_data</i>	A button to zoom to the data
<i>btn_view_global</i>	A button to zoom out to the entire stratigraphic diagram
<i>results_plot</i>	A <i>ResultsPlot</i> to plot the digitized data in a new diagram
<i>table</i>	A <i>PlotControlTable</i> to display visual diagnostics

Methods

<i>refresh()</i>	Refresh from the straditizer
<i>setup_children(item)</i>	Setup the children for this control
<i>zoom_data()</i>	Zoom to the data part
<i>zoom_global()</i>	Zoom out to the full straditgraphic diagram

***btn_view_data* = None**

A button to zoom to the data (see *zoom_data()*)

***btn_view_global* = None**

A button to zoom out to the entire stratigraphic diagram (see *zoom_global()*)

refresh()

Refresh from the straditizer

***results_plot* = None**

A *ResultsPlot* to plot the digitized data in a new diagram

setup_children(item)

Setup the children for this control

This method is called to setup the children in the *StraditzerWidgets.tree*. By default, it just creates a child QTreeWidgetItem and sets this control as it's widget

Parameters ***item*** (*QTreeWidgetItem*) – The top level item in the *StraditzerWidgets.tree*

***table* = None**

A *PlotControlTable* to display visual diagnostics

zoom_data()

Zoom to the data part

See also:

`straditize.straditzer.Straditzer.show_data_diagram()`

zoom_global()

Zoom out to the full straditgraphic diagram

See also:

```
    straditize.straditizer.Straditzer.show_full_image()

class straditize.widgets.plots.PlotControlTable(straditzer_widgets, *args, **kwargs)
    Bases: straditize.widgets.StraditzerControlBase, PyQt5.QtWidgets.QTableWidget
```

A widget to control the plots

This control widget is a table to plot, remove and toggle the visibility of visual diagnostics for the straditzer. It has two columns: the first to toggle the visibility of the plot, the second to plot and remove the matplotlib artists. The vertical header are the items in the corresponding `plot_funcs`, `can_be_plotted_funcs` and/or `hide_funcs`.

Rows are added to this table using the `add_item()` method which stores the plotting functions in the `plot_funcs` and the functions to hide the plot in the `hide_funcs`. Whether an item can be plotted or not depends on the results of the corresponding callable in the `can_be_plotted_funcs`. **Methods**

<code>add_item(what, get_artists[, plot_func, ...])</code>	Add a plot object to the table
<code>adjust_height()</code>	
<code>can_plot_column_starts()</code>	Test whether the column starts can be visualized
<code>can_plot_data_box()</code>	Test whether the box around the diagram part can be plotted
<code>can_plot_data_reader_color_image()</code>	Test if the reader color image can be plotted
<code>can_plot_full_df()</code>	Test whether the full_df can be plotted
<code>can_plot_potential_samples()</code>	Test whether potential sample regions can be plotted
<code>can_plot_sample_hlines()</code>	Test whether the sample lines can be plotted
<code>can_plot_samples()</code>	Test whether the samples can be plotted
<code>draw_figs(artists)</code>	Draw the figures of the given <i>artists</i>
<code>enable_or_disable_widgets(b)</code>	b is ignored and is always set to True
<code>get_column_start_lines()</code>	Get the artists of the column starts
<code>get_data_box()</code>	Get the plotted straditize.straditzer.Straditzer.data_box
<code>get_data_reader_background()</code>	Get the straditize.binary.DataReader.background
<code>get_data_reader_color_image()</code>	Get the straditize.binary.DataReader.color_plot_im
<code>get_data_reader_image()</code>	Get the straditize.binary.DataReader.plot_im
<code>get_full_df_lines()</code>	Get the artists of the full_df plot
<code>get_potential_samples_lines()</code>	Get the artists of the plot of potential samples
<code>get_sample_hlines()</code>	Get the plotted the sample lines
<code>get_samples_lines()</code>	Get the artists of the plotted samples
<code>get_straditzer_image()</code>	Get the straditize.straditzer.Straditzer.plot_im
<code>plot_column_starts()</code>	Plot horizontal lines for the column starts
<code>plot_data_box()</code>	Plot the data box around the diagram part
<code>plot_data_reader_color_image()</code>	Plot the data reader color image
<code>plot_full_df()</code>	Plot the <code>full_df</code> of the reader
<code>plot_potential_samples()</code>	Highlight the regions with potential samples in the plot
<code>plot_sample_hlines()</code>	Plot the horizontal sample lines of the reader
<code>plot_samples()</code>	Plot the samples of the reader
<code>refresh()</code>	Refresh from the straditzer
<code>remove_column_starts()</code>	Remove the plotted lines of the column starts

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<code>remove_data_box()</code>	Remove the box around the diagram part
<code>remove_data_reader_color_image()</code>	Remove the straditize.binary. DataReader.color_plot_im
<code>remove_full_df_plot()</code>	Remove the plot of the full_df
<code>remove_potential_samples_plot()</code>	Remove the plot of potential samples
<code>remove_sample_hlines_plot()</code>	Remove the sample lines
<code>remove_samples_plot()</code>	Remove the plotted samples
<code>should_be_enabled(w)</code>	Check if a widget should be enabled
<code>sizeHint(self)</code>	

Attributes

<code>can_be_plotted_funcs</code>	A mapping from plot identifier to a callable that returns True if
<code>col_lines</code>	Built-in mutable sequence.
<code>hide_funcs</code>	A mapping from plot identifier to a callable to hide the corresponding
<code>plot_funcs</code>	A mapping from plot identifier to a callable to plot the corresponding
<code>widgets2disable</code>	Built-in mutable sequence.

`add_item(what, get_artists, plot_func=None, remove_func=None, can_be_plotted=None)`

Add a plot object to the table

Parameters

- `what` (`str`) – The description of the plot object
- `get_artists` (`function`) – A function that takes no arguments and returns the artists
- `plot_func` (`function, optional`) – A function that takes no arguments and makes the plot.
- `remove_func` (`function, optional`) – A function that takes no arguments and removes the plot.
- `can_be_plotted` (`function, optional`) – A function that takes no argument and returns True if the plot can be made.

`adjust_height()`

`can_be_plotted_funcs = {}`

A mapping from plot identifier to a callable that returns True if the corresponding function in the `plot_funcs` mapping can be called

`can_plot_column_starts()`

Test whether the column starts can be visualized

See also:

`plot_column_starts()`

`can_plot_data_box()`

Test whether the box around the diagram part can be plotted

See also:

`plot_data_box()`

can_plot_data_reader_color_image()
Test if the reader color image can be plotted

See also:

plot_data_reader_color_image()

can_plot_full_df()
Test whether the full_df can be plotted

See also:

plot_full_df()

can_plot_potential_samples()
Test whether potential sample regions can be plotted

See also:

plot_potential_samples()

can_plot_sample_hlines()
Test whether the sample lines can be plotted

See also:

plot_sample_hlines()

can_plot_samples()
Test whether the samples can be plotted

See also:

plot_samples()

col_lines = []

draw_figs(artists)
Draw the figures of the given *artists*

Parameters **artists** (list of `matplotlib.artist.Artist`) – The artists to draw the canvas from

enable_or_disable_widgets(b)
b is ignored and is always set to True

get_column_start_lines()
Get the artists of the column starts

See also:

plot_column_starts()

get_data_box()
Get the plotted `straditize.straditizer.Straditizer.data_box`

See also:

plot_data_box()

get_data_reader_background()
Get the `straditize.binary.DataReader.background`

get_data_reader_color_image()
Get the `straditize.binary.DataReader.color_plot_im`

See also:

```
plot_data_reader_color_image()  
get_data_reader_image()  
    Get the straditize.binary.DataReader.plot_im  
get_full_df_lines()  
    Get the artists of the full_df plot  
  
    See also:  
        plot_full_df()  
  
get_potential_samples_lines()  
    Get the artists of the plot of potential samples  
  
    See also:  
        plot_potential_samples()  
  
get_sample_hlines()  
    Get the plotted the sample lines  
  
    See also:  
        plot_sample_hlines()  
  
get_samples_lines()  
    Get the artists of the plotted samples  
  
    See also:  
        plot_samples()  
  
get_straditizer_image()  
    Get the straditize.straditizer.Straditizer.plot_im  
  
hide_funcs = {}  
A mapping from plot identifier to a callable to hide the corresponding artists  
  
plot_column_starts()  
Plot horizontal lines for the column starts  
  
See also:  
    remove_column_starts(), get_column_start_lines(), can_plot_column_starts()  
  
plot_data_box()  
Plot the data box around the diagram part  
  
See also:  
    straditize.straditizer.Straditizer.draw_data_box(), remove_data_box(),  
    get_data_box(), can_plot_data_box()  
  
plot_data_reader_color_image()  
Plot the data reader color image  
  
See also:  
    straditize.binary.DataReader.plot_color_image(), remove_data_reader_color_image(),  
    can_plot_data_reader_color_image()  
  
plot_full_df()  
Plot the full_df of the reader  
  
See also:
```

```
straditiz.binary.DataReader.plot_full_df(),      remove_full_df_plot(),
get_full_df_lines(), can_plot_full_df()
```

plot_funcs = {}

A mapping from plot identifier to a callable to plot the corresponding artists

plot_potential_samples()

Highlight the regions with potential samples in the plot

See also:

```
straditiz.binary.DataReader.plot_potential_samples(),
remove_potential_samples_plot(),           get_potential_samples_lines(),
can_plot_potential_samples()
```

plot_sample_hlines()

Plot the horizontal sample lines of the reader

See also:

```
straditiz.binary.DataReader.plot_sample_hlines(),
remove_sample_hlines_plot(), get_sample_hlines(), can_plot_sample_hlines()
```

plot_samples()

Plot the samples of the reader

See also:

```
straditiz.binary.DataReader.plot_samples(),      remove_samples_plot(),
get_samples_lines(), can_plot_samples()
```

refresh()

Refresh from the straditizer

remove_column_starts()

Remove the plotted lines of the column starts

See also:

```
plot_column_starts()
```

remove_data_box()

Remove the box around the diagram part

See also:

```
plot_data_box()
```

remove_data_reader_color_image()

Remove the straditiz.binary.DataReader.color_plot_im

See also:

```
plot_data_reader_color_image()
```

remove_full_df_plot()

Remove the plot of the full_df

See also:

```
plot_full_df()
```

remove_potential_samples_plot()

Remove the plot of potential samples

See also:

`plot_potential_samples()`
`remove_sample_hlines_plot()`
 Remove the sample lines

See also:

`plot_sample_hlines()`
`remove_samples_plot()`
 Remove the plotted samples

See also:

`plot_samples()`
`should_be_enabled(w)`
 Check if a widget should be enabled

This function checks if a given widget `w` from the `widgets2disable` attribute should be enabled or not

Parameters `w` (`QWidget`) – The widget to check

Returns True, if the widget should be enabled

Return type `bool`

`sizeHint(self) → QSize`

property `widgets2disable`
 Built-in mutable sequence.

If no argument is given, the constructor creates a new empty list. The argument must be an iterable if specified.

class straditize.widgets.plots.`ResultsPlot` (straditzer_widgets)
 Bases: `straditize.widgets.StraditzerControlBase`

A widget for plotting the final results

This widgets contains a QPushButton `btn_plot` to plot the results using the `straditize.binary.DataReader.plot_results()` method **Attributes**

<code>btn_plot</code>	The QPushButton to call the <code>plot_results()</code> method
<code>cb_final</code>	A QCheckBox whether the samples or the full digitized data shall be
<code>cb_transformed</code>	A QCheckBox whether x- and y-axis should be translated from pixel to

Methods

<code>plot_results()</code>	Plot the results
<code>refresh()</code>	Refresh from the straditzer
<code>setup_children(item)</code>	Setup the children for this control

`btn_plot = None`
 The QPushButton to call the `plot_results()` method

`cb_final = None`
 A QCheckBox whether the samples or the full digitized data shall be plotted

```
cb_transformed = None
A QCheckBox whether x- and y-axis should be translated from pixel to data units

plot_results()
Plot the results

What is plotted depends on the cb_transformed and the cb_final

cb_transformed and cb_final are checked Plot the straditize.straditizer.Straditizer.final_df

cb_transformed is checked but not cb_final Plot the straditize.straditizer.Straditizer.full_df

cb_transformed is not checked but cb_final Plot the straditize.binary.DataReader.sample_locs

cb_transformed and cb_final are both not checked Plot the straditize.binary.DataReader.full_df

refresh()
Refresh from the straditzer

setup_children(item)
Setup the children for this control

This method is called to setup the children in the StraditizerWidgets.tree. By default, it just creates a child QTreeWidgetItem and sets this control as its widget

Parameters item (QTreeWidgetItem) – The top level item in the StraditizerWidgets.tree
```

straditize.widgets.progress_widget module

Progress widget for the straditzation

The ProgressWidget defined here is a ListWidget to show the current state of the straditization.

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Classes

<code>ColumnNamesTask</code> (parent)	Task to specify the column names
<code>ColumnsTask</code> (parent)	A task to separate columns
<code>DataLimitsTask</code> (parent)	The task to check the data limits
<code>DataReaderTask</code> (parent)	A task for initializing the reader
<code>DigitizeTask</code> (parent)	Task to digitize the data

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<code>ExportTask</code> (parent)	Task to export the final results
<code>InitStraditizerTask</code> (parent)	The task to initialize a straditizer
<code>OccurrencesTask</code> (parent)	Task to handle occurrences
<code>ProgressTask</code> (parent)	The base class for an item that should be shown in the ProgressWidget
<code>ProgressWidget</code> (straditizer_widgets, item)	A widget to show the progress of the straditization
<code>RemoveArtifactsTask</code> (parent)	Task to clean the binary image
<code>RemoveLinesTask</code> (parent)	Task to remove y-axes, x-axes, horizontal and vertical lines
<code>SamplesTask</code> (parent)	Task to edit and find samples
<code>SaveProjectTask</code> (*args, **kwargs)	Task to remember saving the project
<code>SelectExaggerationsTask</code> (parent)	Task to handle exaggerations
<code>XTranslationTask</code> (parent)	Task to specify the x-axes conversion from pixel to data units
<code>YTranslationTask</code> (parent)	Task to specify the y-axis conversion from pixel to data units

```
class straditize.widgets.progress_widget.ColumnNamesTask (parent)
```

Bases: `straditize.widgets.progress_widget.ProgressTask`

Task to specify the column names

Parameters `parent` (`QListWidget`) – The list that holds this item

Attributes

<code>dependencies</code>	Built-in mutable sequence.
<code>done_tooltip</code>	
<code>is_finished</code>	A property that is True, when the task is finished
<code>name</code>	<code>str(object=") -> str</code>
<code>rst_file</code>	<code>str(object=") -> str</code>
<code>summary</code>	<code>str(object=") -> str</code>
<code>task_tooltip</code>	<code>str(object=") -> str</code>

```
dependencies = ['columns']
```

```
property done_tooltip
```

```
property is_finished
```

A property that is True, when the task is finished

```
name = 'column_names'
```

```
rst_file = 'column_names'
```

```
summary = 'Specify the column names'
```

```
task_tooltip = 'Click the <i>Edit column names</i> button to insert the names for each
```

```
class straditize.widgets.progress_widget.ColumnsTask (parent)
```

Bases: `straditize.widgets.progress_widget.ProgressTask`

A task to separate columns

Parameters `parent` (`QListWidget`) – The list that holds this item

Attributes

<code>dependencies</code>	Built-in mutable sequence.
<code>done_tooltip</code>	
<code>is_finished</code>	A property that is True, when the task is finished
<code>name</code>	<code>str(object=") -> str</code>
<code>rst_file</code>	<code>str(object=") -> str</code>
<code>summary</code>	<code>str(object=") -> str</code>
<code>task_tooltip</code>	<code>str(object=") -> str</code>

```
dependencies = ['init_reader']

property done_tooltip

property is_finished
    A property that is True, when the task is finished

name = 'columns'

rst_file = 'select_column_starts'

summary = 'Separate the columns'

task_tooltip = 'Separate the columns (subdiagrams) of the stratigraphic diagram'

class straditize.widgets.progress_widget.DataLimitsTask(parent)
Bases: straditize.widgets.progress_widget.ProgressTask
```

The task to check the data limits

Parameters `parent` (`QListWidget`) – The list that holds this item

Attributes

<code>dependencies</code>	Built-in mutable sequence.
<code>done_tooltip</code>	
<code>is_finished</code>	A property that is True, when the task is finished
<code>name</code>	<code>str(object=") -> str</code>
<code>rst_file</code>	<code>str(object=") -> str</code>
<code>summary</code>	<code>str(object=") -> str</code>
<code>task_tooltip</code>	<code>str(object=") -> str</code>

```
dependencies = ['init_stradi']

property done_tooltip

property is_finished
    A property that is True, when the task is finished

name = 'datalim'

rst_file = 'select_data_part'

summary = 'Limits of the diagram'

task_tooltip = 'Specify the corners for the data part of the diagram by clicking the <

class straditize.widgets.progress_widget.DataReaderTask(parent)
Bases: straditize.widgets.progress_widget.ProgressTask
```

A task for initializing the reader

Parameters `parent` (`QListWidget`) – The list that holds this item

Attributes

<code>dependencies</code>	Built-in mutable sequence.
<code>done_tooltip</code>	
<code>is_finished</code>	A property that is True, when the task is finished
<code>name</code>	<code>str(object=") -> str</code>
<code>rst_file</code>	<code>str(object=") -> str</code>
<code>summary</code>	<code>str(object=") -> str</code>
<code>task_tooltip</code>	<code>str(object=") -> str</code>

```

dependencies = ['datalim']

property done_tooltip
property is_finished
    A property that is True, when the task is finished
name = 'init_reader'

rst_file = 'select_reader'

summary = 'Initialize the diagram reader'

task_tooltip = 'Choose the appropriate reader type and click the <i>Convert image</i> button'

class straditize.widgets.progress_widget.DigitizeTask(parent)
Bases: straditize.widgets.progress_widget.ProgressTask

```

Task to digitize the data

Parameters `parent` (`QListWidget`) – The list that holds this item

Attributes

<code>dependencies</code>	Built-in mutable sequence.
<code>done_tooltip</code>	
<code>is_finished</code>	A property that is True, when the task is finished
<code>name</code>	<code>str(object=") -> str</code>
<code>rst_file</code>	<code>str(object=") -> str</code>
<code>summary</code>	<code>str(object=") -> str</code>
<code>task_tooltip</code>	<code>str(object=") -> str</code>

```

dependencies = ['remove_artifacts', 'remove_lines']

property done_tooltip
property is_finished
    A property that is True, when the task is finished
name = 'digitize'

rst_file = 'digitize'

summary = 'Digitize the diagram'

task_tooltip = 'Click the `Digitize` button to digitize the diagram'

class straditize.widgets.progress_widget.ExportTask(parent)
Bases: straditize.widgets.progress_widget.ProgressTask

```

Task to export the final results

Parameters `parent` (`QListWidget`) – The list that holds this item

Attributes

<code>dependencies</code>	Built-in mutable sequence.
<code>done_tooltip</code>	
<code>is_finished</code>	A property that is True, when the task is finished
<code>name</code>	<code>str(object=") -> str</code>
<code>rst_file</code>	<code>str(object=") -> str</code>
<code>summary</code>	<code>str(object=") -> str</code>
<code>task_tooltip</code>	<code>str(object=") -> str</code>

```
dependencies = ['samples']
```

```
property done_tooltip
```

```
property is_finished
```

A property that is True, when the task is finished

```
name = 'export'
```

```
rst_file = 'export'
```

```
summary = 'Export the final data'
```

```
task_tooltip = 'Export the final data'
```

```
class straditize.widgets.progress_widget.InitStraditizerTask(parent)
```

Bases: `straditize.widgets.progress_widget.ProgressTask`

The task to initialize a straditzer

Parameters `parent` (`QListWidget`) – The list that holds this item

Attributes

<code>done_tooltip</code>	
<code>is_finished</code>	A property that is True, when the task is finished
<code>name</code>	<code>str(object=") -> str</code>
<code>rst_file</code>	<code>str(object=") -> str</code>
<code>summary</code>	<code>str(object=") -> str</code>
<code>task_tooltip</code>	<code>str(object=") -> str</code>

```
property done_tooltip
```

```
property is_finished
```

A property that is True, when the task is finished

```
name = 'init_stradi'
```

```
rst_file = 'load_image'
```

```
summary = 'Load an image or project'
```

```
task_tooltip = 'Load a straditzer image or project to get started'
```

```
class straditize.widgets.progress_widget.OccurrencesTask(parent)
```

Bases: `straditize.widgets.progress_widget.ProgressTask`

Task to handle occurences

Parameters `parent` (`QListWidget`) – The list that holds this item

Attributes

<code>dependencies</code>	Built-in mutable sequence.
<code>done_tooltip</code>	
<code>is_finished</code>	A property that is True, when the task is finished
<code>name</code>	<code>str(object=") -> str</code>
<code>rst_file</code>	<code>str(object=") -> str</code>
<code>summary</code>	<code>str(object=") -> str</code>
<code>task_tooltip</code>	<code>str(object=") -> str</code>

```

dependencies = ['columns']

property done_tooltip
property is_finished
    A property that is True, when the task is finished
name = 'occurences'
rst_file = 'occurences'
summary = 'Select occurrence markers'
task_tooltip = 'Pollen diagrams often have markers for low taxon percentages to show t

class straditize.widgets.progress_widget.ProgressTask(parent)
Bases: PyQt5.QtWidgets.QListWidgetItem

```

The base class for an item that should be shown in the ProgressWidget

Parameters `parent` (`QListWidget`) – The list that holds this item

Attributes

<code>colnames_reader</code>	The column names reader of the straditizer
<code>data_reader</code>	The data reader of the straditizer
<code>dependencies</code>	List of name attributes that this task is depending on
<code>dependencies_tasks</code>	
<code>done</code>	True if <code>is_finished</code> or <code>done_by_user</code>
<code>done_by_user</code>	boolean that is True, when the task is marked as done by the user
<code>done_tooltip</code>	The tooltip when the straditizer is done.
<code>is_finished</code>	A property that is True, when the task is finished
<code>is_ready</code>	Boolean that is True if the task is ready to be solved
<code>name</code>	The name of the task
<code>progress_widget</code>	The progress widget that shows this task
<code>rst_file</code>	rst file that should be displayed on double click. The filename shuould
<code>state</code>	The state of the task
<code>straditizer</code>	The straditizer of the GUI
<code>summary</code>	The summary of the task that is shown in the progress widget
<code>task_tooltip</code>	The tooltip of the item
<code>try_finished</code>	Same as <code>is_finished</code> but catches every exception

Methods

`refresh()`

property colnames_reader
The column names reader of the straditizer

property data_reader
The data reader of the straditizer

dependencies = []
List of `name` attributes that this task is depending on

property dependencies_tasks

property done
True if `is_finished` or `done_by_user`

property done_by_user
boolean that is True, when the task is marked as done by the user

done_tooltip = None
The tooltip when the straditizer is done. If None, the `task_tooltip` is used

property is_finished
A property that is True, when the task is finished

is_ready = True
Boolean that is True if the task is ready to be solved

name = ''
The name of the task

property progress_widget
The progress widget that shows this task

refresh()

rst_file = None
rst file that should be displayed on double click. The filename shuould be without .rst ending.

property state
The state of the task

property straditizer
The straditizer of the GUI

summary = ''
The summary of the task that is shown in the progress widget

task_tooltip = ''
The tooltip of the item

property try_finished
Same as `is_finished` but catches every exception

class straditize.widgets.progress_widget.ProgressBar(straditizer_widgets, item)
Bases: PyQt5.QtWidgets.QWidget, `straditize.widgets.StraditzerControlBase`

A widget to show the progress of the straditization **Attributes**

`combo_display`

A QComboBox to select which tasks to display
(todo, done, not yet ready)

Continued on next page

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<code>info_label</code>	A QLabel to display the tooltip of the selected task
<code>progress_list</code>	A QListWidget to display the <code>ProgressTask</code> instances

Methods

<code>contextMenuEvent(event)</code>	Reimplement Qt method
<code>populate_list()</code>	Populate the <code>progress_list</code>
<code>refresh()</code>	Refresh from the straditizer
<code>setup_menu()</code>	Set up the context menu
<code>show_RST(item[, old])</code>	Show the documentation corresponding to a :class:`ProgressTask`
<code>toggle_done_by_user()</code>	
<code>update_info_label(item[, old])</code>	Update the <code>info_label</code> from a <code>ProgressTask</code>

`combo_display = None`

A QComboBox to select which tasks to display (todo, done, not yet ready or all tasks)

`contextMenuEvent(event)`

Reimplement Qt method

`info_label = None`

A QLabel to display the tooltip of the selected task

`populate_list()`

Populate the `progress_list`

This method adds instances of the `ProgressTask` class (or it's subclasses) to the the `progress_list`

`progress_list = None`

A QListWidget to display the `ProgressTask` instances

`refresh()`

Refresh from the straditizer

`setup_menu()`

Set up the context menu

`show_RST(item, old=None)`

Show the documentation corresponding to a :class:`ProgressTask`

Parameters

- `item` (`ProgressTask`) – The task to display it's `rst_file` in the `psyplot_gui.main.MainWindow.help_explorer`
- `old` (`ProgressTask`) – The old task that has been selected previously (this parameter is ignored)

`toggle_done_by_user()`

`update_info_label(item, old=None)`

Update the `info_label` from a `ProgressTask`

Parameters

- `item` (`ProgressTask`) – The selected task whose tooltip the `info_label` shall display

- **old** ([ProgressTask](#)) – The old task that has been selected previously (this parameter is ignored)

```
class straditize.widgets.progress_widget.RemoveArtifactsTask (parent)
Bases: straditize.widgets.progress_widget.ProgressTask
```

Task to clean the binary image

Parameters **parent** (*QListWidget*) – The list that holds this item

Attributes

<i>dependencies</i>	Built-in mutable sequence.
<i>done_tooltip</i>	
<i>is_finished</i>	A property that is True, when the task is finished
<i>name</i>	<code>str(object=") -> str</code>
<i>rst_file</i>	<code>str(object=") -> str</code>
<i>summary</i>	<code>str(object=") -> str</code>
<i>task_tooltip</i>	<code>str(object=") -> str</code>

```
dependencies = ['columns']

property done_tooltip

property is_finished
    A property that is True, when the task is finished

name = 'remove_artifacts'

rst_file = 'removing_features'

summary = 'Clean the diagram image'

task_tooltip = 'Remove all artifacts in the diagram part that do not represent data. May also remove some lines in the diagram part that do not represent data.'
```

```
class straditize.widgets.progress_widget.RemoveLinesTask (parent)
Bases: straditize.widgets.progress_widget.RemoveArtifactsTask
```

Task to remove y-axes, x-axes, horizontal and vertical lines

Parameters **parent** (*QListWidget*) – The list that holds this item

Attributes

<i>dependencies</i>	Built-in mutable sequence.
<i>done_tooltip</i>	
<i>is_finished</i>	A property that is True, when the task is finished
<i>name</i>	<code>str(object=") -> str</code>
<i>rst_file</i>	<code>str(object=") -> str</code>
<i>summary</i>	<code>str(object=") -> str</code>
<i>task_tooltip</i>	<code>str(object=") -> str</code>

```
dependencies = ['columns']

property done_tooltip

property is_finished
    A property that is True, when the task is finished

name = 'remove_lines'
```

```

rst_file = 'remove_lines'
summary = 'Remove y-, x-axes and other lines'
task_tooltip = 'In order to clean the diagram part, remove all vertical and horizontal
class straditize.widgets.progress_widget.SamplesTask(parent)
Bases: straditize.widgets.progress_widget.ProgressTask

```

Task to edit and find samples

Parameters `parent` (`QListWidget`) – The list that holds this item

Attributes

<code>dependencies</code>	Built-in mutable sequence.
<code>done_tooltip</code>	
<code>is_finished</code>	A property that is True, when the task is finished
<code>name</code>	<code>str(object=") -> str</code>
<code>rst_file</code>	<code>str(object=") -> str</code>
<code>summary</code>	<code>str(object=") -> str</code>
<code>task_tooltip</code>	<code>str(object=") -> str</code>

```

dependencies = ['digitize']
property done_tooltip
property is_finished
A property that is True, when the task is finished
name = 'samples'
rst_file = 'samples'
summary = 'Find and edit the samples'
task_tooltip = 'Find and edit the samples using the `Edit samples` menu'
class straditize.widgets.progress_widget.SaveProjectTask(*args, **kwargs)
Bases: straditize.widgets.progress_widget.ProgressTask

```

Task to remember saving the project

Parameters `parent` (`QListWidget`) – The list that holds this item

Attributes

<code>dependencies</code>	Built-in mutable sequence.
<code>is_finished</code>	True if the project was saved less than 5 minutes ago
<code>name</code>	<code>str(object=") -> str</code>
<code>rst_file</code>	<code>str(object=") -> str</code>
<code>summary</code>	<code>str(object=") -> str</code>
<code>task_tooltip</code>	<code>str(object=") -> str</code>

Methods

<code>refresh_tooltip()</code>
<code>tooltip()</code>

```

dependencies = ['init_stradi']

```

```
property is_finished
    True if the project was saved less than 5 minutes ago

name = 'save_project'

refresh_tooltip()

rst_file = 'save_and_load'

summary = 'Save the straditizer'

property task_tooltip
    str(object="") -> str str(bytes_or_buffer[, encoding[, errors]]) -> str

Create a new string object from the given object. If encoding or errors is specified, then the object must expose a data buffer that will be decoded using the given encoding and error handler. Otherwise, returns the result of object.__str__() (if defined) or repr(object). encoding defaults to sys.getdefaultencoding(). errors defaults to 'strict'.
```

```
tooltip()
```

```
class straditize.widgets.progress_widget.SelectExaggerationsTask(parent)
Bases: straditize.widgets.progress_widget.ProgressTask
```

Task to handle exaggerations

Parameters `parent` (`QListWidget`) – The list that holds this item

Attributes

<code>dependencies</code>	Built-in mutable sequence.
<code>done_tooltip</code>	
<code>is_finished</code>	A property that is True, when the task is finished
<code>name</code>	str(object="") -> str
<code>rst_file</code>	str(object="") -> str
<code>summary</code>	str(object="") -> str
<code>task_tooltip</code>	str(object="") -> str

```
dependencies = ['columns']
```

```
property done_tooltip
```

```
property is_finished
```

A property that is True, when the task is finished

```
name = 'exag'
```

```
rst_file = 'exaggerations'
```

```
summary = 'Select exaggerations'
```

```
task_tooltip = 'Pollen diagrams often display an exaggerated value of of the taxon per
```

```
class straditize.widgets.progress_widget.XTranslationTask(parent)
```

Bases: straditize.widgets.progress_widget.ProgressTask

Task to specify the x-axes conversion from pixel to data units

Parameters `parent` (`QListWidget`) – The list that holds this item

Attributes

<code>dependencies</code>	Built-in mutable sequence.
<code>done_tooltip</code>	
<code>is_finished</code>	A property that is True, when the task is finished
<code>name</code>	<code>str(object="") -> str</code>
<code>rst_file</code>	<code>str(object="") -> str</code>
<code>summary</code>	<code>str(object="") -> str</code>
<code>task_tooltip</code>	<code>str(object="") -> str</code>

```

dependencies = ['columns']

property done_tooltip
property is_finished
    A property that is True, when the task is finished
name = 'xaxes_trans'
rst_file = 'xaxis_translation'
summary = 'Transform the x-axes'

property task_tooltip
    str(object="") -> str str(bytes_or_buffer[, encoding[, errors]]) -> str
        Create a new string object from the given object. If encoding or errors is specified, then the object must expose a data buffer that will be decoded using the given encoding and error handler. Otherwise, returns the result of object.__str__() (if defined) or repr(object). encoding defaults to sys.getdefaultencoding(). errors defaults to 'strict'.

```

class `straditize.widgets.progress_widget.YTranslationTask` (*parent*)
Bases: `straditize.widgets.progress_widget.ProgressTask`

Task to specify the y-axis conversion from pixel to data units

Parameters `parent` (`QListWidget`) – The list that holds this item

Attributes

<code>dependencies</code>	Built-in mutable sequence.
<code>done_tooltip</code>	
<code>is_finished</code>	A property that is True, when the task is finished
<code>name</code>	<code>str(object="") -> str</code>
<code>rst_file</code>	<code>str(object="") -> str</code>
<code>summary</code>	<code>str(object="") -> str</code>
<code>task_tooltip</code>	<code>str(object="") -> str</code>

```

dependencies = ['init_reader']

property done_tooltip
property is_finished
    A property that is True, when the task is finished
name = 'yaxis_trans'
rst_file = 'yaxis_translation'
summary = 'Transform the y-axis'
task_tooltip = 'Transform the y-axis from pixel to data units'

```

straditize.widgets.samples_table module

A table for manipulating samples

This module defines the sample editors, either for editing the samples in the straditizer image (*SingleCrossMarksEditor*) or in a separate Figure (*MultiCrossMarksEditor*)

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Classes

<i>MultiCrossMarksEditor</i> (straditizer[, axes])	An editor for cross marks in multiple axes
<i>MultiCrossMarksModel</i> (marks, columns, straditzer)	A table model to handle multiple connected cross marks in different axes
<i>MultiCrossMarksView</i> (marks, full_df, *args, ...)	A table view set up by cross marks from multiple axes
<i>SingleCrossMarksEditor</i> (straditizer[, axes])	The editor for cross marks on a single axes
<i>SingleCrossMarksModel</i> (*args, **kwargs)	A table model to handle cross marks within one single axis
<i>SingleCrossMarksView</i> (marks, full_df, *args, ...)	A table for visualizing marks from a single axes

```
class straditize.widgets.samples_table.MultiCrossMarksEditor(straditizer,
                                                               axes=None, *args,
                                                               **kwargs)
```

Bases: `psyplot_gui.common.DockMixin, PyQt5.QtWidgets.QWidget`

An editor for cross marks in multiple axes

Parameters

- **straditizer** (`weakref.ref`) – The reference to the straditizer
- **axes** (`matplotlib.axes.Axes`) – The matplotlib axes corresponding to the marks

Attributes

<code>cb_plot_lines</code>	Plot the reconstructed data
<code>straditzer</code>	A weakref to the

Methods

<code>create_view([axes])</code>	Create the <i>MultiCrossMarksView</i> of the editor
<code>maybe_show_selection_only()</code>	
<code>maybe_tabify()</code>	

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<code>maybe_zoom_to_selection()</code>	
<code>save_samples()</code>	Save the samples to the straditizer without removing them
<code>to_dock(main[, title, position, docktype])</code>	
<code>toggle_cb_selection_only()</code>	
<code>toggle_cb_zoom_to_selection()</code>	
<code>toggle_fit2selection()</code>	Enable the fitting so selected digitized data
<code>toggle_fmt_button(text)</code>	
<code>toggle_plot_lines()</code>	
<code>update_format()</code>	Update the format of the table

Classes

<code>dock_cls</code>	The QDockWidget for the DataFrameEditor
-----------------------	---

```

cb_plot_lines = None
    Plot the reconstructed data

create_view(axes=None)
    Create the MultiCrossMarksView of the editor

    Parameters axes (list of matplotlib.axes.Axes) – The matplotlib axes for the marks

dock_cls
    alias of psyplot_gui.dataframeeditor.DataFrameDock

maybe_show_selection_only()

maybe_tabify()

maybe_zoom_to_selection()

save_samples()
    Save the samples to the straditizer without removing them

straditizer = None
    A weakref to the straditizer

to_dock(main, title=None, position=None, docktype='df', *args, **kwargs)

toggle_cb_selection_only()

toggle_cb_zoom_to_selection()

toggle_fit2selection()
    Enable the fitting so selected digitized data

toggle_fmt_button(text)

toggle_plot_lines()

update_format()
    Update the format of the table

class straditize.widgets.samples_table.MultiCrossMarksModel (marks, columns,
                                                 straditizer,
                                                 axes=None,
                                                 occurrences_value=-
                                                 9999)

```

Bases: PyQt5.QtCore.QAbstractTableModel

A table model to handle multiple connected cross marks in different axes

Parameters

- **marks** (list of `straditize.cross_mark.CrossMarks`) – the initial marks
- **columns** (list of `str`) – the column names to use
- **straditizer** (`straditize.straditizer.Straditizer`) – The straditizer that manages the *marks*
- **axes** (list of `matplotlib.axes.Axes`) – The matplotlib axes that contain the *marks*
- **occurrences_value** (`float`) – The value that marks an occurrence

Methods

<code>columnCount([index])</code>	The number of rows in the table
<code>data(self, QModelIndex, role)</code>	
<code>delRow(irow)</code>	
<code>flags(index)</code>	Set flags
<code>get_cell_mark(row, column)</code>	Get the mark for a given cell in the table
<code>get_format()</code>	Return current format
<code>headerData(section, orientation[, role])</code>	Set header data
<code>insertRow(irow[, xa, ya])</code>	Insert a row into the table
<code>load_new_marks(mark)</code>	Add a new mark into the table after they have been added by the user
<code>plot_lines()</code>	Connect the samples through visual lines
<code>remove_lines()</code>	Remove the lines
<code>remove_mark(mark)</code>	Remove a mark from the table after it has been removed by the user
<code>reset()</code>	Reset the model
<code>rowCount([index])</code>	The number of rows in the table
<code>setData(index, value[, role, change_type])</code>	Cell content change
<code>set_format(fmt)</code>	Change display format
<code>set_marks(marks, columns)</code>	Set the <i>marks</i> attribute from the given <i>columns</i>
<code>sort_marks()</code>	Sort the <i>marks</i> based on there y-position
<code>update_after_move(old_pos, mark)</code>	
<code>update_lines()</code>	Update the lines or plot them

Attributes

<code>df</code>	Get the samples_locs
<code>fig</code>	The figure of the cross marks
<code>iter_marks</code>	Iter over all marks in the <i>marks</i> attribute
<code>lines</code>	A list of <code>matplotlib.lines.Line2D</code> that connects the cross marks
<code>marks</code>	A list [<code>float</code> , <code>list</code>] where the first <code>float</code> is the vertical

`columnCount (index=<PyQt5.QtCore.QModelIndex object>)`

The number of rows in the table

`data (self, QModelIndex, role: int = Qt.DisplayRole) → Any`

`delRow (irow)`

```

property df
    Get the samples_locs

property fig
    The figure of the cross marks

flags (index)
    Set flags

get_cell_mark (row, column)
    Get the mark for a given cell in the table

    Parameters
        • row (int) – The row of the cell
        • column (int) – The column of the cell

    Returns The corresponding mark from the marks attribute

    Return type straditize.cross_mark.CrossMarks

get_format ()
    Return current format

headerData (section, orientation, role=0)
    Set header data

insertRow (irow, xa=None, ya=None)
    Insert a row into the table

    Parameters irow (int) – The row index. If irow is equal to the length of the marks, the rows
        will be appended

property iter_marks
    Iter over all marks in the marks attribute

lines = []
    A list of matplotlib.lines.Line2D that connects the cross marks and plots a reconstruction based
    on them

load_new_marks (mark)
    Add a new mark into the table after they have been added by the user

    Parameters mark (straditize.cross_mark.CrossMarks) – The added mark

marks = []
    A list [float, list] where the first float is the vertical position and the second list is a list of
    the corresponding CrossMarks instances

plot_lines ()
    Connect the samples through visual lines

remove_lines ()
    Remove the lines

remove_mark (mark)
    Remove a mark from the table after it has been removed by the user

    Parameters mark (straditize.cross_mark.CrossMarks) – The removed mark

reset ()
    Reset the model

```

rowCount (*index=<PyQt5.QtCore.QModelIndex object>*)

The number of rows in the table

setData (*index, value, role=2, change_type=None*)

Cell content change

set_format (*fmt*)

Change display format

set_marks (*marks, columns*)

Set the *marks* attribute from the given *columns*

Parameters

- **marks** (list of *straditize.cross_mark.CrossMarks*) – the initial marks

- **columns** (list of str) – the column names to use

sort_marks()

Sort the marks based on there y-position

update_after_move (*old_pos, mark*)

update_lines()

Update the *lines* or plot them

See also:

[plot_lines\(\)](#)

class straditize.widgets.samples_table.**MultiCrossMarksView** (*marks, full_df, *args, **kwargs*)

Bases: PyQt5.QtWidgets.QTableView

A table view set up by cross marks from multiple axes

The model for this table is the *MultiCrossMarksModel*

Parameters

- **marks** (list of *straditize.cross_mark.CrossMarks*) – the initial marks

- **full_df** (*pandas.DataFrame*) – The data fame of the full digitized data

Methods

contextMenuEvent (event)	Reimplement Qt method
delete_selected_rows()	
fit2data()	Fit the selected cells to the <i>full_df</i>
init_model (marks, *args, **kwargs)	Initialize the table <i>MultiCrossMarksModel</i>
insert_row_above_selection()	Insert a row above the selection
insert_row_below_selection()	Insert a row below the selection
moveCursor (cursor_action, modifiers)	Update the table position.
resizeEvent (event)	Update the frozen column dimensions.
scrollTo (index, hint)	Scroll the table.
setup_menu()	Setup context menu
show_all_marks()	Show all marks
show_selected_marks_only()	Show only the marks selected in the table
update_section_height (logical_index, ...)	Update the vertical width of the frozen column when a

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<code>update_section_width(logical_index, ...)</code>	Update the horizontal width of the frozen column when a
<code>zoom_to_cells(rows, cols)</code>	Zoom to specific cells in the plot
<code>zoom_to_selection()</code>	Zoom to the selected cells in the plot

Attributes

<code>full_df</code>	The <code>pandas.DataFrame</code> representing the full digitized data
----------------------	--

<code>contextMenuEvent (event)</code>	Reimplement Qt method
<code>delete_selected_rows ()</code>	
<code>fit2data ()</code>	Fit the selected cells to the <code>full_df</code>
<code>full_df = None</code>	The <code>pandas.DataFrame</code> representing the full digitized data from the <code>straditize.binary.DataReader.full_df</code> data frame
<code>init_model (marks, *args, **kwargs)</code>	Initialize the table <code>MultiCrossMarksModel</code>
<code>insert_row_above_selection ()</code>	Insert a row above the selection
<code>insert_row_below_selection ()</code>	Insert a row below the selection
<code>moveCursor (cursor_action, modifiers)</code>	Updates the table position.
	Updates the position along with the frozen column when the cursor (selector) changes its position
<code>resizeEvent (event)</code>	Update the frozen column dimensions.
	Updates takes place when the enclosing window of this table reports a dimension change
<code>scrollTo (index, hint)</code>	Scroll the table.
	It is necessary to ensure that the item at index is visible. The view will try to position the item according to the given hint. This method does not takes effect only if the frozen column is scrolled.
<code>setup_menu ()</code>	Setup context menu
<code>show_all_marks ()</code>	Show all marks
See also:	
<code>show_selected_marks_only ()</code>	
<code>show_selected_marks_only ()</code>	Show only the marks selected in the table
See also:	

`show_all_marks()`

`update_section_height(logical_index, old_size, new_size)`

Update the vertical width of the frozen column when a change takes place on any of the rows

`update_section_width(logical_index, old_size, new_size)`

Update the horizontal width of the frozen column when a change takes place in the first column of the table

`zoom_to_cells(rows, cols)`

Zoom to specific cells in the plot

Parameters

- `rows` (*list of int*) – The row indices of the cells
- `cols` (*list of int*) – The column indices of the cells

`zoom_to_selection()`

Zoom to the selected cells in the plot

class straditize.widgets.samples_table.**SingleCrossMarksEditor**(straditizer,
 axes=None,
 *args, **kwargs)

Bases: *straditize.widgets.samples_table.MultiCrossMarksEditor*

The editor for cross marks on a single axes

Parameters

- `straditizer` (*weakref.ref*) – The reference to the straditizer
- `axes` (*matplotlib.axes.Axes*) – The matplotlib axes corresponding to the marks

Methods

<code>create_view([axes])</code>	Create the <i>SingleCrossMarksView</i> of the editor
----------------------------------	--

<code>save_samples()</code>	Save the samples to the straditizer without removing them
-----------------------------	---

`create_view(axes=None)`

Create the *SingleCrossMarksView* of the editor

Parameters `axes` (*list of matplotlib.axes.Axes*) – The matplotlib axes for the marks

`save_samples()`

Save the samples to the straditizer without removing them

class straditize.widgets.samples_table.**SingleCrossMarksModel**(*args, **kwargs)
Bases: *straditize.widgets.samples_table.MultiCrossMarksModel*

A table model to handle cross marks within one single axis

Parameters

- `column_bounds` (*np.ndarray* of shape (N, 2)) – The column boundaries
- `y0` (*float*) – The upper extent of the data image

Methods

<code>columnCount([index])</code>	The number of rows in the table
<code>delRow(irow)</code>	
<code>get_cell_mark(row, column)</code>	Get the mark for a given cell in the table
<code>insertRow(irow[, xa, ya])</code>	Insert a row into the table
<code>load_new_marks(mark)</code>	Add a new mark into the table after they have been added by the user
<code>plot_lines()</code>	Connect the samples through visual lines
<code>remove_mark(mark)</code>	Remove a mark from the table after it has been removed by the user
<code>set_marks(marks, columns)</code>	Set the <code>marks</code> attribute from the given <code>columns</code>
<code>update_after_move(old_pos, mark)</code>	
<code>update_lines()</code>	Update the lines or plot them

Attributes

<code>df</code>	Get the <code>samples_locs</code>
<code>iter_marks</code>	Iter over all marks in the <code>marks</code> attribute
<code>marks</code>	A list of tuples like <code>(float, mark)</code> where <code>float</code> is the y-pixel

columnCount (`index=<PyQt5.QtCore.QModelIndex object>`)

The number of rows in the table

delRow (`irow`)

property df

Get the `samples_locs`

get_cell_mark (`row, column`)

Get the mark for a given cell in the table

Parameters

- **row** (`int`) – The row of the cell
- **column** (`int`) – The column of the cell

Returns The corresponding mark from the `marks` attribute

Return type `straditiz.cross_mark.CrossMarks`

insertRow (`irow, xa=None, ya=None`)

Insert a row into the table

Parameters `irow` (`int`) – The row index. If `irow` is equal to the length of the `marks`, the rows will be appended

property iter_marks

Iter over all marks in the `marks` attribute

load_new_marks (`mark`)

Add a new mark into the table after they have been added by the user

Parameters `mark` (`straditiz.cross_mark.CrossMarks`) – The added mark

marks = []

A list of tuples like `(float, mark)` where `float` is the y-pixel and `mark` is the corresponding `straditiz.cross_mark.CrossMarks` instance

plot_lines()

Connect the samples through visual lines

remove_mark(mark)

Remove a mark from the table after it has been removed by the user

Parameters `mark (straditize.cross_mark.CrossMarks)` – The removed mark

set_marks(marks, columns)

Set the `marks` attribute from the given `columns`

Parameters

- `marks` (list of `straditize.cross_mark.CrossMarks`) – the initial marks

- `columns` (*list of str*) – the column names to use

update_after_move(old_pos, mark)

update_lines()

Update the lines or plot them

See also:

`plot_lines()`

class `straditize.widgets.samples_table.SingleCrossMarksView(marks, full_df, *args, **kwargs)`

Bases: `straditize.widgets.samples_table.MultiCrossMarksView`

A table for visualizing marks from a single axes

Parameters

- `marks` (list of `straditize.cross_mark.CrossMarks`) – the initial marks

- `full_df` (`pandas.DataFrame`) – The data fame of the full digitized data

Methods

`fit2data()`

Fit the selected cells to the `full_df`

`init_model(marks, *args, **kwargs)`

Initialize the table `SingleCrossMarksModel`

`zoom_to_cells(rows, cols)`

Zoom to specific cells in the plot

`fit2data()`

Fit the selected cells to the `full_df`

`init_model(marks, *args, **kwargs)`

Initialize the table `SingleCrossMarksModel`

Parameters % (`SingleCrossMarksModel.parameters`) s –

`zoom_to_cells(rows, cols)`

Zoom to specific cells in the plot

Parameters

- `rows` (*list of int*) – The row indices of the cells

- `cols` (*list of int*) – The column indices of the cells

straditize.widgets.selection_toolbar module

Module for the selection toolbar

This module defines the selection toolbar that is added to the `psyplot_gui.main.MainWindow` for selecting features in the stratigraphic diagram and the data reader image.

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Classes

<code>PointOrRectangleSelector(ax, onselect[, ...])</code>	RectangleSelector that allows to select points
<code>SelectionToolbar(straditzer_widgets, *args, ...)</code>	A toolbar for selecting features in the straditzer and data image

```
class straditize.widgets.selection_toolbar.PointOrRectangleSelector(ax, on-
                     select,
                     draw-
                     type='box',
                     minspanx=None,
                     minspany=None,
                     use-
                     blit=False,
                     line-
                     props=None,
                     rect-
                     props=None,
                     spanco-
                     ords='data',
                     but-
                     ton=None,
                     maxdist=10,
                     marker_props=None,
                     interac-
                     tive=False,
                     state_modifier_keys=None)
```

Bases: `matplotlib.widgets.RectangleSelector`

RectangleSelector that allows to select points

This class reimplements the `matplotlib.widgets.RectangleSelector` to select points

Create a selector in `ax`. When a selection is made, clear the span and call `onselect` with:

```
onselect(pos_1, pos_2)
```

Methods

`press(*args, **kwargs)`

Button press handler and validator

and clear the drawn box/line. The `pos_1` and `pos_2` are arrays of length 2 containing the x- and y-coordinate. If `minspanx` is not `None` then events smaller than `minspanx` in x direction are ignored (it's the same for y).

The rectangle is drawn with `rectprops`; default:

```
rectprops = dict(facecolor='red', edgecolor = 'black',
                 alpha=0.2, fill=True)
```

The line is drawn with `lineprops`; default:

```
lineprops = dict(color='black', linestyle='-', linewidth = 2, alpha=0.5)
```

Use `drawtype` if you want the mouse to draw a line, a box or nothing between click and actual position by setting `drawtype = 'line'`, `drawtype='box'` or `drawtype = 'none'`. Drawing a line would result in a line from vertex A to vertex C in a rectangle ABCD.

`spancoords` is one of ‘data’ or ‘pixels’. If ‘data’, `minspanx` and `minspany` will be interpreted in the same coordinates as the x and y axis. If ‘pixels’, they are in pixels.

`button` is a list of integers indicating which mouse buttons should be used for rectangle selection. You can also specify a single integer if only a single button is desired. Default is `None`, which does not limit which button can be used.

Note, typically: 1 = left mouse button 2 = center mouse button (scroll wheel) 3 = right mouse button

`interactive` will draw a set of handles and allow you interact with the widget after it is drawn.

`state_modifier_keys` are keyboard modifiers that affect the behavior of the widget.

The defaults are: `dict(move=' ', clear='escape', square='shift', center='ctrl')`

Keyboard modifiers, which: ‘move’: Move the existing shape. ‘clear’: Clear the current shape. ‘square’: Makes the shape square. ‘center’: Make the initial point the center of the shape. ‘square’ and ‘center’ can be combined.

`press (*args, **kwargs)`

Button press handler and validator

```
class straditize.widgets.selection_toolbar.SelectionToolbar(straditzer_widgets,
                                                               *args, **kwargs)
Bases: PyQt5.QtWidgets.QToolBar, straditize.widgets.StraditizerControlBase
```

A toolbar for selecting features in the straditizer and data image

The current data object is set in the `combo` and can be accessed through the `data_obj` attribute. It's either the straditizer or the data_reader that is accessed **Methods**

<code>add_or_remove_pattern()</code>	Enable the removing or adding of the pattern selection
<code>clear_selection()</code>	Clear the current selection
<code>create_actions()</code>	Define the actions for the toolbar and set everything up
<code>disable_actions()</code>	
<code>disconnect(self)</code>	

Continued on next page

Table 129 – continued from previous page

<code>enable_or_disable_widgets(b)</code>	Enable or disable the widgets in this control
<code>end_selection()</code>	Finish the selection and disconnect everything
<code>expand_selection()</code>	Expand the selected areas to select the full labels
<code>get_xy_slice(lastx, lasty, x, y)</code>	Transform x- and y-coordinates to <code>slice</code> objects
<code>invert_selection()</code>	Invert the current selection
<code>on_poly_select(points)</code>	Call the <code>poly_callbacks</code> after a polygon selection
<code>on_rect_select(e0, e1)</code>	Call the <code>rect_callbacks</code> after a rectangle selection
<code>refresh()</code>	Refresh from the straditizer
<code>select_all()</code>	Select all features in the image
<code>select_everything_to_the_right()</code>	Selects everything to the right of the current selection
<code>select_poly(points)</code>	Select the data defined by a polygon
<code>select_rect(slx, sly)</code>	Select the data defined by a rectangle
<code>set_binary_pattern_mode()</code>	Set the current pattern mode to the binary pattern
<code>set_col_wand_mode()</code>	Set the current wand tool to the color wand
<code>set_color_wand_mode()</code>	Set the current wand tool to the color wand
<code>set_grey_pattern_mode()</code>	Set the current pattern mode to the binary pattern
<code>set_label_wand_mode()</code>	Set the current wand tool to the color wand
<code>set_poly_select_mode()</code>	Set the current wand tool to the color wand
<code>set_rect_select_mode()</code>	Set the current wand tool to the color wand
<code>set_row_wand_mode()</code>	Set the current wand tool to the color wand
<code>should_be_enabled(w)</code>	Check if a widget should be enabled
<code>start_pattern_selection()</code>	Open the pattern selection dialog
<code>start_selection([arr, rgba, rect_callbacks, ...])</code>	Start the selection in the current <code>data_obj</code>
<code>toggle_selection()</code>	Activate selection mode
<code>uncheck_pattern_selection()</code>	Disable the pattern selection
<code>update_alpha(i)</code>	Set the transparency of the selection image

Attributes

<code>add_select_action</code>	The action to add to the current selection with the selection tools
<code>ax</code>	The <code>matplotlib.axes.Axes</code> of the <code>data_obj</code>
<code>canvas</code>	The canvas of the <code>data_obj</code>
<code>clear_select_action</code>	An action to clear the current selection
<code>combo</code>	The QCombobox that defines the data object to be used
<code>data</code>	The <code>np.ndarray</code> of the <code>data_obj</code> image
<code>data_obj</code>	The data object as set in the <code>combo</code> .
<code>expand_select_action</code>	An action to expand the current selection to the full feature
<code>fig</code>	The <code>Figure</code> of the <code>data_obj</code>
<code>invert_select_action</code>	An action to invert the current selection
<code>labels</code>	The labeled data that is displayed
<code>new_select_action</code>	The action to make new selection with one of the selection tools
<code>poly_callbacks</code>	The functions to call after the polygon selection

Continued on next page

Table 130 – continued from previous page

<code>rect_callbacks</code>	The functions to call after the rectangle selection.
<code>remove_select_action</code>	An action to remove from the current selection with the selection tools
<code>select_action</code>	The rectangle selection tool
<code>select_all_action</code>	An action to select all features in the <code>data</code>
<code>select_pattern_action</code>	An action to start a pattern selection
<code>select_right_action</code>	An action to select everything in the data column to the right
<code>selected(*args, **kwargs)</code>	A signal that is emitted when something is selected
<code>selector</code>	A <code>PointOrRectangleSelector</code> to select features in the image
<code>toolbar</code>	The toolbar of the canvas
<code>wand_action</code>	The wand selection tool
<code>widgets2disable</code>	Built-in mutable sequence.

`add_or_remove_pattern()`

Enable the removing or adding of the pattern selection

`property add_select_action`

The action to add to the current selection with the selection tools

`property ax`

The `matplotlib.axes.Axes` of the `data_obj`

`property canvas`

The canvas of the `data_obj`

`property clear_select_action`

An action to clear the current selection

`clear_selection()`

Clear the current selection

`combo = None`

The QCombobox that defines the data object to be used

`create_actions()`

Define the actions for the toolbar and set everything up

`property data`

The `np.ndarray` of the `data_obj` image

`property data_obj`

The data object as set in the `combo`.

Either a `Straditizer` or a `straditize.binary.DataReader` instance.

`disable_actions()`**`disconnect(self)`****`enable_or_disable_widgets(b)`**

Enable or disable the widgets in this control

This method enables or disables the `widgets2disable` if the `should_be_enabled()` method evaluates to True

Parameters `b` (`bool`) – If True, enable the widgets, if False, disable them

end_selection()
Finish the selection and disconnect everything

property expand_select_action
An action to expand the current selection to the full feature

expand_selection()
Expand the selected areas to select the full labels

property fig
The `Figure` of the `data_obj`

get_xy_slice(lastx, lasty, x, y)
Transform x- and y-coordinates to `slice` objects

Parameters

- **lastx** (`int`) – The initial x-coordinate
- **lasty** (`int`) – The initial y-coordinate
- **x** (`int`) – The final x-coordinate
- **y** (`int`) – The final y-coordinate

Returns

- `slice` – The `slice(lastx, x)`
- `slice` – The `slice(lasty, y)`

property invert_select_action
An action to invert the current selection

invert_selection()
Invert the current selection

property labels
The labeled data that is displayed

property new_select_action
The action to make new selection with one of the selection tools

on_poly_select(points)
Call the `poly_callbacks` after a polygon selection

Parameters

- **e0** (`matplotlib.backend_bases.Event`) – The initial event
- **e1** (`matplotlib.backend_bases.Event`) – The final event

on_rect_select(e0, e1)
Call the `rect_callbacks` after a rectangle selection

Parameters

- **e0** (`matplotlib.backend_bases.Event`) – The initial event
- **e1** (`matplotlib.backend_bases.Event`) – The final event

property poly_callbacks
The functions to call after the polygon selection

If not set manually, it is the `select_poly()` method. Note that this is cleared at every call of the `end_selection()`.

Callables in this list must accept one argument, a `np.ndarray` of shape `(N, 2)`. This array defines the N x- and y-coordinates of the points of the polygon

property rect_callbacks

The functions to call after the rectangle selection.

If not set manually, it is the `select_rect()` method. Note that this is cleared at every call of the `end_selection()`.

Callables in this list must accept two arguments `(slx, sly)`: the first one is the x-slice, and the second one the y-slice. They both correspond to the `data` attribute.

refresh()

Refresh from the straditizer

property remove_select_action

An action to remove from the current selection with the selection tools

reset_cursor_id = None**property select_action**

The rectangle selection tool

select_all()

Select all features in the image

See also:

`straditize.label_selection.LabelSelection.select_all_labels()`

property select_all_action

An action to select all features in the `data`

select_everything_to_the_right()

Selects everything to the right of the current selection

property select_pattern_action

An action to start a pattern selection

select_poly(points)

Select the data defined by a polygon

Parameters `points (np.ndarray of shape (N, 2))` – The x- and y-coordinates of the vertices of the polygon

See also:

`poly_callbacks()`

select_rect(slx, sly)

Select the data defined by a rectangle

Parameters

- `slx (slice)` – The x-slice of the rectangle
- `sly (slice)` – The y-slice of the rectangle

See also:

`rect_callbacks()`

property select_right_action

An action to select everything in the data column to the right

selected(*args, **kwargs)

A signal that is emitted when something is selected

selector = None

A `PointOrRectangleSelector` to select features in the image

set_binary_pattern_mode()

Set the current pattern mode to the binary pattern

set_col_wand_mode()

Set the current wand tool to the color wand

set_color_wand_mode()

Set the current wand tool to the color wand

set_cursor_id = None

set_grey_pattern_mode()

Set the current pattern mode to the binary pattern

set_label_wand_mode()

Set the current wand tool to the color wand

set_poly_select_mode()

Set the current wand tool to the color wand

set_rect_select_mode()

Set the current wand tool to the color wand

set_row_wand_mode()

Set the current wand tool to the color wand

should_be_enabled(w)

Check if a widget should be enabled

This function checks if a given widget `w` from the `widgets2disable` attribute should be enabled or not

Parameters `w` (`QWidget`) – The widget to check

Returns True, if the widget should be enabled

Return type `bool`

start_pattern_selection()

Open the pattern selection dialog

This method will enable the pattern selection by starting a `straditize.widgets.pattern_selection.PatternSelectionWidget`.

start_selection(arr=None, rgba=None, rect_callbacks=None, poly_callbacks=None, apply_funcs=(), cancel_funcs=(), remove_on_apply=True)

Start the selection in the current `data_obj`

Parameters

- **arr** (`np.ndarray`) – The labeled selection array that is used. If specified, the `enable_label_selection()` method is called of the `data_obj` with the given `arr`. If this parameter is `None`, then we expect that this method has already been called
- **rgba** (`np.ndarray`) – The RGBA image that shall be used for the color selection (see the `set_color_wand_mode()`)
- **rect_callbacks** (`list`) – A list of callbacks that shall be called after a rectangle selection has been made by the user (see `rect_callbacks`)

- **poly_callbacks** (*list*) – A list of callbacks that shall be called after a polygon selection has been made by the user (see [poly_callbacks](#))
- **apply_funcs** (*list*) – A list of callables that shall be connected to the [apply_button](#)
- **cancel_funcs** (*list*) – A list of callables that shall be connected to the [cancel_button](#)
- **remove_on_apply** (*bool*) – If True and the [apply_button](#) is clicked, the selected labels will be removed.

toggle_selection()

Activate selection mode

property toolbar

The toolbar of the [canvas](#)

uncheck_pattern_selection()

Disable the pattern selection

update_alpha(*i*)

Set the transparency of the selection image

Parameters **i** (*int*) – The transparency between 0 and 100

property wand_action

The wand selection tool

property widgets2disable

Built-in mutable sequence.

If no argument is given, the constructor creates a new empty list. The argument must be an iterable if specified.

straditize.widgets.stacked_area_reader module

DataReader for stacked area plots

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Classes

StackedReader(image[, ax, extent, plot, ...])

A DataReader for stacked area plots

```
class straditiz.widgets.stacked_area_reader.StackedReader(image, ax=None, extent=None, plot=True,  
    children=[],  
    parent=None,  
    magni=None,  
    plot_background=False,  
    binary=None)  
  
Bases: straditiz.binary.DataReader,  
StraditizerControlBase
```

A DataReader for stacked area plots

This reader only works within the straditizer GUI because the digitization (see `digitize()`) is interactive. The user has to manually distinguish the stacked variables.

Parameters

- **image** (`PIL.Image.Image`) – The image of the diagram
- **ax** (`matplotlib.axes.Axes`) – The matplotlib axes to plot on
- **extent** (`list`) – List of four number specifying the extent of the image in it's source. This extent will be used for the call of `matplotlib.pyplot.imshow()`
- **children** (list of DataReader) – Child readers for other columns in case the newly created instance is the parent reader
- **parent** (DataReader) – The parent reader.
- **magni** (`straditiz.magnifier.Magnifier`) – The magnifier for the given *ax*
- **plot_background** (`bool`) – If True (and *plot* is True), a white, opaque arc is plotted below the *plot_im*
- **binary** (`None`) – The binary version of the given *image*. If not provided, the `to_binary_pil()` method is used with the given *image*

Methods

<code>add_col()</code>	Create a column out of the current selection
<code>decrease_current_col()</code>	Take the previous column as the current column
<code>digitize()</code>	Digitize the data interactively
<code>enable_or_disable_navigation_buttons()</code>	Enable or disable <code>btn_prev</code> and <code>btn_next</code>
<code>get_binary_for_col(<i>col</i>)</code>	Get the binary array for a specific column
<code>increase_current_col()</code>	Take the next column as the current column
<code>plot_full_df([<i>ax</i>])</code>	Plot the lines for the digitized diagram
<code>plot_potential_samples([<i>excluded</i>, <i>ax</i>, <i>kws</i>])</code>	Plot the ranges for potential samples
<code>reset_lbl_col()</code>	Reset the <code>lbl_col</code> to display the current column
<code>resize_axes(<i>grouper</i>, <i>bounds</i>)</code>	Reimplemented to do nothing
<code>select_and_add_current_column()</code>	Select the features for a column and create it as a new one
<code>select_current_column()</code>	Select the features of the current column
<code>update_col()</code>	Update the current column based on the selection.
<code>update_plotted_full_df()</code>	Update the plotted <code>full_df</code> if it is shown

Attributes

<code>btn_add</code>	A QPushButton to add a new variable to the current ones
<code>btn_edit</code>	A QPushButton to select the features in the image for the current
<code>btn_next</code>	A QPushButton to select the next variable during the digitization
<code>btn_prev</code>	A QPushButton to select the previous variable during the digitization
<code>digitize_child</code>	The QTreeWidgetItem that holds the digitization widgets
<code>lbl_col</code>	A QLabel to display the current column
<code>start_of_current_col</code>	The first x-pixel of the current column
<code>strat_plot_identifier</code>	str(object=”) -> str

`add_col()`

Create a column out of the current selection

`btn_add = None`

A QPushButton to add a new variable to the current ones (see `select_and_add_current_column()`)

`btn_edit = None`

A QPushButton to select the features in the image for the current variable (see `select_current_column()`)

`btn_next = None`

A QPushButton to select the next variable during the digitization (see `increase_current_col()`)

`btn_prev = None`

A QPushButton to select the previous variable during the digitization (see `decrease_current_col()`)

`decrease_current_col()`

Take the previous column as the current column

`digitize()`

Digitize the data interactively

This method creates a new child item for the digitize button in the straditzer control to manually distinguish the variables in the stacked diagram.

`digitize_child = None`

The QTreeWidgetItem that holds the digitization widgets

`enable_or_disable_navigation_buttons()`

Enable or disable `btn_prev` and `btn_next`

Depending on the current column, we disable the navigation buttons `btn_prev` and `btn_next`

`get_binary_for_col(col)`

Get the binary array for a specific column

`increase_current_col()`

Take the next column as the current column

`lbl_col = None`

A QLabel to display the current column

`plot_full_df(ax=None)`

Plot the lines for the digitized diagram

plot_potential_samples (*excluded=False, ax=None, plot_kws={}, *args, **kwargs*)
 Plot the ranges for potential samples

reset_lbl_col()
 Reset the `lbl_col` to display the current column

resize_axes (*grouper, bounds*)
 Reimplemented to do nothing

select_and_add_current_column()
 Select the features for a column and create it as a new one

select_current_column()
 Select the features of the current column

property start_of_current_col
 The first x-pixel of the current column

strat_plot_identifier = 'stacked'

update_col()
 Update the current column based on the selection.
 This method updates the end of the current column and adds or removes the changes from the columns to the right.

update_plotted_full_df()
 Update the plotted full_df if it is shown

See also:
`plot_full_df()`

1.7.2 Submodules

straditize.__main__ module

main module of straditize

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Functions

<code>get_parser([create])</code>	Create an argument parser for the command line handling
<code>main([exec_])</code>	
<code>start_app([fname, output, xlim, ylim, full, ...])</code>	Start the psyplot GUI with the straditzer setup

`straditize.__main__.get_parser(create=True)`
Create an argument parser for the command line handling

This function creates a `funcargparse.FuncArgParser` for the usage from the command line

Parameters `create (bool)` – If True, the `funcargparse.FuncArgParser.create_arguments()` method is called

`straditize.__main__.main(exec_=True)`

`straditize.__main__.start_app(fname=None, output=None, xlim=None, ylim=None, full=False, reader_type='area', **kwargs)`

Start the psyplot GUI with the straditizer setup

Parameters

- `fname (str)` – Either the path to a picture to digitize or a previously saved straditizer project (ending with '.pkl')
- `output (str)` – The path to the csv file where to save the digitized diagram
- `xlim (list of int of length 2)` – The x-limits of the data part of the diagram
- `ylim (list of int of length 2)` – The y-limits of the data part of the diagram
- `full (bool)` – If True, the image is digitized and x- and ylim are set to the entire share of the array
- `reader_type ({ 'area' / 'bars' / 'rounded bars' / 'stacked area' / 'line' })` – Specify the reader type
- `name (list of str)` – The variable names to plot if the `output` parameter is set
- `dims (dict)` – A mapping from coordinate names to integers if the `project` is not given
- `plot_method (str)` – The name of the plot_method to use
- `project (str)` – If set, the project located at the given file name is loaded
- `engine (str)` – The engine to use for opening the dataset (see `psyplot.data.open_dataset()`)
- `formatoptions (dict)` – A dictionary of formatoption that is applied to the data visualized by the chosen `plot_method`
- `tight (bool)` – If True/set, it is tried to figure out the tight bbox of the figure and adjust the paper size of the `output` to it
- `rc_file (str)` – The path to a yaml configuration file that can be used to update the `rcParams`
- `encoding (str)` – The encoding to use for loading the project. If None, it is automatically determined by pickle. Note: Set this to 'latin1' if using a project created with python2 on python3.
- `enable_post (bool)` – Enable the `post` processing formatoption. If True/set, post processing scripts are enabled in the given `project`. Only set this if you are sure that you can trust the given project file because it may be a security vulnerability.
- `seaborn_style (str)` – The name of the style of the seaborn package that can be used for the `seaborn.set_style()` function
- `output_project (str)` – The name of a project file to save the project to
- `concat_dim (str)` – The concatenation dimension if multiple files in `fnames` are provided

- **chname** (*dict*) – A mapping from variable names in the project to variable names in the datasets that should be used instead
- **backend** (*None or str*) – The backend to use. By default, the 'gui.backend' key in the `rcParams` dictionary is used. Otherwise it can be `None` to use the standard matplotlib backend or a string identifying the backend
- **new_instance** (*bool*) – If `True`/set and the `output` parameter is not set, a new application is created
- **rc_gui_file** (*str*) – The path to a yaml configuration file that can be used to update the `rcParams`
- **include_plugins** (*list of str*) – The plugin widget to include. Can be either `None` to load all that are not explicitly excluded by `exclude_plugins` or a list of plugins to include. List items can be either module names, plugin names or the module name and widget via '`<module_name>:<widget>`'
- **exclude_plugins** (*list of str*) – The plugin widgets to exclude. Can be either '`'all'`' to exclude all plugins or a list like in `include_plugins`.
- **offline** (*bool*) – If `True`/set, psyplot will be started in offline mode without intersphinx and remote access for the help explorer
- **pwd** (*str*) – The path to the working directory to use. Note if you do not provide any `fnames` or `project`, but set the `pwd`, it will switch the `pwd` of the current GUI.
- **script** (*str*) – The path to a python script that shall be run in the GUI. If the GUI is already running, the commands will be executed in this GUI
- **command** (*str*) – Python commands that shall be run in the GUI. If the GUI is already running, the commands will be executed in this GUI
- **use_all** (*bool*) – If `True`, use all variables. Note that this is the default if the `output` is specified and not `name`
- **exec_** (*bool*) – If `True`, the main loop is entered.
- **callback** (*str*) – A unique identifier for the method that should be used if psyplot is already running. Set this parameter to `None` to avoid sending
- **openglImplementation** ({'software', 'desktop', 'gles', 'automatic'}) – OpenGL implementation to pass to Qt. Possible options are 'software', 'desktop', 'gles' and 'automatic' (which let's PyQt decide).

straditize.binary module

A module to read in and digitize the pollen diagram

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Classes

<code>BarDataReader(*args, **kwargs)</code>	A DataReader for digitizing bar pollen diagrams
<code>DataReader(image[, ax, extent, plot, ...])</code>	A class to read in and digitize the data files of the pollen diagram
<code>LineDataReader(image[, ax, extent, plot, ...])</code>	A data reader for digitizing line diagrams
<code>RoundedBarDataReader(*args, **kwargs)</code>	A bar data reader that can be used for rounded bars

Functions

<code>groupby_arr(arr)</code>	Groupby a boolean array
<code>only_parent(func)</code>	Call the given <i>func</i> only from the parent reader

`class straditiz.binary.BarDataReader(*args, **kwargs)`

Bases: `straditiz.binary.DataReader`

A DataReader for digitizing bar pollen diagrams

Compared to the base `DataReader` class, this reader implements a different strategy in digitizing and finding the samples. When digitizing the full diagram, we try to find the distinct bars using the `get_bars()` method. These bars might have to be splitted manually if they are not easy to distinguish. One key element to distinguish to adjacent bars is the specified *tolerance*.

The base class works for rectangular bars. If you require rounded bars, use the `RoundedBarDataReader`

Parameters `tolerance (int)` – If x_0 is the value in a pixel row y and x_1 the value in the next pixel row $y+1$, then the two pixel rows are considered as belonging to different bars if $\text{abs}(x_1 - x_0) > \text{tolerance}$ (see the `get_bars()` method and the `tolerance` attribute)

Methods

<code>create_grouper(ds, columns, *args, **kwargs)</code>	Create the grouper that plots the results
<code>digitize([do_split, inplace])</code>	Reimplemented to ignore the rows between the bars
<code>find_potential_samples(col[, min_len, ...])</code>	Find the bars in the column
<code>from_dataset(ds, *args, **kwargs)</code>	Create a new <code>DataReader</code> from a <code>xarray.Dataset</code>
<code>get_bars(arr[, do_split])</code>	Find the distinct bars in an array
<code>shift_vertical(pixels)</code>	Shift the columns vertically.
<code>to_dataset([ds])</code>	All the necessary data as a <code>xarray.Dataset</code>

Attributes

<code>min_fract</code>	The minimum fraction of overlap for two bars to be considered as the
<code>nc_meta</code>	<code>dict() -> new empty dictionary</code>
<code>samples_at_boundaries</code>	There should not be samples at the boundaries because the first
<code>tolerance</code>	Tolerance to distinguish bars.

`create_grouper(ds, columns, *args, **kwargs)`

Create the grouper that plots the results

Parameters

- **ds** (`xarray.Dataset`) – The dataset with the data
- **columns** (`list of int`) – The numbers of the columns for which the grouper should be created
- **fig** (`matplotlib.figure.Figure`) – The matplotlib figure to plot on
- **x0** (`float`) – The left boundary of the larger Bbox of the stratigraphic diagram
- **y0** (`int`) – The upper boundary of the larger Bbox of the stratigraphic diagram
- **width** (`float`) – The width of the final axes between 0 and 1
- **height** (`float`) – The height of the final axis between 0 and 1
- **ax0** (`matplotlib.axes.Axes`) – The larger matplotlib axes whose bounding box shall be used.
- **transformed** (`bool`) – If True, y-axes and x-axes have been translated (see the `px2data_x()` and `px2data_y()` methods)
- **colnames** (`list of str`) – The column names to use in the plot
- ****kwargs** – any other keyword argument that is passed to the `psy_strat.stratplot.StratGroup.from_dataset()` method

Returns The grouper that visualizes the given *columns* in the *fig*

Return type `psy_strat.stratplot.StratGroup`

digitize (`do_split=False, inplace=True`)

Reimplemented to ignore the rows between the bars

Parameters

- **do_split** (`bool`) – If True and a bar is 1.7 times longer than the mean, it is splitted into two.
- **inplace** (`bool`) – If True (default), the `full_df` attribute is updated. Otherwise a DataFrame is returned

find_potential_samples (`col, min_len=None, max_len=None, filter_func=None`)

Find the bars in the column

This method gets the bars in the given *col* and returns the distinct indices

Parameters

- **col** (`int`) – The column for which to find the extrema
- **min_len** (`int`) – The minimum length of one extremum. If the width of the interval where we found an extrumum is smaller than that, the extremum is ignored. If None, this parameter does not have an effect (i.e. `min_len=1`).
- **max_len** (`int`) – The maximum length of one extremum. If the width of the interval where we found an extrumum is greater than that, the extremum is ignored. If None, this parameter does not have an effect.
- **filter_func** (`function`) – A function to filter the extreme. It must accept one argument which is a list of integers representing the indices of the extremum in *a*

Returns

- *list of list of int of shape (N, 2)* – The list of N extremum locations. Each tuple in this list represents an interval a where one extremum might be located
- *list of list of int* – The excluded extremum locations that are ignored because we could not find a change of sign in the slope.

See also:

`find_samples()`

classmethod `from_dataset` (*ds, *args, **kwargs*)
Create a new *DataReader* from a `xarray.Dataset`

Parameters

- **`ds`** (`xarray.Dataset`) – The dataset that has been stored with the `to_dataset()` method
- **`*args, **kwargs`** – Any other arguments passed to the *DataReader* constructor

Returns The reader recreated from *ds*

Return type `DataReader`

get_bars (*arr, do_split=False*)
Find the distinct bars in an array

Parameters

- **`arr`** (`np.ndarray`) – The array to find the bars in
- **`do_split`** (`bool`) – If True and a bar is 1.7 times longer than the mean, it is splitted into two.

Returns

- *list of list of ints* – The list of the distinct positions of the bars
- *list of floats* – The heights for each of the bars
- *list of list of ints* – The indices of bars that are longer than 1.7 times the mean of the other bars and should be splitted. If *do_split* is True, they have been splitted already

max_len = None

min_fract = 0.9

The minimum fraction of overlap for two bars to be considered as the same sample (see `unique_bars()`)

min_len = None

nc_meta = {'bars{reader}_bars': {'dims': ('bars{reader}_bar', 'limit'), 'long_name':

samples_at_boundaries = False

There should not be samples at the boundaries because the first sample is in the middle of the first bar

shift_vertical (pixels)

Shift the columns vertically.

Parameters pixels (*list of floats*) – The y-value for each column for which to shift the values. Note that these values have to be greater than or equal to 0

to_dataset (ds=None)

All the necessary data as a `xarray.Dataset`

Parameters ds (`xarray.Dataset`) – The dataset in which to insert the data. If None, a new one will be created

Returns Either the given `ds` or a new `xarray.Dataset` instance

Return type `xarray.Dataset`

tolerance = 2

Tolerance to distinguish bars. If `x0` is the value in a pixel row `y` and `x1` the value in the next pixel row `y+1`, then the two pixel rows are considered as belonging to different bars if `abs(x1 - x0) > tolerance`

class `straditize.binary.DataReader(image, ax=None, extent=None, plot=True, children=[], parent=None, magni=None, plot_background=False, binary=None)`

Bases: `straditize.label_selection.LabelSelection`

A class to read in and digitize the data files of the pollen diagram

The source image is stored in the `image` attribute, the binary array of it is stored in the `binary` attribute. A labeled version created by the `skimage.morphology.label()` function, is stored in the `labels` attribute and can regenerated using the `reset_labels()` method.

Subclasses of this class should reimplement the `digitize()` method that digitizes the diagram, and the `find_potential_samples()` method.

There is always one parent reader stored in the `parent` attribute. This is then the reader that is accessible through the `straditize.straditizer.Straditizer.data_reader` attribute and holds the references to other readers in it's `children` attribute

Parameters

- `image (PIL.Image.Image)` – The image of the diagram
- `ax (matplotlib.axes.Axes)` – The matplotlib axes to plot on
- `extent (list)` – List of four number specifying the extent of the image in it's source. This extent will be used for the call of `matplotlib.pyplot.imshow()`
- `children (list of DataReader)` – Child readers for other columns in case the newly created instance is the parent reader
- `parent (DataReader)` – The parent reader.
- `magni (straditize.magnifier.Magnifier)` – The magnifier for the given `ax`
- `plot_background (bool)` – If True (and `plot` is True), a white, opaque area is plotted below the `plot_im`
- `binary (None)` – The binary version of the given `image`. If not provided, the `to_binary_pil()` method is used with the given `image`

Methods

<code>add_samples(samples[, rough_locs])</code>	Add samples to the found ones
<code>close()</code>	
<code>color_labels([categorize])</code>	The labels of the colored array
<code>create_exaggerations_reader(factor[, cls])</code>	Create a new exaggerations reader for this reader
<code>create_grouper(ds, columns, fig, x0, y0, ...)</code>	Create the grouper that plots the results
<code>create_variable(ds, vname, data, **kwargs)</code>	Insert the data into a variable in an <code>xr.Dataset</code>
<code>digitize([use_sum, inplace])</code>	Digitize the binary image to create the full dataframe
<code>digitize_exaggerated([fraction, absolute, ...])</code>	Merge the exaggerated values into the original digitized result
<code>disable_label_selection(*args, **kwargs)</code>	Disable the label selection

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<code>draw_figure()</code>	Draw the matplotlib <code>fig</code> and the magni figure
<code>end_column_selection()</code>	End the column selection and remove the artists
<code>estimated_column_starts([threshold])</code>	The estimated column starts as <code>numpy.ndarray</code> .
<code>find_potential_samples(col[, min_len, ...])</code>	Find potential samples in an array
<code>find_samples([min_fract, pixel_tol])</code>	Find the samples in the diagram
<code>found_extrema_per_row()</code>	Calculate how many columns have a potential sample in each pixel row
<code>from_dataset(ds, *args, **kwargs)</code>	Create a new <code>DataReader</code> from a <code>xarray.Dataset</code>
<code>get_bbox_for_cols(columns, x0, y0, width, height)</code>	Get the boundary boxes for the columns of this reader in the results
<code>get_binary_for_col(col)</code>	Get the binary array for a specific column
<code>get_cross_column_features([min_px])</code>	Get features that are contained in two or more columns
<code>get_disconnected_parts([fromlast, from0, ...])</code>	Identify parts in the binary data that are not connected
<code>get_labeled_array()</code>	Create a connectivity-based labeled array of the binary data
<code>get_occurrences()</code>	Extract the positions of the occurrences from the selection
<code>get_parts_at_column_ends([npixels])</code>	Identify parts in the binary data that touch the next column
<code>get_reader_for_col(col)</code>	Get the reader for a specific column
<code>get_surrounding_slopes(indices, arr)</code>	
<code>image_array()</code>	The RGBA values of the colored image
<code>is_obstacle(indices, arr)</code>	Check whether the found extrema is only an obstacle of the picture
<code>mark_as_exaggerations(mask)</code>	Mask the given array as exaggerated
<code>merge_close_samples(locs[, rough_locs, ...])</code>	
<code>merge_occurrences(locs)</code>	
<code>merged_binaries()</code>	Get the binary data from all children and merge them into one array
<code>merged_labels()</code>	Get the labeled binary data from all children merged into one array
<code>new_child_for_cols(columns, cls[, plot])</code>	Create a new child reader for specific columns
<code>plot_background([ax])</code>	Plot a white layer below the <code>plot_im</code>
<code>plot_color_image([ax])</code>	Plot the colored image on a matplotlib axes
<code>plot_full_df([ax])</code>	Plot the lines for the digitized diagram
<code>plot_image([ax])</code>	Plot the binary data image on a matplotlib axes
<code>plot_other_potential_samples([tol, ...])</code>	Plot potential samples that are not yet in the samples
<code>plot_potential_samples([excluded, ax, plot_kw])</code>	Plot the ranges for potential samples
<code>plot_results(df[, ax, fig, transformed])</code>	Plot the reconstructed diagram
<code>plot_sample_hlines([ax])</code>	Plot one horizontal line per sample in the <code>sample_locs</code>
<code>plot_samples([ax])</code>	Plot the diagram as lines reconstructed from the samples
<code>px2data_x(coord)</code>	Transform the pixel coordinates into data coordinates

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<code>recognize_hlines([fraction, min_lw, max_lw, ...])</code>	Recognize horizontal lines in the plot and subtract them
<code>recognize_vlines([fraction, min_lw, max_lw, ...])</code>	Recognize horizontal lines in the plot and subtract them
<code>recognize_xaxes([fraction, min_lw, max_lw, ...])</code>	Recognize (and potentially remove) x-axes at bottom and top
<code>recognize_yaxes([fraction, min_lw, max_lw, ...])</code>	Find (and potentially remove) y-axes in the image
<code>remove_in_children(arr, amask)</code>	Update the child reader images after having removed binary data
<code>remove_plots()</code>	Remove all plotted artists by this reader
<code>reset_column_starts()</code>	Reset the column starts, <code>full_df</code> , <code>shifted</code>
<code>reset_image(image[, binary])</code>	Reset the image for this straditizer
<code>reset_labels()</code>	Reset the <code>labels</code> array
<code>reset_samples()</code>	Reset the samples
<code>resize_axes(grouper, bounds)</code>	Resize the axes based on column boundaries
<code>set_as_parent()</code>	Set this instance as the parent reader
<code>set_hline_locs_from_selection([selection])</code>	Save the locations of horizontal lines
<code>set_vline_locs_from_selection([selection])</code>	Save the locations of vertical lines
<code>shift_vertical(pixels[, draw])</code>	Shift the columns vertically.
<code>show_cross_column_features([min_px, remove])</code>	Highlight and maybe remove cross column features
<code>show_disconnected_parts([fromlast, from0, ...])</code>	Highlight or remove disconnected parts
<code>show_parts_at_column_ends([npixels, remove])</code>	Highlight or remove features that touch the column ends
<code>show_small_parts([n, remove])</code>	Highlight and potentially remove small features in the image
<code>start_column_selection([use_all])</code>	Enable the user to select columns
<code>to_binary_pil(image[, threshold])</code>	Convert an image to a binary
<code>to_dataset([ds])</code>	All the necessary data as a <code>xarray.Dataset</code>
<code>to_grey_pil(image[, threshold])</code>	Convert an image to a greyscale image
<code>unique_bars([min_fract, asdict])</code>	Estimate the unique bars
<code>update_image(arr, amask)</code>	Update the image after having removed binary data
<code>update_rgba_image(arr, mask)</code>	Update the RGBA image from the given 3D-array

Attributes

<code>all_column_bounds</code>	The boundaries for the data columns
<code>all_column_ends</code>	1D numpy array with the ends for all column (including child reader)
<code>all_column_starts</code>	1D numpy array with the ends for all column (including child reader)
<code>ax</code>	The matplotlib axes where the <code>plot_im</code> is plotted on
<code>background</code>	White rectangle that represents the background of the binary image.
<code>binary</code>	A 2D numpy array representing the binary version of the image
<code>children</code>	Child readers for specific columns.

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<code>column_bounds</code>	The boundaries for the data columns
<code>column_ends</code>	1D numpy array with the ends for each column of this reader
<code>column_starts</code>	1D numpy array with the starts for each column of this reader
<code>columns</code>	The indices of the columns that are handled by this reader
<code>exaggerated_reader</code>	The reader that represents the exaggerations
<code>extent</code>	The extent of the <code>plot_im</code>
<code>fig</code>	The matplotlib figure of the <code>ax</code>
<code>full_df</code>	The full <code>pandas.DataFrame</code> of the digitized image
<code>hline_locs</code>	<code>list</code> or floats. The indexes of horizontal lines
<code>image</code>	<code>PIL.Image.Image</code> of the diagram part with mode RGBA
<code>is_exaggerated</code>	Exaggeration factor that is not 0 if this reader represents exaggeration
<code>iter_all_readers</code>	Iter through the parent reader and it's children
<code>label_arrs</code>	Built-in mutable sequence.
<code>labels</code>	A connectivity-based labeled version of the binary data
<code>magni</code>	the <code>straditize.magnifier.Magnifier</code> for the <code>ax</code>
<code>magni_background</code>	White rectangle that represents the background of the binary image in the magnifier.
<code>magni_plot_im</code>	magnified <code>plot_im</code>
<code>min_fract</code>	The minimum fraction of overlap for two bars to be considered as the
<code>nc_meta</code>	A mapping from variable name to meta information
<code>non_exaggerated_reader</code>	The reader that represents the exaggerations
<code>num_labels</code>	The maximum label in the <code>labels</code> array
<code>occurrences</code>	A set of tuples marking the position of an occurrence
<code>occurrences_dict</code>	A mapping from column number to an numpy array with the indices of
<code>occurrences_value</code>	The value that is given to the occurrences in the measurements
<code>parent</code>	Parent reader for this instance.
<code>plot_im</code>	the matplotlib image artist
<code>rough_locs</code>	The <code>pandas.DataFrame</code> with rough locations for the samples.
<code>sample_locs</code>	The <code>pandas.DataFrame</code> with locations and values of the
<code>samples_at_boundaries</code>	a boolean flag that shall indicate if we assume that the first and last
<code>shifted</code>	The number of pixels the columns have been shifted
<code>strat_plot_identifier</code>	<code>str(object=") -> str</code>
<code>vline_locs</code>	<code>list</code> or floats. The indexes of vertical lines
<code>xaxis_px</code>	The x indices in column pixel coordinates that are used for x-axes

`add_samples(samples, rough_locs=None)`

Add samples to the found ones

Parameters

- **`samples`** (*series, 1d-array or DataFrame*) – The samples. If it is series, we assume that the index represents the y-value of the sample and the value the x-position (see `xcolumns`). In case of a 1d-array, we assume that the data represents the y-values of the samples. In case of a DataFrame, we assume that the columns correspond to columns in the `full_df` attribute and are True where we have a sample.

Note that the y-values must be in image coordinates (see `extent` attribute).

- **`rough_locs`** (*DataFrame*) – The rough locations of the new samples (see the `rough_locs` attribute)

See also:

`samples()`, `rough_locs()`, `find_samples()`, `sample_locs()`

`property all_column_bounds`

The boundaries for the data columns

`property all_column_ends`

1D numpy array with the ends for all column (including child reader)

See also:

`all_column_starts` The starts for all column

`all_column_bounds` The (start, end)-tuple for all of the columns

`column_ends` The ends for this specific reader

reader

`property all_column_starts`

1D numpy array with the ends for all column (including child reader)

See also:

`all_column_ends` The ends for all column

`all_column_bounds` The (start, end)-tuple for all of the columns

`column_starts` The starts for this specific reader

reader

`ax = None`

The matplotlib axes where the `plot_im` is plotted on

`background = None`

White rectangle that represents the background of the binary image. This is only plotted by the parent reader

`binary = None`

A 2D numpy array representing the binary version of the `image`

`children = []`

Child readers for specific columns. Is not empty if and only if the `parent` attribute is this instance

`close()`

`color_labels (categorize=1)`

The labels of the colored array

property column_bounds

The boundaries for the data columns

property column_ends

1D numpy array with the ends for each column of this reader

See also:

[column_starts](#) The starts for each column

[column_bounds](#) The (start, end)-tuple for each of the columns

[all_column_ends](#) The ends for all columns, including child

reader

property column_starts

1D numpy array with the starts for each column of this reader

See also:

[column_ends](#) The ends for each column

[column_bounds](#) The (start, end)-tuple for each of the columns

[all_column_starts](#) The starts for all columns, including child

reader

property columns

The indices of the columns that are handled by this reader

create_exaggerations_reader(factor, cls=None)

Create a new exaggerations reader for this reader

Parameters

- **factor** (*float*) – The exaggeration factor
- **cls** (*type*) – The [DataReader](#) subclass

Returns The new exaggerated reader

Return type instance of *cls*

create_grouper(ds, columns, fig, x0, y0, width, height, ax0=None, transformed=True, colnames=None, **kwargs)

Create the grouper that plots the results

Parameters

- **ds** (*xarray.Dataset*) – The dataset with the data
- **columns** (*list of int*) – The numbers of the columns for which the grouper should be created
- **fig** (*matplotlib.figure.Figure*) – The matplotlib figure to plot on
- **x0** (*float*) – The left boundary of the larger Bbox of the stratigraphic diagram
- **y0** (*int*) – The upper boundary of the larger Bbox of the stratigraphic diagram
- **width** (*float*) – The width of the final axes between 0 and 1
- **height** (*float*) – The height of the final axis between 0 and 1

- **ax0** (`matplotlib.axes.Axes`) – The larger matplotlib axes whose bounding box shall be used.
- **transformed** (`bool`) – If True, y-axes and x-axes have been translated (see the `px2data_x()` and `px2data_y()` methods)
- **colnames** (`list of str`) – The column names to use in the plot
- ****kwargs** – any other keyword argument that is passed to the `psy_strat.stratplot.StratGroup.from_dataset()` method

Returns The grouper that visualizes the given *columns* in the *fig*

Return type `psy_strat.stratplot.StratGroup`

create_variable (*ds*, *vname*, *data*, ****kwargs**)

Insert the data into a variable in an `xr.Dataset`

Parameters

- **ds** (`xarray.Dataset`) – The destination dataset
- **vname** (`str`) – The name of the variable in the `nc_meta` mapping. This name might include `{reader}` which will then be replaced by the number of the reader in the `iter_all_readers` attribute
- **data** (`np.ndarray`) – The numpy array to store in the variable specified by *vname*
- ****kwargs** – A mapping from dimension to slicer that should be used to slice the dataset

Returns The resolved *vname* that has been used in the dataset

Return type `str`

digitize (*use_sum=False*, *inplace=True*)

Digitize the binary image to create the full dataframe

Parameters

- **use_sum** (`bool`) – If True, the sum of cells that are not background are used for each column, otherwise the value of the cell is used that has the maximal distance to the column start for each row
- **inplace** (`bool`) – If True (default), the `full_df` attribute is updated. Otherwise a DataFrame is returned

Returns The digitization result if *inplace* is True, otherwise None

Return type None or `pandas.DataFrame`

digitize_exaggerated (*fraction=0.05*, *absolute=8*, *inplace=True*, *return_mask=False*)

Merge the exaggerated values into the original digitized result

Parameters

- **fraction** (*float between 0 and 1*) – The fraction under which the exaggerated data should be used. Set this to 0 to ignore it.
- **absolute** (`int`) – The absolute value under which the exaggerated data should be used. Set this to 0 to ignore it.
- **inplace** (`bool`) – If True (default), the `full_df` attribute is updated. Otherwise a DataFrame is returned
- **return_mask** (`bool`) – If True, a boolean 2D array is returned indicating where the exaggerations have been used

Returns

- *pandas.DataFrame or None* – If *inplace* is False, the digitized result. Otherwise, if *return_mask* is True, the mask where the exaggerated results have been used. Otherwise None
- *pandas.DataFrame, optionally* – If *inplace* is False and *return_mask* is True, a pandas.DataFrame containing the boolean mask where the exaggerated results have been used. Otherwise, this is skipped

disable_label_selection(*args, **kwargs)

Disable the label selection

This will disconnect the *pick_event* and remove the selection images**Parameters** **remove** (*bool*) – Whether to remove the selection image from the plot. If None, the *_remove* attribute is used**See also:**`enable_label_selection()`, `remove_selected_labels()`**draw_figure**()Draw the matplotlib *fig* and the *magni* figure**end_column_selection**()

End the column selection and remove the artists

estimated_column_starts(*threshold=None*)The estimated column starts as `numpy.ndarray`.We assume a new column a pixel column i if

1. the previous pixel column $i-1$ did not contain any data ($D(i-1) = 0$)
2. THE amount of data points doubled compared to $i-1$ ($D(i) \geq 2 \cdot D(i-1)$)
3. the amount of data points steadily increases within the next few columns to a value twice as large as the previous column ($D(i+n) \geq 2 \cdot D(i-1)$ with $n > 0$ and $D(i+j) \geq D(i)$ for all $0 < j \leq n$)

Each potential column starts must also be covered by a given *threshold*.**Parameters** **threshold** (*float between 0 and 1*) – The fraction that has to be covered to assume a valid column start. By default, 0.1 (i.e. 10 percent)**Returns** The starts for each column**Return type** `np.ndarray`**property exaggerated_reader**

The reader that represents the exaggerations

property extentThe extent of the *plot_im***property fig**The matplotlib figure of the *ax***find_potential_samples**(*col, min_len=None, max_len=None, filter_func=None*)

Find potential samples in an array

This method finds extrema in an array and returns the indices where the extremum might be. The algorithm thereby filters out obstacles by first going over the array, making sure, that there is a change of sign in the slope in the found extremum, and if not, ignores it and flattens it out.

Parameters

- **col** (*int*) – The column for which to find the extrema
- **min_len** (*int*) – The minimum length of one extremum. If the width of the interval where we found an extrum is smaller than that, the extremum is ignored. If None, this parameter does not have an effect (i.e. `min_len=1`).
- **max_len** (*int*) – The maximum length of one extremum. If the width of the interval where we found an extrum is greater than that, the extremum is ignored. If None, this parameter does not have an effect.
- **filter_func** (*function*) – A function to filter the extreme. It must accept one argument which is a list of integers representing the indices of the extremum in *a*

Returns

- *list of list of int of shape (N, 2)* – The list of N extremum locations. Each tuple in this list represents an interval *a* where one extremum might be located
- *list of list of int* – The excluded extremum locations that are ignored because we could not find a change of sign in the slope.

See also:

`find_samples()`

find_samples (*min_fract=None*, *pixel_tol=5*, **args*, ***kwargs*)

Find the samples in the diagram

This function finds the samples using the `find_potential_samples()` function. It combines the found extrema from all columns and estimates the exact location using an interpolation of the slope

Parameters

- **min_fract** (*float*) – The minimum fraction between 0 and 1 that two bars have to overlap such that they are considered as representing the same sample. If None, the `min_fract` attribute is used
- **min_len** (*int*) – The minimum length of one extremum. If the width of the interval where we found an extrum is smaller than that, the extremum is ignored. If None, this parameter does not have an effect (i.e. `min_len=1`).
- **max_len** (*int*) – The maximum length of one extremum. If the width of the interval where we found an extrum is greater than that, the extremum is ignored. If None, this parameter does not have an effect.
- **filter_func** (*function*) – A function to filter the extreme. It must accept one argument which is a list of integers representing the indices of the extremum in *a*

Returns

- *pandas.DataFrame* – The x- and y-locations of the samples. The index is the y-location, the columns are the columns in the `full_df`.
- *pandas.DataFrame* – The rough locations of the samples. The index is the y-location of the columns, the values are lists of the potential sample locations.

found_extrema_per_row()

Calculate how many columns have a potential sample in each pixel row

Returns A series with one entry per pixel row. The values are the number of columns in the diagram that have a potential sample noted in the `rough_locs`

Return type `pandas.Series`

classmethod from_dataset (*ds, *args, **kwargs*)
Create a new *DataReader* from a *xarray.Dataset*

Parameters

- **ds** (*xarray.Dataset*) – The dataset that has been stored with the *to_dataset()* method
- ***args, **kwargs** – Any other arguments passed to the *DataReader* constructor

Returns The reader recreated from *ds*

Return type *DataReader*

property full_df
The full *pandas.DataFrame* of the digitized image

get_bbox_for_cols (*columns, x0, y0, width, height*)
Get the boundary boxes for the columns of this reader in the results plot

This method is used by the *plot_results()* method to get the Bbox for a *psy_strat.stratplot.StratGroup* grouper

Parameters

- **columns** (*list of int*) – The column numbers to use
- **x0** (*float*) – The left boundary of the larger Bbox of the stratigraphic diagram
- **y0** (*int*) – The upper boundary of the larger Bbox of the stratigraphic diagram
- **width** (*float*) – The width of the final axes between 0 and 1
- **height** (*float*) – The height of the final axis between 0 and 1

Returns The boundary box for the given *columns* in the matplotlib figure

Return type *matplotlib.transforms.Bbox*

See also:

plot_results()

get_binary_for_col (*col*)
Get the binary array for a specific column

get_cross_column_features (*min_px=50*)
Get features that are contained in two or more columns

Parameters *min_px* (*int*) – The number of pixels that have to be contained in each column

Returns The 2D boolean mask with the same shape as the *binary* array that is True if a data pixel is considered as to belong to a cross column feature

Return type *np.ndarray* of dtype bool

get_disconnected_parts (*fromlast=5, from0=10, cross_column=False*)
Identify parts in the *binary* data that are not connected

Parameters

- **fromlast** (*int*) – A pixel $x_1 > x_0$ is considered as disconnected, if it is at least $x_1 - x_0 \geq \text{fromlast}$. If this is 0, it is ignored and only *from0* is considered.
- **from0** (*int*) – A pixel is considered as disconnected if it is more than *from0* pixels away from the column start. If this is 0, it is ignored and only *fromlast* is considered

- **`cross_column (bool)`** – If False, disconnected features are only marked in the column where the disconnection has been detected. Otherwise the entire feature is marked

Returns The 2D boolean mask with the same shape as the `binary` array that is True if a data pixel is considered as to be disconnected

Return type np.ndarray of dtype bool

`get_labeled_array()`

Create a connectivity-based labeled array of the `binary` data

`get_occurrences()`

Extract the positions of the occurrences from the selection

`get_parts_at_column_ends (npixels=2)`

Identify parts in the `binary` data that touch the next column

Parameters `npixels (int)` – If a data pixel is less than `npixels` away from the column end, it is considered to be at the column end and marked

Returns A boolean mask with the same shape as the `binary` data that is True where a pixel is considered to be at the column end

Return type np.ndarray of dtype bool

`get_reader_for_col (col)`

Get the reader for a specific column

Parameters `col (int)` – The column of interest

Returns Either the reader or None if no reader could be found

Return type `DataReader` or None

`get_surrounding_slopes (indices, arr)`

`hline_locs = None`

`list` or floats. The indexes of horizontal lines

`image = None`

PIL.Image.Image of the diagram part with mode RGBA

`image_array()`

The RGBA values of the colored image

`is_exaggerated = 0`

Exaggeration factor that is not 0 if this reader represents exaggeration plots

`is_obstacle (indices, arr)`

Check whether the found extrema is only an obstacle of the picture

`property iter_all_readers`

Iter through the `parent` reader and it's `children`

`label_arcs = ['binary', 'labels', 'image_array']`

`labels = None`

A connectivity-based labeled version of the `binary` data

`magni = None`

the `straditize.magnifier.Magnifier` for the `ax`

`magni_background = None`

White rectangle that represents the background of the binary image in the magnifier. This is only plotted by the parent reader

`magni_color_plot_im = None`

`magni_plot_im = None`
magnified `plot_im`

`mark_as_exaggerations (mask)`

Mask the given array as exaggerated

Parameters `mask` (*2D np.ndarray of dtype bool*) – A mask with the same shape as the `binary` array that is True if a cell should be interpreted as the visualization of an exaggeration

`merge_close_samples (locs, rough_locs=None, pixel_tol=5)`

`merge_occurrences (locs)`

`merged_binaries ()`

Get the binary data from all children and merge them into one array

Returns The binary image with the same shape as the `binary` data

Return type np.ndarray of dtype int

`merged_labels ()`

Get the labeled binary data from all children merged into one array

Returns The labeled binary image with the same shape as the `label` data

Return type np.ndarray of dtype int

`min_fract = 0.9`

The minimum fraction of overlap for two bars to be considered as the same sample (see `unique_bars ()`)

`nc_meta = {'binary': {'dims': ('reader', 'ydata', 'xdata'), 'long_name': 'Binary im...}}`
A mapping from variable name to meta information

`new_child_for_cols (columns, cls, plot=True)`

Create a new child reader for specific columns

Parameters

- `columns` (*list of int*) – The columns for the new reader
- `cls` (*type*) – The `DataReader` subclass
- `plot` (`bool`) – Plot the binary image

Returns The new reader for the specified `columns`

Return type instance of `cls`

`property non_exaggerated_reader`

The reader that represents the exaggerations

`property num_labels`

The maximum label in the `labels` array

`property occurrences`

A set of tuples marking the position of an occurrence

An occurrence, motivated by pollen diagrams, just highlights the existence at a certain point without giving the exact value. In pollen diagrams, these are usually taxa that were found but have a percentage of less than 0.5 %.

This set of tuples (x, y) contains the coordinates of the occurrences. The first value in each tuple is the y-value, the second the x-value.

See also:

`occurrences_dict` A mapping from column number to occurrences

property `occurrences_dict`

A mapping from column number to an numpy array with the indices of an occurrence

`occurrences_value = -9999`

The value that is given to the occurrences in the measurements

`parent = None`

Parent reader for this instance. Might be the instance itself

`plot_background(ax=None, **kwargs)`

Plot a white layer below the `plot_im`

Parameters

- `ax (matplotlib.axes.Axes)` – The matplotlib axes to plot on. If not given, the `ax` attribute is used
- `**kwargs` – Any other keyword that is given to the `matplotlib.pyplot.imshow()` function

`plot_color_image(ax=None, **kwargs)`

Plot the colored `image` on a matplotlib axes

Parameters

- `ax (matplotlib.axes.Axes)` – The matplotlib axes to plot on. If not given, the `ax` attribute is used
- `**kwargs` – Any other keyword that is given to the `matplotlib.pyplot.imshow()` function

`plot_full_df(ax=None, *args, **kwargs)`

Plot the lines for the digitized diagram

Parameters

- `ax (matplotlib.axes.Axes)` – The matplotlib axes to plot on
- `*args, **kwargs` – Any other argument and keyword argument that is passed to the `matplotlib.pyplot.plot()` function

`plot_im = None`

the matplotlib image artist

`plot_image(ax=None, **kwargs)`

Plot the `binary` data image on a matplotlib axes

Parameters

- `ax (matplotlib.axes.Axes)` – The matplotlib axes to plot on. If not given, the `ax` attribute is used and (if this is None, too) a new figure is created
- `**kwargs` – Any other keyword that is given to the `matplotlib.pyplot.imshow()` function

`plot_other_potential_samples(tol=1, already_found=None, *args, **kwargs)`

Plot potential samples that are not yet in the `samples` attribute

Parameters

- **tol** (`int`) – The pixel tolerance for a sample. If the distance between a potential sample and all already existing sample is greater than tolerance, the potential sample will be plotted
- **already_found** (`np.ndarray`) – The pixel rows of samples that have already been found. If not specified, the index of the `sample_locs` is used
- **excluded** (`bool`) – If True, plot the excluded samples instead of the included samples (see the return values in `find_potential_samples()`)
- **ax** (`matplotlib.axes.Axes`) – The matplotlib axes to plot on
- **plot_kws** (`dict`) – Any other keyword argument that is passed to the `matplotlib.pyplot.plot()` function. By default, this is equal to `{'marker': '+'}`
- **min_len** (`int`) – The minimum length of one extremum. If the width of the interval where we found an extrumum is smaller than that, the extremum is ignored. If None, this parameter does not have an effect (i.e. `min_len=1`).
- **max_len** (`int`) – The maximum length of one extremum. If the width of the interval where we found an extrumum is greater than that, the extremum is ignored. If None, this parameter does not have an effect.
- **filter_func** (`function`) – A function to filter the extreme. It must accept one argument which is a list of integers representing the indices of the extremum in `a`

`plot_potential_samples(excluded=False, ax=None, plot_kws={}, *args, **kwargs)`

Plot the ranges for potential samples

This method plots the rough locations of potential samples (see `find_potential_samples()`)

Parameters

- **excluded** (`bool`) – If True, plot the excluded samples instead of the included samples (see the return values in `find_potential_samples()`)
- **ax** (`matplotlib.axes.Axes`) – The matplotlib axes to plot on
- **plot_kws** (`dict`) – Any other keyword argument that is passed to the `matplotlib.pyplot.plot()` function. By default, this is equal to `{'marker': '+'}`
- **min_len** (`int`) – The minimum length of one extremum. If the width of the interval where we found an extrumum is smaller than that, the extremum is ignored. If None, this parameter does not have an effect (i.e. `min_len=1`).
- **max_len** (`int`) – The maximum length of one extremum. If the width of the interval where we found an extrumum is greater than that, the extremum is ignored. If None, this parameter does not have an effect.
- **filter_func** (`function`) – A function to filter the extreme. It must accept one argument which is a list of integers representing the indices of the extremum in `a`

`plot_results(df, ax=None, fig=None, transformed=True)`

Plot the reconstructed diagram

This method plots the reconstructed diagram using the psy-strat module.

Parameters

- **df** (`pandas.DataFrame`) – The data to plot. E.g. the `sample_locs` or the `straditize.straditizer.Straditizer.final_df` data

- **ax** (`matplotlib.axes.Axes`) – The axes to plot on. If None, a new one is created inside the given `fig`
- **fig** (`matplotlib.figure.Figure`) – The matplotlib figure to plot on. If not given, the current figure (see `matplotlib.pyplot.gcf()`) is used
- **transformed** (`bool`) – If True, y-axes and x-axes have been translated (see the `px2data_x()` and `px2data_y()` methods)

Returns

- `psyplot.project.Project` – The newly created psyplot project with the plotters
- list of `psy_strat.stratplot.StratGroup` instances – The groupers for the different columns

plot_sample_hlines (`ax=None, **kwargs`)Plot one horizontal line per sample in the `sample_locs`**Parameters**

- **ax** (`matplotlib.axes.Axes`) – The matplotlib axes to plot on
- ***args, **kwargs** – Any other keyword argument that is passed to the `matplotlib.pyplot.hlines()` function

plot_samples (`ax=None, *args, **kwargs`)

Plot the diagram as lines reconstructed from the samples

Parameters

- **ax** (`matplotlib.axes.Axes`) – The matplotlib axes to plot on
- ***args, **kwargs** – Any other argument and keyword argument that is passed to the `matplotlib.pyplot.plot()` function

px2data_x (`coord`)

Transform the pixel coordinates into data coordinates

Parameters `coord` (`1D np.ndarray`) – The coordinate values in pixels**Returns** The numpy array starting from 0 with transformed coordinates**Return type** `np.ndarray`**Notes**

Since the x-axes for stratigraphic plots are usually interrupted, the return values here are relative and therefore always start from 0

recognize_hlines (`fraction=0.3, min_lw=1, max_lw=None, remove=False, **kwargs`)

Recognize horizontal lines in the plot and subtract them

This method removes horizontal lines in the data diagram, i.e. rows whose non-background cells cover at least the specified `fraction` of the row.

Parameters

- **fraction** (`float`) – The fraction (between 0 and 1) that has to be covered to recognize a horizontal line
- **min_lw** (`int`) – The minimum line width for a line
- **max_lw** (`int`) – The maximum line width for a line or None if it should be ignored

- **remove** (`bool`) – If True, they will be removed immediately, otherwise they are displayed using the `enable_label_selection()` method and can be removed through the `remove_selected_labels()` method

Other Parameters “`kwargs`“** – Additional keywords are parsed to the `enable_label_selection()` method in case `remove` is False

Notes

This method has to be called before the `digitize()` method!

recognize_vlines (`fraction=0.3, min_lw=1, max_lw=None, remove=False, **kwargs`)

Recognize horizontal lines in the plot and subtract them

This method removes horizontal lines in the data diagram, i.e. rows whose non-background cells cover at least the specified `fraction` of the row.

Parameters

- **fraction** (`float`) – The fraction (between 0 and 1) that has to be covered to recognize a horizontal line
- **min_lw** (`int`) – The minimum line width for a line
- **max_lw** (`int`) – The maximum line width for a line or None if it should be ignored
- **remove** (`bool`) – If True, they will be removed immediately, otherwise they are displayed using the `enable_label_selection()` method and can be removed through the `remove_selected_labels()` method

Other Parameters “`kwargs`“** – Additional keywords are parsed to the `enable_label_selection()` method in case `remove` is False

Notes

This method should be called before the column starts are set

recognize_xaxes (`fraction=0.3, min_lw=1, max_lw=None, remove=False, **kwargs`)

Recognize (and potentially remove) x-axes at bottom and top

Parameters

- **fraction** (`float`) – The fraction (between 0 and 1) that has to be covered to recognize an x-axis
- **min_lw** (`int`) – The minimum line width of an axis
- **max_lw** (`int`) – The maximum line width of an axis. If not specified, it will be ignored
- **remove** (`bool`) – If True, they will be removed immediately, otherwise they are displayed using the `enable_label_selection()` method and can be removed through the `remove_selected_labels()` method

recognize_yaxes (`fraction=0.3, min_lw=0, max_lw=None, remove=False`)

Find (and potentially remove) y-axes in the image

Parameters

- **fraction** (`float`) – The fraction (between 0 and 1) that has to be covered to recognize a y-axis
- **min_lw** (`int`) – The minimum line width of an axis

- **max_lw** (`int`) – The maximum line width of an axis. If not specified, the median if the axes widths is taken
- **remove** (`bool`) – If True, they will be removed immediately, otherwise they are displayed using the `enable_label_selection()` method and can be removed through the `remove_selected_labels()` method

remove_in_children (`arr, amask`)

Update the child reader images after having removed binary data

Calls the `update_image()` and `update_rgba_image()` methods for all `children`

remove_plots ()

Remove all plotted artists by this reader

reset_column_starts ()

Reset the column starts, `full_df`, `shifted` and `occurrences`

reset_image (`image, binary=False`)

Reset the image for this straditizer

Parameters

- **image** (`PIL.Image.Image`) – The new image
- **binary** (`bool`) – If True, then the `image` is considered as the binary image and the `image` attribute is not touched

reset_labels ()

Reset the `labels` array

reset_samples ()

Reset the samples

resize_axes (`grouper, bounds`)

Resize the axes based on column boundaries

This method sets the x-limits for the different columns to the given `bounds` and resizes the axes

Parameters

- **grouper** (`psy_strat.stratplot.StratGroup`) – The grouper that manages the plot
- **bounds** (`np.ndarray of shape (N, 2)`) – The boundaries for the columns handled by the `grouper`

property rough_locs

The `pandas.DataFrame` with rough locations for the samples. It has one row per sample in the `sample_locs` dataframe and `ncols * 2` columns, where `ncols` is the number of columns in the `sample_locs`.

If the potential sample `sample_locs.iloc[i, col]` ranges `j` to `k` (see the `find_potential_samples()` method), the cell at `rough_locs.iloc[i, col * 2]` specifies the first y-pixel (`j`) and `rough_locs.iloc[i, col * 2 + 1]` the last y-pixel (+1), i.e. `k` where this sample might be located

property sample_locs

The `pandas.DataFrame` with locations and values of the samples

samples_at_boundaries = True

a boolean flag that shall indicate if we assume that the first and last rows shall be a sample if they contain non-zero values

```
set_as_parent()
    Set this instance as the parent reader

set_hline_locs_from_selection(selection=None)
    Save the locations of horizontal lines

    This methods takes every pixel row in the hline_locs attribute where at least 30% is selected. The
    digitize method will interpolate at these indices.

set_vline_locs_from_selection(selection=None)
    Save the locations of vertical lines

    This methods takes every pixel column in the vline_locs attribute where at least 30% is selected.

shift_vertical(pixels, draw=True)
    Shift the columns vertically.
```

Parameters

- **pixels** (*list of floats*) – The y-value for each column for which to shift the values. Note that these values have to be greater than or equal to 0
- **draw** (`bool`) – If True, the `ax` is drawn at the end

```
shifted = None
```

The number of pixels the columns have been shifted

```
show_cross_column_features(min_px=50, remove=False, **kwargs)
    Highlight and maybe remove cross column features
```

Parameters

- **min_px** (`int`) – The number of pixels that have to be contained in each column
- **remove** (`bool`) – If True, remove the data in the `binary` array, etc. If False, the `enable_label_selection()` method is envoked and the user can select the features to remove
- **select_all** (`bool`) – If True and `remove` is False, all labels in `arr` will be selected and the given `selection` is ignored
- **selection** (*np.ndarray of dtype bool*) – A boolean mask with the same shape as `arr` that is True where a pixel should be selected. If `remove` is True, only this mask will be used.
- **img** (*matplotlib image*) – The image for the selection. If not provided, a new image is created
- **set_picker** (`bool`) – If True, connect the `matplotlib` `pick_event` to the `pick_label()` method

```
show_disconnected_parts(fromlast=5, from0=10, remove=False, **kwargs)
    Highlight or remove disconnected parts
```

Parameters

- `% (DataReader.get_disconnected_parts.parameters.
fromlast|from0)s -`
- `% (DataReader._show_parts2remove.parameters.no_arr)s -`

```
show_parts_at_column_ends(npixels=2, remove=False, **kwargs)
    Highlight or remove features that touch the column ends
```

Parameters

- `% (DataReader.get_parts_at_column_ends.parameters)s -`
- `% (DataReader._show_parts2remove.parameters.no_arr)s -`

`show_small_parts (n=10, remove=False, **kwargs)`
Highlight and potentially remove small features in the image

Parameters

- `n (int)` – The maximal size of a feature to be considered as small
- `remove (bool)` – If True, remove the data in the `binary` array, etc. If False, the `enable_label_selection()` method is envoked and the user can select the features to remove
- `select_all (bool)` – If True and `remove` is False, all labels in `arr` will be selected and the given `selection` is ignored
- `selection (np.ndarray of dtype bool)` – A boolean mask with the same shape as `arr` that is True where a pixel should be selected. If `remove` is True, only this mask will be used.
- `img (matplotlib image)` – The image for the selection. If not provided, a new image is created
- `set_picker (bool)` – If True, connect the matplotlib `pick_event` to the `pick_label()` method

See also:

`skimage.morphology.remove_small_objects()`

`start_column_selection (use_all=False)`
Enable the user to select columns

Parameters `use_all (bool)` – If True, all columns can be selected. Otherwise only the columns in the `columns` attribute can be selected

`strat_plot_identifier = 'percentages'`
`static to_binary_pil (image, threshold=690)`
Convert an image to a binary

Parameters

- `image (PIL.Image.Image)` – The RGBA image file
- `threshold (float)` – If the multiplied RGB values in a cell are above the threshold, the cell is regarded as background and will be set to 0

Returns The binary image of integer type

Return type np.ndarray of ndim 2

`to_dataset (ds=None)`
All the necessary data as a `xarray.Dataset`

Parameters `ds (xarray.Dataset)` – The dataset in which to insert the data. If None, a new one will be created

Returns Either the given `ds` or a new `xarray.Dataset` instance

Return type `xarray.Dataset`

`static to_grey_pil (image, threshold=690)`
Convert an image to a greyscale image

Parameters

- **image** (*PIL.Image.Image*) – The RGBA image file
- **threshold** (*float*) – If the multiplied RGB values in a cell are above the threshold, the cell is regarded as background and will be set to 0

Returns The greyscale image of integer type

Return type np.ndarray of ndim 2

unique_bars (*min_fract=None, asdict=True, *args, **kwargs*)

Estimate the unique bars

This method puts the overlapping bars of the different columns together

Parameters

- **min_fract** (*float*) – The minimum fraction between 0 and 1 that two bars have to overlap such that they are considered as representing the same sample. If None, the *min_fract* attribute is used
- **asdict** (*bool*) – If True, dictionaries are returned

Returns A list of the bar locations. If asdict is True (default), each item in the returned list is a dictionary whose keys are the column indices and whose values are the indices for the corresponding column. Otherwise, a list of *_Bar* objects is returned

Return type list

update_image (*arr, amask*)

Update the image after having removed binary data

This method is in the *remove_callbacks* mapping and is called after a pixel has been removed from the *binary* data. It mainly just calls the *reset_labels()* method and updates the plot

update_rgba_image (*arr, mask*)

Update the RGBA image from the given 3D-array

This method is in the *remove_callbacks* mapping and is called after a pixel has been removed from the *binary* data. It updates the *image* attribute

Parameters

- **arr** (3D np.ndarray of dtype float) – The image array
- **mask** (boolean mask of the same shape as *arr*) – The mask of features that shall be set to 0 in *arr*

vline_locs = None

list or floats. The indexes of vertical lines

xaxis_data = None

property xaxis_px

The x indices in column pixel coordinates that are used for x-axes translations

class straditize.binary.**LineDataReader** (*image, ax=None, extent=None, plot=True, children=[], parent=None, magni=None, plot_background=False, binary=None*)

Bases: *straditize.binary.DataReader*

A data reader for digitizing line diagrams

This class does not have a significantly different behaviour than the base *DataReader* class, but might be improved with more specific features in the future

Parameters

- **image** (`PIL.Image.Image`) – The image of the diagram
- **ax** (`matplotlib.axes.Axes`) – The matplotlib axes to plot on
- **extent** (`list`) – List of four number specifying the extent of the image in it's source. This extent will be used for the call of `matplotlib.pyplot.imshow()`
- **children** (list of `DataReader`) – Child readers for other columns in case the newly created instance is the parent reader
- **parent** (`DataReader`) – The parent reader.
- **magni** (`straditize.magnifier.Magnifier`) – The magnifier for the given `ax`
- **plot_background** (`bool`) – If True (and `plot` is True), a white, opaque area is plotted below the `plot_im`
- **binary** (`None`) – The binary version of the given `image`. If not provided, the `to_binary_pil()` method is used with the given `image`

Attributes

<code>strat_plot_identifier</code>	<code>str(object=") -> str</code>
------------------------------------	--------------------------------------

```
strat_plot_identifier = 'default'

class straditize.binary.RoundedBarDataReader(*args, **kwargs)
    Bases: straditize.binary.BarDataReader
```

A bar data reader that can be used for rounded bars

Parameters tolerance (`int`) – If `x0` is the value in a pixel row `y` and `x1` the value in the next pixel row `y+1`, then the two pixel rows are considered as belonging to different bars if `abs(x1 - x0) > tolerance` (see the `get_bars()` method and the `tolerance` attribute)

Attributes

<code>tolerance</code>	<code>int([x]) -> integer</code>
------------------------	-------------------------------------

```
tolerance = 10

straditize.binary.groupby_arr(arr)
    Groupby a boolean array

Parameters arr (np.ndarray of ndim 1 of dtype bool) – An array that can be converted to a numeric array
```

Returns

- **keys** (`np.ndarray`) – The keys in the array
- **starts** (`np.ndarray`) – The index of the first element that correspond to the key in `keys`

```
straditize.binary.only_parent(func)
    Call the given func only from the parent reader
```

straditize.colnames module

Module for text recognition

Disclaimer

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Classes

<i>Bbox</i>	A bounding box for a column name
<i>ColNamesReader</i> (image, bounds[, rotate, ...])	A class to recognize the text in an image

class straditize.colnames.**Bbox**
Bases: straditize.colnames._Bbox

A bounding box for a column name

Create new instance of _Bbox(x, y, w, h) **Attributes**

<i>bottom</i>	The bottom of the box
<i>bounds</i>	A list [x, y, width, height]
<i>corners</i>	A np.ndarray of shape (4, 2) with the corners of the box
<i>crop_extents</i>	The extents necessary for PIL.Image.crop
<i>extents</i>	A list [x0, x1, y0, y1] with x0 <= x1 and y0 <= y1
<i>height</i>	The (positive) height
<i>left</i>	The left edge of the box
<i>right</i>	The right edge of the box
<i>top</i>	The top of the box
<i>width</i>	The (positive) width
<i>x0</i>	The left edge
<i>x1</i>	The right edge
<i>y0</i>	The lower (bottom) edge
<i>y1</i>	The upper (top) edge

Methods

<i>from_dict</i> (d)	Construct a box from the dictionary
----------------------	-------------------------------------

property bottom
The bottom of the box

property bounds
A list [x, y, width, height]

property corners
A np.ndarray of shape (4, 2) with the corners of the box

```

property crop_extents
    The extents necessary for PIL.Image.crop

property extents
    A list [x0, x1, y0, y1] with x0 <= x1 and y0 <= y1

classmethod from_dict(d)
    Construct a box from the dictionary

property height
    The (positive) height

property left
    The left edge of the box

property right
    The right edge of the box

property top
    The top of the box

property width
    The (positive) width

property x0
    The left edge

property x1
    The right edge

property y0
    The lower (bottom) edge

property y1
    The upper (top) edge

class straditize.colnames.ColNamesReader(image, bounds, rotate=45, mirror=False,
                                         flip=False, highres_image=None,
                                         data_ylim=None)
Bases: object

```

A class to recognize the text in an image

This object handles the column names in the `column_names` attribute. It also implements several algorithms to automatically read in the column names using the tesserocr package. In particular these are the `recognize_text()` method to read in one small image and the `find_colnames()` method to find the column names automatically.

Parameters

- `image (PIL.Image.Image)` – The RGBA image that has the same shape as the original stratigraphic diagram
- `bounds (np.ndarray of shape (N, 2))` – The boundaries for each column. These are essential for the `find_colnames()` and the `highlight_column()` methods
- `rotate (float)` – An angle between 0 and 90 that corresponds to the rotation of the column names
- `mirror (bool)` – If True, the image is mirrored (horizontally)
- `flip (bool)` – If True, the image is flipped (vertically)

- **highres_image** (*PIL.Image.Image*) – A high resolution version of the *image* with the same width-to-height ratio
- **data_ylim** (*tuple (y0, y1)*) – The vertical data limits of the data part that should be ignored in the *find_colnames()* method if the *ignore_data_part* is True

Methods

<code>close()</code>	Close the column names reader
<code>create_variable(ds, vname, data, **kwargs)</code>	Insert the data into a variable in an <code>xr.Dataset</code>
<code>find_colnames([extents])</code>	Find the names for the columns using tesserocr
<code>from_dataset(ds)</code>	Create a <code>ColNamesReader</code> for a <code>xarray.Dataset</code>
<code>get_colpic(x0, y0, x1, y1)</code>	Extract the picture of the column name
<code>highlight_column(col, ax)</code>	Highlight the column in the given axes displaying the
<code>navigate_to_col(col, ax)</code>	Navigate to the specified column
<code>recognize_text(image)</code>	Recognize the text in an image using tesserocr
<code>rotate_image(image)</code>	Modify an image with <code>rotate</code> , <code>flip</code> , <code>mirror</code>
<code>to_dataset([ds])</code>	All the necessary data as a <code>xarray.Dataset</code>
<code>transform_point(x, y[, invert, image])</code>	Transform a point between un-rotated and rotated co-ordinate system

Attributes

<code>colpics</code>	The pictures of the column names
<code>column_names</code>	The names of the columns
<code>data_ylim</code>	The vertical data limits of the data part that shall be exluded in the
<code>highres_image</code>	The <code>image</code> attribute with higher resolution and with masked out data part if the <code>ignore_data_part</code> attribute is True and the <code>data_ylim</code> attribute is not None.
<code>ignore_data_part</code>	Boolean flag.
<code>image</code>	The RGBA <code>PIL.Image.Image</code> that stores the column names
<code>nc_meta</code>	<code>dict() -> new empty dictionary</code>
<code>rotated_image</code>	The rotated image based on the <code>rotate_image()</code> method

```
close()
    Close the column names reader

property colpics
    The pictures of the column names

property column_names
    The names of the columns

create_variable(ds, vname, data, **kwargs)
    Insert the data into a variable in an xr.Dataset

data_ylim = None
    The vertical data limits of the data part that shall be exluded in the highres_image if the ignore_data_part is True

find_colnames(extents=None)
    Find the names for the columns using tesserocr
```

Parameters `extents` (*list of floats (x0, y0, x1, y1)*) – The extents to crop the `rotated_image`. We only look for column names in this image

Returns

- `dict` – A mapping from column number to a string (the column name)
- `dict` – A mapping from column number to a `PIL.Image.Image` (the image of the column name)
- `dict` – A mapping from column number to a `Box` (the bounding box of the corresponding column name)

classmethod `from_dataset(ds)`

Create a `ColNamesReader` for a `xarray.Dataset`

Parameters `ds` (`xarray.Dataset`) – The dataset as obtained from the `to_dataset()` method

get_colpic(x0, y0, x1, y1)

Extract the picture of the column name

Parameters

- `x0` (`int`) – The left edge
- `y0` (`int`) – The upper edge
- `x1` (`int`) – The right edge
- `y1` (`int`) – The lower edge

Returns The part of the rotated `highres_image` cropped out from the given parameters

Return type `PIL.Image.Image`

highlight_column(col, ax)

Highlight the column in the given axes displaying the `rotated_image`

This method draws a rotated rectangle highlighting the given column `col` in the given `ax`.

Parameters

- `col` (`int`) – The column number
- `ax` (`matplotlib.axes.Axes`) – The matplotlib axes on which to plot the rectangle.
This `ax` is expected to show the `rotated_image`

property highres_image

The `image` attribute with higher resolution and with masked out data part if the `ignore_data_part` attribute is True and the `data_ylim` attribute is not None. The data part is then set to white with 0 alpha

ignore_data_part = True

Boolean flag. If True, the data part is masked out in the `highres_image`

image = None

The RGBA `PIL.Image.Image` that stores the column names

navigate_to_col(col, ax)

Navigate to the specified column

Change the x- and y-limits of the `ax` to display the given `col` based on the `column_bounds`

Parameters

- `col` (`int`) – The column number

- **ax** (`matplotlib.axes.Axes`) – The matplotlib axes for which to update the limits.
This `ax` is expected to show the `rotated_image`

nc_meta = {'colname': { 'dims': 'column', 'long_name': 'Name of the columns'}, 'colname'}

recognize_text (`image`)

Recognize the text in an image using tesserocr

This method uses the `tesserocr.image_to_text()` to read in the text in a given `image`

Parameters `image` (`PIL.Image.Image`) – The image to read in

Returns The text found in it without newline characters

Return type `str`

rotate_image (`image`)

Modify an image with rotate, flip, mirror

This method rotated, mirrors and/or flips the given `image` based on the `rotate`, `mirror` and `flip` attributes

Parameters `image` (`PIL.Image.Image`) – The source image

Returns The target image

Return type `PIL.Image.Image`

property rotated_image

The rotated `image` based on the `rotate_image()` method

to_dataset (`ds=None`)

All the necessary data as a `xarray.Dataset`

Parameters `ds` (`xarray.Dataset`) – The dataset in which to insert the data. If None, a new one will be created

Returns Either the given `ds` or a new `xarray.Dataset` instance

Return type `xarray.Dataset`

transform_point (`x, y, invert=False, image=None`)

Transform a point between un-rotated and rotated coordinate system

Parameters

- **x** (`float`) – The x-coordinate of the point in the source coordinate system
- **y** (`float`) – The y-coordinate of the point in the source coordinate system
- **invert** (`bool`) – If True, the source coordinate system is the rotated one (i.e. this method transform from the `rotated_image` to the coordinate system of the `image`), other wise from the `image` to the `rotated_image`
- **image** (`PIL.Image.Image`) – The unrotated source image. If None, the `image` is used. This image defines the source coordinate system (or the target coordinate system if `invert` is True)

Returns

- `float` – The transformed x-coordinate
- `float` – The transformed y-coordinate

straditize.common module

Module of commonly use python objects

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Functions

<code>rgba2rgb(image[, color])</code>	Alpha composite an RGBA Image with a specified color.
---------------------------------------	---

`straditize.common.rgba2rgb (image, color=(255, 255, 255))`

Alpha composite an RGBA Image with a specified color.

Source: <http://stackoverflow.com/a/9459208/284318>

Parameters

- `image (PIL.Image)` – The PIL RGBA Image object
- `color (tuple)` – The rgb color for the background

Returns The rgb image

Return type PIL.Image

straditize.cross_mark module

Module for a cross mark to select one point in a matplotlib axes

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Classes

<code>CrossMarkText(*args, **kwargs)</code>	A CrossMarks that opens a QInputDialog after changing the position
<code>CrossMarks([pos, ax, selectable, draggable, ...])</code>	A set of draggable marks in a matplotlib axes
<code>DraggableHLine(y[, ax])</code>	A draggable horizontal line
<code>DraggableHLineText(*args, **kwargs)</code>	A CrossMarks that opens a QInputDialog after changing the position
<code>DraggableVLine(x[, ax])</code>	A draggable vertical line
<code>DraggableVLineText(*args, **kwargs)</code>	A CrossMarks that opens a QInputDialog after changing the position

`class straditize.cross_mark.CrossMarkText (*args, **kwargs)`

Bases: `straditize.cross_mark.CrossMarks`

A CrossMarks that opens a QInputDialog after changing the position

Parameters

- `pos` (*tuple of 2 arrays*) – The initial positions of the crosses. The first item marks the x-coordinates of the points, the second the y-coordinates
- `ax` (*matplotlib.axes.Axes*) – The axes object to draw to. If not specified and `draw_lines` is True, the current axes object is used
- `selectable` (*list of { 'x', 'y' }*) – Determine whether only the x-, y-, or both lines should be selectable
- `draggable` (*list of { 'x', 'y' }*) – Determine whether only the x-, y-, or both lines should be draggable
- `idx_h` (*pandas.Index*) – The index for the horizontal coordinates. If not provided, we use a continuous movement along x.
- `idx_v` (*pandas.Index*) – The index for the vertical coordinates. If not provided, we use a continuous movement along y.
- `xlim` (*tuple of floats (xmin, xmax)*) – The minimum and maximum x value for the lines
- `ylim` (*tuple for floats (ymin, ymax)*) – The minimum and maximum y value for the lines
- `select_props` (*color*) – The line properties for selected marks
- `auto_hide` (*bool*) – If True, the lines are hidden if they are not selected.
- `connected_artists` (*list of artists*) – List of artists whose properties should be changed to `select_props` when this marks is selected
- `lock` (*bool*) – If True, at most one mark can be selected at a time
- `draw_lines` (*bool*) – If True, the cross mark lines are drawn. Otherwise, you must call the `draw_lines` method explicitly
- `hide_vertical` (*bool*) – Boolean to control whether the vertical lines should be hidden. If None, the default class attribute is used
- `hide_horizontal` (*bool*) – Boolean to control whether the horizontal lines should be hidden. If None, the default class attribute is used
- `**kwargs` – Any other keyword argument that is passed to the `matplotlib.pyplot.plot()` function

- **dtype** (*object*) – The data type for the data conversion
- **message** (*str*) – The message to display in the dialog
- **label** (*str*) – The label to how this value should be named
- **value** (*float*) – The initial value to use

Methods

<code>ask_for_value([val, label])</code>	Ask for a value for the cross mark
<code>on_release(event, *args, **kwargs)</code>	Release the mark and unselect it

Attributes

<code>value</code>	The value of this cross mark
--------------------	------------------------------

ask_for_value (*val=None, label=None*)

Ask for a value for the cross mark

This method opens a QInputDialog to ask for a new *value***Parameters**

- **val** (*float*) – The initial value
- **label** (*str*) – the name of what to ask for

on_release (*event, *args, **kwargs*)

Release the mark and unselect it

Parameters

- **event** (*matplotlib.backend_bases.MouseEvent*) – The mouseevent that releases the mark
- **force** (*bool*) – If True, the mark is released although it does not contain the *event*
- **connected** (*bool*) – If True, connected marks that should maintain a constant x- and y-distance are released, too
- **draw** (*bool*) – If True, the figure is drawn
- ****kwargs** (**args,*) – Any other parameter that is passed to the connected lines

value = None

The value of this cross mark

```
class straditize.cross_mark.CrossMarks (pos=(0, 0), ax=None, selectable=['h', 'v'], dragable=['h', 'v'], idx_h=None, idx_v=None, xlim=None, ylim=None, select_props={'c': 'r'}, auto_hide=False, connected_artists=[], lock=True, draw_lines=True, hide_vertical=None, hide_horizontal=None, **kwargs)
```

Bases: *object*

A set of draggable marks in a matplotlib axes

Parameters

- **pos** (*tuple of 2 arrays*) – The initial positions of the crosses. The first item marks the x-coordinates of the points, the second the y-coordinates

- **ax** (`matplotlib.axes.Axes`) – The axes object to draw to. If not specified and `draw_lines` is True, the current axes object is used
- **selectable** (`list of {'x', 'y'}`) – Determine whether only the x-, y-, or both lines should be selectable
- **draggable** (`list of {'x', 'y'}`) – Determine whether only the x-, y-, or both lines should be draggable
- **idx_h** (`pandas.Index`) – The index for the horizontal coordinates. If not provided, we use a continuous movement along x.
- **idx_v** (`pandas.Index`) – The index for the vertical coordinates. If not provided, we use a continuous movement along y.
- **xlim** (`tuple of floats (xmin, xmax)`) – The minimum and maximum x value for the lines
- **ylim** (`tuple for floats (ymin, ymax)`) – The minimum and maximum y value for the lines
- **select_props** (`color`) – The line properties for selected marks
- **auto_hide** (`bool`) – If True, the lines are hidden if they are not selected.
- **connected_artists** (`list of artists`) – List of artists whose properties should be changed to `select_props` when this marks is selected
- **lock** (`bool`) – If True, at most one mark can be selected at a time
- **draw_lines** (`bool`) – If True, the cross mark lines are drawn. Otherwise, you must call the `draw_lines` method explicitly
- **hide_vertical** (`bool`) – Boolean to control whether the vertical lines should be hidden. If None, the default class attribute is used
- **hide_horizontal** (`bool`) – Boolean to control whether the horizontal lines should be hidden. If None, the default class attribute is used
- ****kwargs** – Any other keyword argument that is passed to the `matplotlib.pyplot.plot()` function

Attributes

<code>ax</code>	The matplotlib axes to plot on
<code>block_signals</code>	Block the emitting of signals of this instance
<code>connected_artists</code>	a list of <code>matplotlib.artist.Artist</code> whose colors are changed
<code>fig</code>	The <code>matplotlib.figure.Figure</code> that this mark plots on
<code>hide_horizontal</code>	Boolean to control whether the horizontal lines should be hidden
<code>hide_vertical</code>	Boolean to control whether the vertical lines should be hidden
<code>hline</code>	The current horizontal line
<code>hlines</code>	the list of horizontal lines
<code>idx_h</code>	The index for vertical lines
<code>idx_v</code>	The index for horizontal lines
<code>line_connections</code>	The line connections to the current position

Continued on next page

Table 152 – continued from previous page

<code>lock</code>	Class attribute that is set to a CrossMark instance to lock the
<code>moved</code>	A signal that is emitted when the mark is moved.
<code>other_connections</code>	All other connections to the current position
<code>points</code>	The x-y-coordinates of the points as a (N, 2)-shaped array
<code>pos</code>	The position of the current line
<code>show_connected_artists</code>	A boolean to control whether the connected artists should be shown
<code>vline</code>	The current vertical line
<code>vlines</code>	the list of vertical lines
<code>x</code>	The x-position of the mark
<code>xlim</code>	The x-limits of the <code>hlines</code>
<code>y</code>	The y-position of the mark
<code>ylim</code>	The x-limits of the <code>vlines</code>

Methods

<code>connect()</code>	Connect the marks matplotlib events
<code>connect_marks(marks[, visible])</code>	Connect multiple marks to each other
<code>connect_to_marks(marks[, visible, append])</code>	Append other marks that should be considered for aligning the lines
<code>contains(event)</code>	Test if the mark is selected by the given <code>event</code>
<code>disconnect()</code>	Disconnect all the stored connection ids
<code>draw_lines(**kwargs)</code>	Draw the vertical and horizontal lines
<code>is_selected_by(event[, buttons])</code>	Test if the given <code>event</code> selects the mark
<code>maintain_x(marks)</code>	Connect marks and maintain a constant horizontal distance
<code>maintain_y(marks)</code>	Connect marks and maintain a constant vertical distance between them
<code>on_motion(event[, force, move_connected, ...])</code>	Move the lines of this mark
<code>on_press(event[, force, connected])</code>	Select the mark
<code>on_release(event[, force, connected, draw])</code>	Release the mark and unselect it
<code>remove([artists])</code>	Remove all lines and disconnect the mark
<code>set_connected_artists(artists)</code>	Set the connected artists
<code>set_connected_artists_visible(visible)</code>	Set the visibility of the connected artists
<code>set_current_point(x, y[, nearest])</code>	Set the current point that is selected
<code>set_pos(pos)</code>	Move the point(s) to another position
<code>set_visible(b)</code>	Set the visibility of the mark

ax = None

The matplotlib axes to plot on

property block_signals

Block the emitting of signals of this instance

connect()

Connect the marks matplotlib events

static connect_marks (marks, visible=False)

Connect multiple marks to each other

Parameters

- **marks** (*list of CrossMarks*) – A list of marks
- **visible** (*bool*) – If True, the marks are connected through visible lines

Notes

Different from the `connect_to_marks()` method, this static function connects each of the marks to the others.

`connect_to_marks(marks, visible=False, append=True)`

Append other marks that should be considered for aligning the lines

Parameters

- **marks** (*list of CrossMarks*) – A list of other marks
- **visible** (*bool*) – If True, the marks are connected through visible lines
- **append** (*bool*) – If True, the marks are appended. This is important if the mark will be moved by the `set_pos` method

Notes

This method can only be used to connect other marks with this mark. If you want to connect multiple marks within each other, use the `connect_marks()` static method

`connected_artists = []`

a list of `matplotlib.artist.Artist` whose colors are changed when this mark is selected

`contains(event)`

Test if the mark is selected by the given *event*

Parameters `event` (`ButtonPressEvent`) – The ButtonPressEvent that has been triggered

`disconnect()`

Disconnect all the stored connection ids

`draw_lines(**kwargs)`

Draw the vertical and horizontal lines

Parameters `**kwargs` – An keyword that is passed to the `matplotlib.pyplot.plot()` function

`property fig`

The `matplotlib.figure.Figure` that this mark plots on

`hide_horizontal = False`

Boolean to control whether the horizontal lines should be hidden

`hide_vertical = False`

Boolean to control whether the vertical lines should be hidden

`property hline`

The current horizontal line

`hlines = []`

the list of horizontal lines

`property idx_h`

The index for vertical lines

property idx_v

The index for horizontal lines

is_selected_by (event, buttons=[1])

Test if the given *event* selects the mark

Parameters

- **event** (`matplotlib.backend_bases.MouseEvent`) – The matplotlib event
- **button** (`list of int`) – Possible buttons to select this mark

Returns True, if it is selected

Return type `bool`

property line_connections

The line connections to the current position

lock = None

Class attribute that is set to a `CrossMark` instance to lock the selection of marks

static maintain_x (marks)

Connect marks and maintain a constant horizontal distance

Parameters **marks** (`list of CrossMarks`) – A list of marks. If one of the marks is moved horizontally, the others are, too

static maintain_y (marks)

Connect marks and maintain a constant vertical distance between them

Parameters **marks** (`list of CrossMarks`) – A list of marks. If one of the marks is moved vertically, the others are, too

moved

A signal that is emitted when the mark is moved. Connected function are expected to accept two arguments.
One tuple with the old position and the `CrossMarks` instance itself

on_motion (event, force=False, move_connected=True, restore=True)

Move the lines of this mark

Parameters

- **event** (`matplotlib.backend_bases.MouseEvent`) – The mouseevent that moves the mark
- **force** (`bool`) – If True, the mark is moved although it does not contain the *event*
- **move_connected** (`bool`) – If True, connected marks that should maintain a constant x- and y-distance are moved, too
- **restore** (`bool`) – If True, the axes background is restored

on_press (event, force=False, connected=True)

Select the mark

Parameters

- **event** (`matplotlib.backend_bases.MouseEvent`) – The mouseevent that selects the mark
- **force** (`bool`) – If True, the mark is selected although it does not contain the *event*
- **connected** (`bool`) – If True, connected marks that should maintain a constant x- and y-distance are selected, too

on_release(*event*, *force=False*, *connected=True*, *draw=True*, **args*, ***kwargs*)

Release the mark and unselect it

Parameters

- **event** (*matplotlib.backend_bases.MouseEvent*) – The mouseevent that releases the mark
- **force** (*bool*) – If True, the mark is released although it does not contain the *event*
- **connected** (*bool*) – If True, connected marks that should maintain a constant x- and y-distance are released, too
- **draw** (*bool*) – If True, the figure is drawn
- ****kwargs** (**args*,) – Any other parameter that is passed to the connected lines

property other_connections

All other connections to the current position

property points

The x-y-coordinates of the points as a (N, 2)-shaped array

property pos

The position of the current line

remove(*artists=True*)

Remove all lines and disconnect the mark

Parameters **artists** (*bool*) – If True, the *connected_artists* list is cleared and the corresponding artists are removed as well

set_connected_artists(*artists*)

Set the connected artists

Parameters **artists** (*matplotlib.artist.Artist*) – The artists (e.g. other lines) that should be connected and highlighted if this mark is selected

set_connected_artists_visible(*visible*)

Set the visibility of the connected artists

Parameters **visible** (*bool*) – True, show the connected artists, else don't

set_current_point(*x*, *y*, *nearest=False*)

Set the current point that is selected

Parameters

- **x** (*int*) – The index of the x-value in the *xa* attribute
- **y** (*int*) – The index of the y-value in the *ya* attribute
- **nearest** (*bool*) – If not None, *x* and *y* are interpreted as x- and y-values and we select the closest one

set_pos(*pos*)

Move the point(s) to another position

Parameters **pos** (*tuple of 2 arrays*) – The positions of the crosses. The first item marks the x-coordinates of the points, the second the y-coordinates

set_visible(*b*)

Set the visibility of the mark

Parameters **b** (*bool*) – If False, hide all horizontal and vertical lines, and the *connected_artists*

```

show_connected_artists = True
    A boolean to control whether the connected artists should be shown at all

property vline
    The current vertical line

vlines = []
    the list of vertical lines

property x
    The x-position of the mark

xlim = None
    The x-limits of the hlines

property y
    The y-position of the mark

ylim = None
    The x-limits of the vlines

class straditize.cross_mark.DraggableHLine(y, ax=None, *args, **kwargs)
Bases: straditize.cross_mark.CrossMarks

A draggable horizontal line

Parameters

- y (float) – The y-position for the horizontal line
- ax (matplotlib.axes.Axes) – The matplotlib axes
- idx_h (pandas.Index) – The index for the horizontal coordinates. If not provided, we use a continuous movement along x.
- idx_v (pandas.Index) – The index for the vertical coordinates. If not provided, we use a continuous movement along y.
- xlim (tuple of floats (xmin, xmax)) – The minimum and maximum x value for the lines
- ylim (tuple for floats (ymin, ymax)) – The minimum and maximum y value for the lines
- select_props (color) – The line properties for selected marks
- auto_hide (bool) – If True, the lines are hidden if they are not selected.
- connected_artists (list of artists) – List of artists whose properties should be changed to select_props when this marks is selected
- lock (bool) – If True, at most one mark can be selected at a time
- draw_lines (bool) – If True, the cross mark lines are drawn. Otherwise, you must call the draw_lines method explicitly
- hide_vertical (bool) – Boolean to control whether the vertical lines should be hidden. If None, the default class attribute is used
- hide_horizontal (bool) – Boolean to control whether the horizontal lines should be hidden. If None, the default class attribute is used
- **kwargs – Any other keyword argument that is passed to the matplotlib.pyplot.plot() function

```

Attributes

<code>hide_vertical</code>	<code>bool(x) -> bool</code>
<code>x</code>	The x-position of the mark

Methods

<code>set_visible(b)</code>	Set the visibility of the mark
-----------------------------	--------------------------------

`hide_vertical = True`

`set_visible(b)`

Set the visibility of the mark

Parameters `b` (`bool`) – If False, hide all horizontal and vertical lines, and the `connected_artists`

`property x`

The x-position of the mark

class straditize.cross_mark.`DraggableHLineText` (*args, **kwargs)
Bases: `straditize.cross_mark.DraggableHLine`

A CrossMarks that opens a QInputDialog after changing the position

Parameters

- `y` (`float`) – The y-position for the horizontal line
- `ax` (`matplotlib.axes.Axes`) – The matplotlib axes
- `idx_h` (`pandas.Index`) – The index for the horizontal coordinates. If not provided, we use a continuous movement along x.
- `idx_v` (`pandas.Index`) – The index for the vertical coordinates. If not provided, we use a continuous movement along y.
- `xlim` (`tuple of floats (xmin, xmax)`) – The minimum and maximum x value for the lines
- `ylim` (`tuple for floats (ymin, ymax)`) – The minimum and maximum y value for the lines
- `select_props` (`color`) – The line properties for selected marks
- `auto_hide` (`bool`) – If True, the lines are hidden if they are not selected.
- `connected_artists` (`list of artists`) – List of artists whose properties should be changed to `select_props` when this marks is selected
- `lock` (`bool`) – If True, at most one mark can be selected at a time
- `draw_lines` (`bool`) – If True, the cross mark lines are drawn. Otherwise, you must call the `draw_lines` method explicitly
- `hide_vertical` (`bool`) – Boolean to control whether the vertical lines should be hidden. If None, the default class attribute is used
- `hide_horizontal` (`bool`) – Boolean to control whether the horizontal lines should be hidden. If None, the default class attribute is used
- `**kwargs` – Any other keyword argument that is passed to the `matplotlib.pyplot.plot()` function

- **dtype** (*object*) – The data type for the data conversion
- **message** (*str*) – The message to display in the dialog
- **label** (*str*) – The label to how this value should be named
- **value** (*float*) – The initial value to use

Methods

<code>ask_for_value([val, label])</code>	Ask for a value for the cross mark
<code>on_release(event, *args, **kwargs)</code>	Release the mark and unselect it

ask_for_value (*val=None, label=None*)

Ask for a value for the cross mark

This method opens a QInputDialog to ask for a new value

Parameters

- **val** (*float*) – The initial value
- **label** (*str*) – the name of what to ask for

on_release (*event, *args, **kwargs*)

Release the mark and unselect it

Parameters

- **event** (*matplotlib.backend_bases.MouseEvent*) – The mouseevent that releases the mark
- **force** (*bool*) – If True, the mark is released although it does not contain the *event*
- **connected** (*bool*) – If True, connected marks that should maintain a constant x- and y-distance are released, too
- **draw** (*bool*) – If True, the figure is drawn
- ****kwargs** (**args,*) – Any other parameter that is passed to the connected lines

class straditize.cross_mark.DraggableVLine (*x, ax=None, *args, **kwargs*)

Bases: straditize.cross_mark.CrossMarks

A dragable vertical line

Parameters

- **x** (*float*) – The x-position for the vertical line
- **ax** (*matplotlib.axes.Axes*) – The matplotlib axes
- **idx_h** (*pandas.Index*) – The index for the horizontal coordinates. If not provided, we use a continuous movement along x.
- **idx_v** (*pandas.Index*) – The index for the vertical coordinates. If not provided, we use a continuous movement along y.
- **xlim** (*tuple of floats (xmin, xmax)*) – The minimum and maximum x value for the lines
- **ylim** (*tuple for floats (ymin, ymax)*) – The minimum and maximum y value for the lines
- **select_props** (*color*) – The line properties for selected marks

- **auto_hide** (`bool`) – If True, the lines are hidden if they are not selected.
- **connected_artists** (*list of artists*) – List of artists whose properties should be changed to `select_props` when this marks is selected
- **lock** (`bool`) – If True, at most one mark can be selected at a time
- **draw_lines** (`bool`) – If True, the cross mark lines are drawn. Otherwise, you must call the `draw_lines` method explicitly
- **hide_vertical** (`bool`) – Boolean to control whether the vertical lines should be hidden. If None, the default class attribute is used
- **hide_horizontal** (`bool`) – Boolean to control whether the horizontal lines should be hidden. If None, the default class attribute is used
- ****kwargs** – Any other keyword argument that is passed to the `matplotlib.pyplot.plot()` function

Attributes

<code>hide_horizontal</code>	<code>bool(x) -> bool</code>
<code>y</code>	The y-position of the mark

Methods

<code>set_visible(b)</code>	Set the visibility of the mark
-----------------------------	--------------------------------

`hide_horizontal = True`

`set_visible(b)`

Set the visibility of the mark

Parameters `b` (`bool`) – If False, hide all horizontal and vertical lines, and the `connected_artists`

`property y`

The y-position of the mark

`class straditize.cross_mark.DraggableVLineText(*args, **kwargs)`

Bases: `straditize.cross_mark.DraggableVLine`

A CrossMarks that opens a QDialog after changing the position

Parameters

- **x** (`float`) – The x-position for the vertical line
- **ax** (`matplotlib.axes.Axes`) – The matplotlib axes
- **idx_h** (`pandas.Index`) – The index for the horizontal coordinates. If not provided, we use a continuous movement along x.
- **idx_v** (`pandas.Index`) – The index for the vertical coordinates. If not provided, we use a continuous movement along y.
- **xlim** (*tuple of floats (xmin, xmax)*) – The minimum and maximum x value for the lines
- **ylim** (*tuple for floats (ymin, ymax)*) – The minimum and maximum y value for the lines
- **select_props** (`color`) – The line properties for selected marks

- **auto_hide** (`bool`) – If True, the lines are hidden if they are not selected.
- **connected_artists** (`list of artists`) – List of artists whose properties should be changed to `select_props` when this marks is selected
- **lock** (`bool`) – If True, at most one mark can be selected at a time
- **draw_lines** (`bool`) – If True, the cross mark lines are drawn. Otherwise, you must call the `draw_lines` method explicitly
- **hide_vertical** (`bool`) – Boolean to control whether the vertical lines should be hidden. If None, the default class attribute is used
- **hide_horizontal** (`bool`) – Boolean to control whether the horizontal lines should be hidden. If None, the default class attribute is used
- ****kwargs** – Any other keyword argument that is passed to the `matplotlib.pyplot.plot()` function
- **dtype** (`object`) – The data type for the data conversion
- **message** (`str`) – The message to display in the dialog
- **label** (`str`) – The label to how this value should be named
- **value** (`float`) – The initial value to use

Methods

<code>ask_for_value([val, label])</code>	Ask for a value for the cross mark
<code>on_release(event, *args, **kwargs)</code>	Release the mark and unselect it

`ask_for_value(val=None, label=None)`

Ask for a value for the cross mark

This method opens a QInputDialog to ask for a new value

Parameters

- **val** (`float`) – The initial value
- **label** (`str`) – the name of what to ask for

`on_release(event, *args, **kwargs)`

Release the mark and unselect it

Parameters

- **event** (`matplotlib.backend_bases.MouseEvent`) – The mouseevent that releases the mark
- **force** (`bool`) – If True, the mark is released although it does not contain the `event`
- **connected** (`bool`) – If True, connected marks that should maintain a constant x- and y-distance are released, too
- **draw** (`bool`) – If True, the figure is drawn
- ****kwargs** (`*args,`) – Any other parameter that is passed to the connected lines

straditiz.evaluator module

Evaluator class for the straditiz algorithms

Classes

<code>BaselineScenario([output_dir])</code>	The baseline evaluation scenario for straditiz with data from POLNET
<code>BlackWhiteScenario([output_dir])</code>	An evaluation scenario with a binary (black and white) image
<code>DPI150Scenario([output_dir])</code>	Another evaluation scenario but with a resolution of 150 dpi
<code>DPI600Scenario([output_dir])</code>	Another evaluation scenario but with a resolution of 600 dpi
<code>ExaggerationsEvaluator(*args, **kwargs)</code>	An evaluator with exaggerations
<code>ExaggerationsScenario([output_dir])</code>	An evaluation scenario with an exaggerated plot of low percentages
<code>NoVerticalsEvaluator(data, *args[, name, ...])</code>	An evaluator for an image without y-axis
<code>NoVerticalsScenario([output_dir])</code>	An evaluation scenario without y-axes in the plot
<code>StraditizEvaluator(data, *args[, name, ...])</code>	An evaluator for the straditiz components

Functions

<code>print_progressbar(iteration, total[, ...])</code>	Print iterations progress
<code>rmse(sim, ref)</code>	Calculate the root mean squared error between simulation and reference

`class straditiz.evaluator.BaselineScenario (output_dir='.)`

Bases: `object`

The baseline evaluation scenario for straditiz with data from POLNET

This class uses the default settings of the `StraditizEvaluator` and runs the analysis for a given dataset from POLNET. **Methods**

<code>export_evaluator(evaluator, *args, **kwargs)</code>	
<code>init_evaluator(name, data, *args, **kwargs)</code>	Initialize an evaluator for a given data set
<code>run(data[, processes])</code>	

Attributes

<code>index_names</code>	Built-in mutable sequence.
--------------------------	----------------------------

```
export_evaluator (evaluator, *args, **kwargs)
index_names = ['e_', 'ntaxa', 'nsamples']
init_evaluator (name, data, *args, **kwargs)
    Initialize an evaluator for a given data set
run (data, processes=None)
```

`class straditiz.evaluator.BlackWhiteScenario (output_dir='.)`

Bases: `straditiz.evaluator.BaselineScenario`

An evaluation scenario with a binary (black and white) image **Methods**

<code>init_evaluator(*args, **kwargs)</code>	Initialize an evaluator for a given data set
--	--

`init_evaluator(*args, **kwargs)`
Initialize an evaluator for a given data set

`class straditize.evaluator.DPI150Scenario(output_dir='')`

Bases: `straditize.evaluator.BaselineScenario`

Another evaluation scenario but with a resolution of 150 dpi **Methods**

<code>export_evaluator(*args, **kwargs)</code>
--

`export_evaluator(*args, **kwargs)`

`class straditize.evaluator.DPI600Scenario(output_dir='')`

Bases: `straditize.evaluator.BaselineScenario`

Another evaluation scenario but with a resolution of 600 dpi **Methods**

<code>export_evaluator(*args, **kwargs)</code>
--

`export_evaluator(*args, **kwargs)`

`class straditize.evaluator.ExaggerationsEvaluator(*args, **kwargs)`

Bases: `straditize.evaluator.StraditizeEvaluator`

An evaluator with exaggerations

Parameters

- `df (pandas.DataFrame)` – The dataframe containing the data to plot.
- `group_func (function)` – A function that groups the columns in the input `df` together. It must accept the name of a column and return the corresponding group name:

```
def group_func(col_name: str):
    return "name of it's group"
```

If this parameter is not specified, each column will be assigned to the ‘`nogroup`’ group that can then be used in the other parameters, such as `formatoptions` and `percentages`. Each group may also be divided into `subgroups` (see below), in this case, the `group_func` should return the corresponding subgroup.

- `formatoptions (dict)` – The formatoption for each group. Depending on the chosen plot method, this contains the formatoptions for the psyplot plotter.
- `ax (matplotlib.axes.Axes)` – The matplotlib axes to plot on. New axes will be created that cover all the space of the given axes. If this parameter is not specified and `fig` is None, a new matplotlib figure is created with a new matplotlib axes.
- `thresh (float)` – A minimum number between 0 and 100 (by default 1%) that a `percentages` column has to fullfil in order to be included in the plot. If a variable is always below this threshold, it will not be included in the figure
- `percentages (list of str or bool)` – The group names (see `group_func`) that represent percentage values. This variables will be visualized using an area plot and can be rescaled to sum up to 100% using the `calculate_percentages` parameter. This parameter can also be set to True if all groups shall be considered as percentage data

- **exclude** (*list of str*) – Either group names of column names in *df* that should be excluded in the plot
- **widths** (*dict*) – A mapping from group name to its relative width in the plot. The values of this mapping should sum up to 1, e.g.:

```
widths = {'group1': 0.3, 'group2': 0.5, 'group3': 0.2}
```

- **calculate_percentages** (*bool or list of str*) – If True, rescale the groups mentioned in the *percentages* parameter to sum up to 100%. In case of a list of str, this parameter represents the group (or variable) names that shall be used for the normalization
- **min_percentage** (*float*) – The minimum percentage (between 0 and 100) that should be covered by variables displaying *percentages* data. Each plot in one of the *percentages* groups will have at least have a xlim from 0 to *min_percentage*
- **trunc_height** (*float*) – A float between 0 and 1. The fraction of the *ax* that should be reserved for the group titles.
- **fig** (*matplotlib.Figure*) – The matplotlib figure to draw the plot on. If neither *ax* nor *fig* is specified, a new figure will be created.
- **all_in_one** (*list of str*) – The groups mentioned in this parameter will all be plotted in one single axes whereas the default is to plot each variable in a separate plot
- **stacked** (*list of str*) – The groups mentioned in this parameter will all be plotted in one single axes, stacked onto each other
- **summed** (*list of str*) – The groups (or subgroups) mentioned in this parameter will be summed and an extra plot will be appended to the right of the stratigraphic diagram
- **use_bars** (*list of str or bool*) – The variables specified in this parameter (or all variables if *use_bars* is True) will be visualized by a bar diagram, instead of a line or area plot.
- **subgroups** (*dict*) – A mapping from group name to a list of subgroups, e.g.:

```
subgroups = {'Pollen': ['Trees', 'Shrubs']}
```

to divide an overarching group into subgroups.

Methods

`init_stradi(*args, **kwargs)`

`init_stradi(*args, **kwargs)`

class straditize.evaluator.**ExaggerationsScenario** (*output_dir=’.’*)
Bases: straditize.evaluator.*BaselineScenario*

An evaluation scenario with an exaggerated plot of low percentages **Methods**

`init_evaluator(name, data, *args, **kwargs)` Initialize an evaluator for a given data set

`init_evaluator(name, data, *args, **kwargs)`

Initialize an evaluator for a given data set

class straditize.evaluator.**NoVerticalsEvaluator** (*data, *args, name=’data’, axis linestyle={‘bottom’: ‘-’, ‘left’: ‘-’, ‘right’: ‘-’, ‘top’: ‘-’}, **kwargs*)

Bases: `straditize.evaluator.StraditizeEvaluator`

An evaluator for an image without y-axis

Parameters

- **`df`** (`pandas.DataFrame`) – The dataframe containing the data to plot.
- **`group_func`** (`function`) – A function that groups the columns in the input `df` together. It must accept the name of a column and return the corresponding group name:

```
def group_func(col_name: str):
    return "name of it's group"
```

If this parameter is not specified, each column will be assigned to the ‘`nogroup`’ group that can then be used in the other parameters, such as `formatoptions` and `percentages`. Each group may also be divided into `subgroups` (see below), in this case, the `group_func` should return the corresponding subgroup.

- **`formatoptions`** (`dict`) – The `formatoption` for each group. Depending on the chosen plot method, this contains the `formatoptions` for the `psyplot` plotter.
- **`ax`** (`matplotlib.axes.Axes`) – The `matplotlib` axes to plot on. New axes will be created that cover all the space of the given axes. If this parameter is not specified and `fig` is `None`, a new `matplotlib` figure is created with a new `matplotlib` axes.
- **`thresh`** (`float`) – A minimum number between 0 and 100 (by default 1%) that a `percentages` column has to fullfil in order to be included in the plot. If a variable is always below this threshold, it will not be included in the figure
- **`percentages`** (`list of str or bool`) – The group names (see `group_func`) that represent percentage values. This variables will be visualized using an area plot and can be rescaled to sum up to 100% using the `calculate_percentages` parameter. This parameter can also be set to `True` if all groups shall be considered as percentage data
- **`exclude`** (`list of str`) – Either group names of column names in `df` that should be excluded in the plot
- **`widths`** (`dict`) – A mapping from group name to it’s relative width in the plot. The values of this mapping should some up to 1, e.g.:

```
widths = {'group1': 0.3, 'group2': 0.5, 'group3': 0.2}
```

- **`calculate_percentages`** (`bool or list of str`) – If `True`, rescale the groups mentioned in the `percentages` parameter to sum up to 100%. In case of a list of str, this parameter represents the group (or variable) names that shall be used for the normalization
- **`min_percentage`** (`float`) – The minimum percentage (between 0 and 100) that should be covered by variables displaying `percentages` data. Each plot in one of the `percentages` groups will have at least have a `xlim` from 0 to `min_percentage`
- **`trunc_height`** (`float`) – A float between 0 and 1. The fraction of the `ax` that should be reserved for the group titles.
- **`fig`** (`matplotlib.Figure`) – The `matplotlib` figure to draw the plot on. If neither `ax` nor `fig` is specified, a new figure will be created.
- **`all_in_one`** (`list of str`) – The groups mentioned in this parameter will all be plotted in one single axes whereas the default is to plot each variable in a separate plot
- **`stacked`** (`list of str`) – The groups mentioned in this parameter will all be plotted in one single axes, stacked onto each other

- **summed** (*list of str*) – The groups (or subgroups) mentioned in this parameter will be summed and an extra plot will be appended to the right of the stratigraphic diagram
- **use_bars** (*list of str or bool*) – The variables specified in this parameter (or all variables if *use_bars* is True) will be visualized by a bar diagram, instead of a line or area plot.
- **subgroups** (*dict*) – A mapping from group name to a list of subgroups, e.g.:

```
subgroups = {'Pollen': ['Trees', 'Shrubs']}
```

to divide an overarching group into subgroups.

Methods

```
evaluate_column_starts([close, base])
evaluate_yaxes_removal([close])
export(*args, **kwargs)
```

```
evaluate_column_starts(close=True, base='starts_')
```

```
evaluate_yaxes_removal(close=True)
```

```
export(*args, **kwargs)
```

```
class straditize.evaluator.NoVerticalsScenario(output_dir='.'
```

Bases: *straditize.evaluator.BaselineScenario*

An evaluation scenario without y-axes in the plot

```
init_evaluator(name, data, *args, **kwargs) Initialize an evaluator for a given data set
```

```
init_evaluator(name, data, *args, **kwargs)
```

Initialize an evaluator for a given data set

```
class straditize.evaluator.StraditizeEvaluator(data, *args, name='data', axis-
linestyle={'bottom': '-', 'left': '-', 'right': '-', 'top': '-'}, **kwargs)
```

Bases: *object*

An evaluator for the straditize components

Parameters

- **df** (*pandas.DataFrame*) – The dataframe containing the data to plot.
- **group_func** (*function*) – A function that groups the columns in the input *df* together. It must accept the name of a column and return the corresponding group name:

```
def group_func(col_name: str):
    return "name of it's group"
```

If this parameter is not specified, each column will be assigned to the ‘nogroup’ group that can then be used in the other parameters, such as *formatoptions* and *percentages*. Each group may also be divided into *subgroups* (see below), in this case, the *group_func* should return the corresponding subgroup.

- **formatoptions** (*dict*) – The formatoption for each group. Depending on the chosen plot method, this contains the formatoptions for the psyplot plotter.

- **ax** (`matplotlib.axes.Axes`) – The matplotlib axes to plot on. New axes will be created that cover all the space of the given axes. If this parameter is not specified and `fig` is `None`, a new matplotlib figure is created with a new matplotlib axes.
- **thresh** (`float`) – A minimum number between 0 and 100 (by default 1%) that a *percentages* column has to fullfil in order to be included in the plot. If a variable is always below this threshold, it will not be included in the figure
- **percentages** (`list of str or bool`) – The group names (see `group_func`) that represent percentage values. This variables will be visualized using an area plot and can be rescaled to sum up to 100% using the `calculate_percentages` parameter. This parameter can also be set to True if all groups shall be considered as percentage data
- **exclude** (`list of str`) – Either group names or column names in `df` that should be excluded in the plot
- **widths** (`dict`) – A mapping from group name to it's relative width in the plot. The values of this mapping should some up to 1, e.g.:

```
widths = {'group1': 0.3, 'group2': 0.5, 'group3': 0.2}
```

- **calculate_percentages** (`bool or list of str`) – If True, rescale the groups mentioned in the `percentages` parameter to sum up to 100%. In case of a list of str, this parameter represents the group (or variable) names that shall be used for the normalization
- **min_percentage** (`float`) – The minimum percentage (between 0 and 100) that should be covered by variables displaying *percentages* data. Each plot in one of the `percentages` groups will have at least have a xlim from 0 to `min_percentage`
- **trunc_height** (`float`) – A float between 0 and 1. The fraction of the `ax` that should be reserved for the group titles.
- **fig** (`matplotlib.Figure`) – The matplotlib figure to draw the plot on. If neither `ax` nor `fig` is specified, a new figure will be created.
- **all_in_one** (`list of str`) – The groups mentioned in this parameter will all be plotted in one single axes whereas the default is to plot each variable in a separate plot
- **stacked** (`list of str`) – The groups mentioned in this parameter will all be plotted in one single axes, stacked onto each other
- **summed** (`list of str`) – The groups (or subgroups) mentioned in this parameter will be summed and an extra plot will be appended to the right of the stratigraphic diagram
- **use_bars** (`list of str or bool`) – The variables specified in this parameter (or all variables if `use_bars` is True) will be visualized by a bar diagram, instead of a line or area plot.
- **subgroups** (`dict`) – A mapping from group name to a list of subgroups, e.g.:

```
subgroups = {'Pollen': ['Trees', 'Shrubs']}
```

to divide an overarching group into subgroups.

Attributes

Continued on next page

Table 171 – continued from previous page

<code>data</code>	
<code>data_xlim</code>	
<code>data_ylim</code>	
<code>dpi</code>	
<code>full_df</code>	
<code>height</code>	
<code>results</code>	
<code>results_column</code>	The column name in <code>all_results</code>
<code>summed_perc</code>	
<code>transformed_data</code>	The <code>data</code> in pixel coordinates
<code>width</code>	

Methods

<code>close()</code>	
<code>evaluate_column_starts([close, base])</code>	
<code>evaluate_full([close])</code>	
<code>evaluate_sample_accuracy([close, stradi, base])</code>	
<code>evaluate_sample_position([close, stradi, base])</code>	
<code>evaluate_yaxes_removal([close])</code>	
<code>export(filepath[, dpi, labels])</code>	
<code>from_polnet(data, *args, **kwargs)</code>	
<code>init_stradi([datalim, columns, names, ...])</code>	
<code>run()</code>	Run all evaluations
<code>set_xtranslation(stradi)</code>	

```
property all_results
close()
property column_bounds
property column_ends
property column_starts
property data
property data_xlim
property data_ylim
property dpi
evaluate_column_starts(close=True, base='starts_')
evaluate_full(close=True)
evaluate_sample_accuracy(close=True, stradi=None, base='samples_')
evaluate_sample_position(close=True, stradi=None, base='samples_')
evaluate_yaxes_removal(close=True)
export(filepath, dpi=300, labels={})
classmethod from_polnet(data, *args, **kwargs)
```

```

property full_df
property height
init_stradi(datalim=True, columns=True, names=True, digitize=True, samples=True,
axes=False)
property results
property results_column
    The column name in all_results
run()
    Run all evaluations
set_xtranslation(stradi)
property summed_perc
property transformed_data
    The data in pixel coordinates
property width
straditize.evaluator.print_progressbar(iteration, total, prefix="", suffix="", length=100,
                                         fill'')
Print iterations progress

```

Taken from <https://stackoverflow.com/a/34325723>

Parameters

- **iteration** (*int*) – current iteration
- **total** (*int*) – total iterations
- **prefix** (*str*) – prefix string
- **suffix** (*str*) – suffix string
- **decimals** (*int*) – positive number of decimals in percent complete
- **length** (*int*) – character length of bar
- **fill** (*str*) – bar fill character

```
straditize.evaluator.rmse(sim, ref)
```

Calculate the root mean squared error between simulation and reference

Parameters

- **sim** (*np.ndarray*) – The simulated data
- **ref** (*np.ndarray*) – The reference data

straditize.label_selection module

Module for the *LabelSelection* class

This module defines the *LabelSelection* class, a base class for the *straditize.straditzer.Straditzer* *straditize.binary.DataReader* classes. This class implements the features to select parts of an image and deletes them. The *straditize.widgets.selection_toolbar.SelectionToolbar* interfaces with instances of this class.

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Classes

<i>LabelSelection</i>	Class to provide selection functionalities for an image
-----------------------	---

class straditize.label_selection.**LabelSelection**

Bases: *object*

Class to provide selection functionalities for an image

This class provides functionalities to select features in an image. A new selection can be started through *enable_label_selection()* method and selected parts can be removed through the *remove_selected_labels()* method.

A 2D boolean mask of the selected pixels can be accessed through the *selected_part* attribute.

This class generally assumes that the array for the selection is a 2D integer array, e.g. obtained from the *skimage.morphology.label()* function.

The selection of labels is handled through the colormap. The selection is displayed as a matplotlib image on the *ax* attribute of this instance. If the color for one label is equal to the *cselect* color, it is considered as selected. Additionally every cell that has a value greater than the original number of labels is considered to be selected.

Cells with a value of -1 are not selected and cells with a value of 0 cannot be selected. **Attributes**

<i>cselect</i>	The RGBA color for selected polygons
<i>cunselect</i>	The RGBA color for unselected polygons
<i>label_arrs</i>	List of attribute names of arrays that should be modified if the labels are about to be removed.
<i>remove_callbacks</i>	Functions that shall be called before the labels are removed.
<i>selected_labeled_part</i>	The selected part as a 2D boolean mask
<i>selected_labels</i>	A list of selected labels in the selection array
<i>selected_part</i>	The selected part as a 2D boolean mask

Methods

<i>copy_cmap(cmap_src, colors)</i>	Copy a colormap with replaced colors
<i>disable_label_selection([remove])</i>	Disable the label selection
<i>enable_label_selection(arr, ncolors[, img, ...])</i>	Start the selection of labels
<i>get_default_cmap(ncolors)</i>	The default colormap for binary images
<i>highlight_small_selections([n])</i>	Highlight labels that cover only a small portion of cells
<i>pick_label(event)</i>	Pick the label selected by the given mouseevent

Continued on next page

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<code>remove_selected_labels([disable])</code>	Remove the selected parts of the diagram
<code>remove_small_selection_ellipses()</code>	Remove the ellipses for small features
<code>select_all_labels()</code>	Select the entire array
<code>select_all_other_labels()</code>	Invert the selection
<code>select_labels(selected)</code>	Select a list of labels
<code>unselect_all_labels()</code>	Clear the selection

cid_select = None

static copy_cmap(cmap_src, colors)
Copy a colormap with replaced colors

This function creates a method that has the same name and the same *under*, *over* and *bad* values as the given *cmap_src* but with replaced colors

Parameters

- **cmap_src** (`matplotlib.colors.Colormap`) – The source colormap
- **colors** (`np.ndarray`) – The colors for the colormap

Returns The new colormap

Return type `matplotlib.colors.Colormap`

cselect = [1.0, 0.0, 0.0, 1.0]

The RGBA color for selected polygons

cunselect = [0.0, 0.0, 0.0, 0.0]

The RGBA color for unselected polygons

disable_label_selection(remove=None)

Disable the label selection

This will disconnect the *pick_event* and remove the selection images

Parameters `remove(bool)` – Whether to remove the selection image from the plot. If None, the `_remove` attribute is used

See also:

`enable_label_selection(), remove_selected_labels()`

enable_label_selection(arr, ncolors, img=None, set_picker=False, **kwargs)

Start the selection of labels

Parameters

- **arr** (2D `np.ndarray` of `dtype int`) – The labeled array that contains the features to select.
- **ncolors** (`int`) – The maximum of the labels in *arr*
- **img** (`matplotlib image`) – The image for the selection. If not provided, a new image is created
- **set_picker** (`bool`) – If True, connect the `matplotlib pick_event` to the `pick_label()` method

See also:

`disable_label_selection(), remove_selected_labels()`

get_default_cmap(*ncolors*)

The default colormap for binary images

highlight_small_selections(*n=20*)

Highlight labels that cover only a small portion of cells

This method uses the `skimage.morphology.remove_small_objects()` to detect and highlight small features in the diagram. Each feature will be highlighted through an ellipsis around it.

See also:

`remove_small_selection_ellipses()`

label_attrs = []

List of attribute names of arrays that should be modified if the labels are about to be removed. The attributes might be callable and should then provide the array

pick_label(*event*)

Pick the label selected by the given mouseevent

remove_callbacks = None

Functions that shall be called before the labels are removed. The keys must be the attributes in the `label_attrs` list, values must be list of function that accept two arguments, the array and the boolean mask highlighting the cells that will be set to 0

remove_selected_labels(*disable=False*)

Remove the selected parts of the diagram

This method will call the callbacks in the `remove_callbacks` attribute for all the attributes in the `label_attrs` list.

Parameters **disable**(*bool*) – If True, call the `disable_label_selection()` method at the end

See also:

`enable_label_selection()`, `disable_label_selection()`

remove_small_selection_ellipses()

Remove the ellipses for small features

Removes the ellipses plotted by the `highlight_small_selections()` method

select_all_labels()

Select the entire array

select_all_other_labels()

Invert the selection

select_labels(*selected*)

Select a list of labels

Parameters **selected**(*np.ndarray*) – The numpy array of labels that should be selected

property selected_labeled_part

The selected part as a 2D boolean mask

property selected_labels

A list of selected labels in the selection array

property selected_part

The selected part as a 2D boolean mask

unselect_all_labels()

Clear the selection

straditize.magnifier module

Magnifier class for an image

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Classes

<code>Magnifier(ax_src[, ax])</code>	A magnification of a matplotlib axes
--------------------------------------	--------------------------------------

class straditize.magnifier.**Magnifier**(*ax_src*, *ax=None*, **args*, ***kwargs*)
Bases: `object`

A magnification of a matplotlib axes

It zooms into the region where the mouse pointer is, when it enters the source axes. The appearance of the plot is defined by the `make_plot()` method. **Methods**

<code>adjust_limits(zoom_val)</code>	
<code>close()</code>	Close the magnifier and the associated plots
<code>disconnect()</code>	
<code>enable_zoom()</code>	
<code>make_plot(image, *args, **kwargs)</code>	
<code>onenter(event)</code>	
<code>onleave(event)</code>	
<code>onmotion(event)</code>	

Attributes

<code>dx</code>
<code>dy</code>

```
adjust_limits(zoom_val)
ax = None
cid_enter = None
cid_leave = None
cid_motion = None
close()
Close the magnifier and the associated plots
disconnect()
```

```
property dx
property dy
enable_zoom()
make_plot(image, *args, **kwargs)
onenter(event)
onleave(event)
onmotion(event)
```

straditize.straditizer module

Core module of the Straditizer class

This module defines the *Straditizer* class, the main object to digitize a stratigraphic diagram

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Classes

<i>Straditizer</i> (image[, ax, plot, attrs])	An object to digitize a stratigraphic diagram
---	---

Functions

<i>format_coord_func</i> (ax, ref)	Create a function that can replace the
------------------------------------	--

class straditize.straditizer.**Straditizer**(image, ax=None, plot=True, attrs=None)

Bases: *straditize.label_selection.LabelSelection*

An object to digitize a stratigraphic diagram

One instance of a straditizer manages the digitization of a *PIL.Image.Image* hold in the *image* attribute.

To create a new Straditizer instance, you can either provide a *PIL.Image.Image* to the class constructor (i.e. *stradi = Straditizer(image)*) or you use the *from_dataset()* method.

The reader for the diagram part can be accessed through the *data_reader* attribute, the reader for the column names through the *colnames_reader* attribute.

Parameters

- **image** (*PIL.Image.Image* or *np.ndarray*) – The image file to process. A numpy array should be 3D with shape (Y, X, 4), where the last channel [..., -1] should represent the alpha channel. A *PIL.Image.Image* will be converted to a RGBA image (if not already)

- **ax** (`matplotlib.axes.Axes`) – The matplotlib axes. If None, a new one will be created
- **attrs** (dict or `pandas.DataFrame`) – The attributes for this straditzer

Methods

<code>adjust_lims()</code>	
<code>adjust_lims_after_resize([event])</code>	
<code>adjust_lims_after_zoom(ax)</code>	
<code>align_columns()</code>	Shift the columns after the marks have been moved
<code>close()</code>	
<code>create_magni_marks(marks)</code>	Copy the created marks to the magnifiers axes
<code>create_variable(ds, vname, data, **kwargs)</code>	Insert the data into a variable in an <code>xr.Dataset</code>
<code>data2px_y(coord)</code>	Transform the data coordinates into pixel coordinates
<code>digitize_diagram()</code>	
<code>draw_data_box()</code>	
<code>draw_figure()</code>	
<code>from_dataset(ds[, ax, plot])</code>	Create a new <code>Straditizer</code> from a dataset
<code>get_attr(key)</code>	
<code>get_labels([categorize])</code>	
<code>get_reader_for_column(col)</code>	
<code>guess_data_lims([fraction])</code>	Guess the limits of the diagram part
<code>image_array()</code>	
<code>init_reader([reader_type, ax])</code>	
<code>load(fname[, ax, plot])</code>	
<code>marks_for_column_ends([threshold])</code>	
<code>marks_for_column_starts([threshold])</code>	
<code>marks_for_data_selection([nums, fraction, ...])</code>	
<code>marks_for_occurrences()</code>	Create marks for editing the occurrences
<code>marks_for_samples()</code>	
<code>marks_for_samples_sep([nrows])</code>	
<code>marks_for_vertical_alignment()</code>	Create marks for vertical alignment of the columns
<code>marks_for_x_values([at_col_start])</code>	Create two marks for selecting the x-values
<code>marks_for_y_values()</code>	Create two marks for selecting the x-values
<code>plot_image([ax])</code>	
<code>px2data_y(coord)</code>	Transform the pixel coordinates into data coordinates
<code>remove_data_box()</code>	Remove the data_box
<code>remove_marks()</code>	Remove any drawn marks
<code>reset_image(image[, reader])</code>	Reset the straditzer image
<code>save(fname)</code>	Dump the <code>Straditizer</code> instance to a file
<code>set_attr(key, value)</code>	Update an attribute in the attrs
<code>show_data_diagram()</code>	
<code>show_full_image()</code>	
<code>to_dataset([ds])</code>	All the necessary data as a <code>xarray.Dataset</code>
<code>update_column_ends()</code>	
<code>update_column_starts()</code>	
<code>update_data_part()</code>	
<code>update_image(arr, mask)</code>	Update the image from the given 3D-array
<code>update_occurrences([remove])</code>	Set the occurrences from the given marks
<code>update_samples([remove])</code>	

Continued on next page

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`update_samples_sep([remove])`

`update_xvalues()`

`update_yvalues()`

Attributes

<code>adjusting</code>	True if xlim and ylim are adjusted
<code>attrs</code>	<code>pandas.DataFrame</code> . The attributes of this straditizer
<code>attrs_dict</code>	
<code>ax</code>	The matplotlib axes
<code>block_signals</code>	Block the emitting of signals of this instance
<code>colnames_reader</code>	The <code>straditize.colnames.ColNamesReader</code> for reading the column
<code>column_indexes</code>	The horizontal indexes for each column
<code>data_reader</code>	The <code>straditize.binary.DataReader</code> instance to digitize the
<code>data_xlim</code>	
<code>data_ylim</code>	
<code>fig</code>	
<code>final_df</code>	
<code>full_df</code>	
<code>indexes</code>	
<code>label_arrs</code>	Built-in mutable sequence.
<code>magni</code>	The <code>straditize.magnifier.Magnifier</code> for the diagram image
<code>mark_added</code>	A signal that is emitted if a mark has been added.
<code>mark_cids</code>	set() -> new empty set object
<code>mark_removed</code>	A signal that is emitted, if a mark has been removed.
<code>nc_meta</code>	dict() -> new empty dictionary
<code>valid_attrs</code>	
<code>yaxis_px</code>	

`adjust_lims()``adjust_lims_after_resize(event=None)``adjust_lims_after_zoom(ax)`

property adjusting

True if xlim and ylim are adjusted

`align_columns()`

Shift the columns after the marks have been moved

This method should be called after the `marks_for_vertical_alignment()` method to align the columns

`attrs = None`

`pandas.DataFrame`. The attributes of this straditzer

`property attrs_dict``ax = None`

The matplotlib axes

```

property block_signals
    Block the emitting of signals of this instance

close()

property colnames_reader
    The straditize.colnames.ColNamesReader for reading the column names

property column_indexes
    The horizontal indexes for each column

create_magni_marks(marks)
    Copy the created marks to the magnifiers axes

create_variable(ds, vname, data, **kwargs)
    Insert the data into a variable in an xr.Dataset

data2px_y(coord)
    Transform the data coordinates into pixel coordinates

        Parameters coord (1D np.ndarray) – The coordinate values

        Returns The numpy array with transformed coordinates

        Return type np.ndarray

data_reader = None
    The straditize.binary.DataReader instance to digitize the data

property data_xlim
property data_ylim
digitize_diagram()
draw_data_box()
draw_figure()
property fig
fig_h = None
fig_w = None
property final_df

classmethod from_dataset(ds, ax=None, plot=True)
    Create a new Straditzer from a dataset

    This method uses a dataset that has been exported with the to_dataset() method to initialize a new reader

property full_df
get_attr(key)
get_labels(categorize=1)
get_reader_for_column(col)
guess_data_lims(fraction=0.7)
    Guess the limits of the diagram part

        Parameters fraction (float) – The smallest fraction that has to be covered for a cell to be considered as a corner

        Returns

```

- `np.array` – xmin and xmax of the diagram part (see `data_xlim`)
- `np.array` – ymin and ymax of the diagram part (see `data_ylim`)

image_array()

property indexes

init_reader(`reader_type='area'`, `ax=None`, `**kwargs`)

label_arrs = ['image_array']

classmethod load(`fname`, `ax=None`, `plot=True`)

magni = None
The `straditize.magnifier.Magnifier` for the diagram image

mark_added
A signal that is emitted if a mark has been added. Functions are expected to accept one argument, the newly created `straditize.cross_mark.CrossMarks` instance

mark_cids = {}

mark_removed
A signal that is emitted, if a mark has been removed. Functions are expected to accept one argument, the removed `straditize.cross_mark.CrossMarks` instance

marks_for_column_ends(`threshold=None`)

marks_for_column_starts(`threshold=None`)

marks_for_data_selection(`nums=2`, `fraction=0.7`, `guess_lims=True`)

marks_for_occurrences()
Create marks for editing the occurrences

marks_for_samples()

marks_for_samples_sep(`nrows=3`)

marks_for_vertical_alignment()
Create marks for vertical alignment of the columns
This method creates one mark for each column. These marks should then be moved to positions that should be on the same vertical level. After that, the `align_columns()` method has to be called

marks_for_x_values(`at_col_start=True`)
Create two marks for selecting the x-values
Parameters `at_col_start` (`bool`) – If True, and no translation has yet been performed, create a mark at the column start and ask for the corresponding value

marks_for_y_values()
Create two marks for selecting the y-values

nc_meta = {'axis': {'long_name': 'Axis coordinate'}, 'data_lims': {'dims': ('axis')}

plot_image(`ax=None`, `**kwargs`)

px2data_y(`coord`)
Transform the pixel coordinates into data coordinates
Parameters `coord` (`1D np.ndarray`) – The coordinate values in pixels
Returns The numpy array with transformed coordinates
Return type `np.ndarray`

remove_data_box()
Remove the data_box

remove_marks()
Remove any drawn marks

reset_image(image, reader=False)
Reset the straditzer image

Parameters

- **image** (*PIL.Image.Image*) – The new image to use
- **reader** (*bool*) – If True, the image of the data reader will be replaced, too

save(fname)
Dump the *Straditizer* instance to a file

Parameters fname (str) – The file name where to save the instance

set_attr(key, value)
Update an attribute in the *attrs*

show_data_diagram()

show_full_image()

to_dataset(ds=None)
All the necessary data as a *xarray.Dataset*

Parameters ds (xarray.Dataset) – The dataset in which to insert the data. If None, a new one will be created

Returns Either the given *ds* or a new *xarray.Dataset* instance

Return type *xarray.Dataset*

update_column_ends()

update_column_starts()

update_data_part()

update_image(arr, mask)
Update the image from the given 3D-array

Parameters

- **arr** (3D *np.ndarray* of *dtype float*) – The image array
- **mask** (boolean mask of the same shape as *arr*) – The mask of features that shall be set to 0 in *arr*

update_occurrences(remove=True)
Set the occurrences from the given marks

update_samples(remove=True)

update_samples_sep(remove=True)

update_xvalues()

update_yvalues()

property valid_attrs

yaxis_data = None

property yaxis_px

straditize.straditizer.**format_coord_func**(ax, ref)

Create a function that can replace the `matplotlib.axes.Axes.format_coord()`

Parameters

- **ax** (`matplotlib.axes.Axes`) – The axes instance
- **ref** (`weakref.weakref`) – The reference to the `Straditizer` instance

Returns The function that can be used to replace `ax.format_coord`

Return type function

straditize.version module

Version string of straditize

Disclaimer

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1.8 Changelog

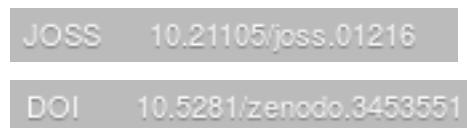
1.8.1 v0.1.3

Patch that forces the diagram limits to be integers.

1.8.2 v0.1.2

This release contains several small bug fixes, mainly for the bar data reader and for duplicated samples and occurrences.

1.8.3 v0.1.1



This release has been approved by the Journal of Open Source Software (JOSS) in <https://github.com/openjournals/joss-reviews/issues/1216>

Added

- Changelog

Changed

- Thanks to the feedbacks from [@ixjlyons](#) and [@sgrieve](#) the installation instructions and some other documentation instructions have been improved (see the issues <https://github.com/Chilipp/straditize/issues/1>, <https://github.com/Chilipp/straditize/issues/2>, <https://github.com/Chilipp/straditize/issues/3>, <https://github.com/Chilipp/straditize/issues/4> and <https://github.com/Chilipp/straditize/issues/8>)

**CHAPTER
TWO**

HOW TO CITE STRADITIZE

When using straditize, you should at least cite the publication in [the Journal of Open Source Software](#):

Sommer, Philipp, Dilan Rech, Manuel Chevalier, and Basil A. S. Davis. Straditize: Digitizing Stratigraphic Diagrams. *Journal of Open Source Software*, vol. 4, no. 34, 34, The Open Journal, Feb. 2019, p. 1216, doi:10.21105/joss.01216, <https://doi.org/10.21105/joss.00363>.

BibTex - EndNote

Furthermore, each release of straditize is associated with a DOI using [zenodo.org](#). If you want to cite a specific version or plugin, please refer to the [releases page of straditize](#).

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THREE

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- genindex
- modindex
- search

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[Basil2007] Davis, Basil A. S., and A. C. Stevenson. “The 8.2ka Event and Early-Mid Holocene Forests, Fires and Flooding in the Central Ebro Desert, NE Spain.” *Quat. Sci. Rev.* , vol. 26, no. 13-14, 2007, pp. 1695-712, doi:10.1016/j.quascirev.2007.04.007.

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