
Mangrove-Surface-Python-SDK Documentation

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Mangrove

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Contents

1	Documentation	3
2	Installation	5
3	(<i>optional</i>) Setup your environment variables	7
4	Test your installation	9
5	Support	11
6	Contents	13
	Python Module Index	37

CHAPTER 1

Documentation

A complete documentation is available there: [documentation](#)

CHAPTER 2

Installation

Install the python SDK package:

```
pip install mangrove-surface
```

or using the git repository:

```
git clone https://github.com/mangroveai/mangrove-surface-python-sdk  
python setup.py install
```


CHAPTER 3

(optional) Setup your environment variables

You can use the SDK with an explicit configuration of url instance and token or you can provide them as environment variables: `SURFACE_URL` and `SURFACE_TOKEN`.

For example on unix-like system:

```
$ export SURFACE_URL=http://your_mangrove.ai_url/api
$ export SURFACE_TOKEN='eyJ0eXAiOiJKV1QiLCJhbGciOiJ...'
```

Warning: Mangrove Surface URL have to end with `/api`.

Note: The explicit configuration overrides the implicit one.

Note: Token can be provided by the administrator using GUI (see documentation) or using the SDK (see `mangrove.Surface._Admin.create_token()`).

CHAPTER 4

Test your installation

You can run tests with the following python lines:

- Test if it is properly installed:

```
>>> import mangrove_surface
>>> mangrove_surface.__version__
'2.0.0'
```

The python SDK is properly connected to your Mangrove Surface:

```
>>> from mangrove_surface import SurfaceClient
>>> # if environment variables are setup
>>> client = SurfaceClient()
>>> # otherwise
>>> # client = SurfaceClient(url="...", token="...")
>>> client.admin.versions()
[
  {
    'name': 'atlas',
    'version': '1.0.0'
  }, {
    'name': 'license_authority',
    'version': u'1.5.0'
  }, {
    'name': 'dmgr',
    'version': '1.0.0'
  }, {
    'name': 'modeler',
    'version': '1.0.0'
  }, {
    'name': 'exporter',
    'version': '1.0.0'
  }, {
    'name': 'mangrove-surface-sdk',
    'version': '1.0.0'
  }
]
```

```
}  
]
```

It is well configured! Congratulation!

Let's begin with *Surface*.

CHAPTER 5

Support

Please refer to *Logger* section if you have any unexpected behavior using the SDK.

6.1 Surface

class mangrove_surface.**SurfaceClient** (*url=None, token=None, username=None, password=None*)

Instantiate Mangrove.ai python SDK with instance url and identity access

Parameters

- **url** – (*optional*) url of the Mangrove.ai instance (by default environment variable SURFACE_URL is used)
- **token** – (*optional*) access token used to secure connection (by default, if username/password are given then those are used to generate an access token; otherwise environment variable SURFACE_TOKEN is used)
- **username** – (*optional*) username used (with password) to sign in (by default, a token is expected)
- **password** – (*optional*) password used to (with username) sign in (by default, a token is expected)

Note: Surface URL or access token can be explicitly provided as parameters or implicitly using environment variables (SURFACE_URL and SURFACE_TOKEN)

Raises

- **IOError** – if the endpoint doesn't answer correctly
- **AttributeError** – if url, username-password or token are not provided

Load Surface python SDK:

```
>>> from mangrove_surface import SurfaceClient
```

Instantiate with url and token are environment variables:

```
>>> client = SurfaceClient()
```

Or with url as environment variable and an explicit token:

```
>>> client = SurfaceClient(token='eyJ0eXAiOiJKV1QiLCJhbGciOiJ...')
```

Or with explicit url and token:

```
>>> client = SurfaceClient(  
...     url='http://my.surface/api',  
...     token='eyJ0eXAiOiJKV1QiLCJhbGciOiJ...'  
... )
```

create_project (*name*, *schema*, *description*=", *schema_test*=None, *tags*=[], *default_classifier*=True, *force*=False)

Create a new project

Parameters

- **name** – project name
- **description** – (*optional*) project description
- **schema** – (*optional*) a data schema which contains train data sets and relations, like:

```
{  
  "tags": ["dataset", "tag"],  
  "datasets": [  
    {  
      "name": "Dataset Name",  
      "filepath": "/path/to/dataset.csv",  
      "tags": ["optional", "tags"],  
      "central": True | False,  
      "keys": ["index"], # optional if there is only  
                        # one dataset  
      "separator": ",", # could be `|`, ` `, `;` or ` `   
    }, ...  
  ],  
  "outcome": "FIELD TARGETED",  
  "outcome_modality": "main value targeted"  
}
```

filepath is an absolute filepath or it could be a S3 uri, like:

```
{  
  "type": "s3",  
  "bucket": "mang-model-producer-samples",  
  "key": "CAR_INSURANCE/CHAT_SESSION_CONTENT_TRAIN.csv"  
}
```

- **tags** – (*optional*) list of project tags
- **schema_test** – (*optional*) a data schema which contains test data sets and relations, like **schema**
- **default_classifier** – (*default*: True) indicates if a default classifier it provided at the project creation


```
>>> mang.admin.license_information()
[{'expires_at': '2018-08-29 00:00:00 UTC',
  'service_level': 'full',
  'system_information': None,
  'updated_at': '2018-03-29 09:49:34 UTC'}]

>>> mang.admin.license_information(request_code=True)
[{'expires_at': '2018-08-29 00:00:00 UTC',
  'service_level': 'full',
  'system_information': {'request_code': 'kzho-isiA-8dwy-gZyq-gFNb-EC5l-od7s-JBai-RcaF-2hMb-
↪cirj-52rS-P4M3-2sRg-fuZa-/S5W-FkRn-RDSO-srVa-0x1X-q7KO-NkMY-380Y-dmW4-JfHG-
↪Q01x-so3N-NhdO-MoMj-Xw+B-bUdb-Q7VI-K+Hy-gSMF-kVpD-kCkO-v3Ay-a2/f-To9v-Lnxw-
↪3EdE-FEPa-yVMI-x/U4-EsUV-T1eq-LQsM-C88E-yPOS-RtVp-vDtD-zwEn-PAS7-/pSl-MGJ+-
↪jnUq-JllG-ux0+-seDZ-6X+v-rXBI-zHUx-go3p-K2ZO'},
  'updated_at': '2018-03-29 09:49:34 UTC'}]
```

Raises `MangroveError` – if the license has expired

```
{
  "status": 402,
  "status_text": "Payment Required",
  "errors": [
    {
      "code": "LICENSE-402-002",
      "type": "license",
      "metadata": {
        "reason": "License not found",
        "request_code": "P6jC-ns9H-X5ph-KZVg-finF-ttgV-2Jtl-ygPn-Ie/z-
↪VUBc-hYYz-GeLT-yTb4-UrkS-tr/s-w3uf-lzlG-Av34-3fnx-0gl2-8SnL-jaQt-0BJ+-bhKU-
↪zgWl-tu6j-kM4r-i84u-s2Qo-SR4P-JyEH-AIRh-psnw-d0zd-R+Nf-zrl+-8hWy-l3Db-HeD9-
↪6DY7-gwlO-lZjp-Opvu-pp5I-mxWQ-qDtS-WWTo-xj1K-hE9q-sukL-YEeK-OPWz-aaJl-0ZzB-
↪0sN2-6Gqz-soPd-lEXR-USDl-vDzJ-JltE-RWX+-HfZs-2Njd",
        "type": "private",
        "autofetch": False
      }
    }
  ]
}
```

`new_license` (*license_code*)

Update the licence

Parameters `license_code` – string provided by the Mangrove team

```
>>> mang.admin.new_license('fxbE-D9pK-h0t7-x+5r-B3G7-bY+x-AG6x-dzYW-tccq-HAt1-
↪Bkzb-JPVw-jsFd-zcvN-Nr15-vkIZ-ZK4J-yafW-niK9-RbaV-FGS9-oks5-zsLJ-yweZ-fg3K-
↪SAeT-jDWP-pDnj-bJ8P-ZjKh-Tskp-I/1A-Ymow-fV6s-fvXK-dliu-CHCY-lOrf-pBY0-VDgm-
↪IBaP-3Dz3-CiYS-4MVR-hQsO-KNfu-WK7d-7/6w-CTNW-A0HA-9rnB-im62-evcd-j7HS-KnnL-
↪K/aD-UNlU-5vO5-K9g=')
{
  'expires_at': '2018-03-30 00:00:00 UTC',
```

```
'service_level': 'full',
'system_information': None,
'updated_at': '2018-03-31 11:47:54 UTC'
}
```

Raises `HTTPError` – if you provide a wrong license

(Please use GUI)

tokens ()

List all tokens

```
>>> mang.admin.tokens()
[{'created_at': '2018-03-29T09:45:06.067Z',
  'expires_at': 1577833200,
  'id': '40a6a8b6-65b1-465e-b6c7-6c3021c30952',
  'name': 'token_jbp',
  'token': 'eyJ0eXAiOiJKV1QiLCJhbGciOiJIUzI1NiJ9.eyJzdWIiOiI0OTg4NC05M2UwLTQwNmMtYmQ3OS02NTUuIiwiaWF0IjoxNTIwODUyOTUwLmV5YXNjaWwifQ.looosUk2TuXOVXREmAvPoVnOxOkLaSLOT4TlOMK_yTA',
  'updated_at': '2018-03-29T09:45:06.067Z'
}]
```

```
users ( )
```

Retrieve all users

```
>>> mang.admin.users()
[admin(admin), Toto, Gillou]
```

```
versions()
```

Version information

```
>>> mang.admin.versions()
[
  {
    'name': 'atlas',
    'version': '0.0.1-alpha.1'
  }, {
    'name': 'license_authority',
    'version': u'1.3.2-rc1'
  }, {
    'name': 'dmgr',
    'version': '0.0.1'
  }, {
    'name': 'modeler',
    'version': '0.0.1-alpha.1'
  }, {
    'name': 'exporter',
    'version': '0.0.1'
  }, {
    'name': 'mang_sdk',
    'version': '1.2.1-30-ge792bea-dirty'
  }
]
```

6.2 Project

class mangrove_surface.wrapper.project.ProjectWrapper

Project resource

classifier (*name*)

Return classifier named *name*

Parameters *name* – project name

Raises **ClassifierDoesNotExist** – if there is no classifier named *name*

classifiers ()

List all classifiers

```
>>> pj.classifiers()
[
    Project_2018-03-20T15:39:18.120Z,
    Project_2018-03-20T15:40:02.880Z,
    Project_2018-03-20T15:40:45.242Z,
    MyClassifier
]
```

collection (*name*)

Return the collection named *name*

Parameters *name* – collection name

collections ()

List all collections

create_collection ()

Create a new collection

A collection stores similar schemas of data sets.

Warning: Expert method: it should be only use to store new data set schemas

default_feature_set ()

Return the default feature set

(see: mangrove.wrapper.feature_set)

description ()

Project description

schemas (*type=None*)

List all schemas

Parameters *type* – (*optional*) *type* could be `train`, `test` or `export`. It is to filter schemas of type *type*. By default all schemas are listed.

tags ()

Return the project tags

update_description (*new_description*)

Update the project description

Parameters *new_description* – the new project description

update_name (*new_name*)

Update the project name

Parameters **new_name** – the new project name

update_tags (*new_tags=[]*)

Update the project tags

Parameters **new_tags** – the list of new tags

6.3 Classifier

class mangrove_surface.wrapper.classifier.ClassifierWrapper

Classifier resource

A classifier provides

- the list relevant features (including level, weight, discretization attributes)
- the assessments over each train/test schemas
- method to export scores over
- method to improve classifier

add_schema (*type_schm, schema, name=None*)

Upload a new schema of datasets

Parameters

- **type_schm** – train, test or export
- **schema** – a python dictionary recording datasets like this

```
{
  "tags": ["dataset", "tag"],
  "datasets": [
    {
      "name": "Dataset Name",
      "filepath": "/path/to/dataset.csv",
      "tags": ["optional", "tags"],
      "central": True | False,
      "keys": ["index"], # optional if there is only
                        # one dataset
      "separator": ",", # could be `|`, ` `, `;` or ``
    }, ...
  ]
}
```

add_schema_and_export (*schema, name=None, modalities=[], bin_format='label', raw_variables=[], binned_variables=[], predicted_modality=False*)

Upload a new schema and export it

Parameters

- **schema** – a python dictionary of datasets (see `add_schema()` :)
- **name** – (*optional*) the schema name
- **modalities** – (*optional*) the modalities scored. If no modality is provided then scores are not provided (only variables)

- **raw_variables** – the list of variables to export as raw value
- **binned_variables** – the list of variables to export as binned value
- **bin_format** – (*default: label*) select how to express the binned variables. *label* (*default*) to express value as its intervals or groups, or *id* to express value as a concise value
- **predicted_modality** – provided a column with the predicted value if `predicted_modality==True` (*default* `predicted_modality==False`)

compatible_schemas (*test=True, export=True*)

List compatible schemas (with there type)

compute_assessments (*schm_name, outcome_modality=None*)

Compute assessment over schema named `schm_name` (focus on modality `outcome_modality`)

Parameters

- **schm_name** – name of the schema used to compute assessments
- **outcome_modality** – the modality used to compute assessments (by *default* assessments is computed over the main modality)

compute_export (*schm_name, export_name=None, modalities=[], bin_format='label', raw_variables=[], binned_variables=[], predicted_modality=False*)

Compute a new export

Parameters

- **schm_name** – the dataset schema which is exported
- **export_name** – name of the export
- **modalities** – (*optional*) the modalities scored. If no modality is provided then scores are not provided (only variables)
- **raw_variables** – the list of variables to export as raw value
- **binned_variables** – the list of variables to export as binned value
- **bin_format** – (*default: label*) select how to express the binned variables. *label* (*default*) to express value as its intervals or groups, or *id* to express value as a concise value
- **predicted_modality** – provided a column with the predicted value if `predicted_modality==True` (*default* `predicted_modality==False`)

discretization_attribute (**args, **kwargs*)

Return the discretization attribute of the contributive feature name

Parameters **name** – feature name

```
>>> classifier.discretization_attribute("Car_Type")
[
  {
    'coverage': 0.0248497,
    'frequency': 529,
    'target_distribution': {
      '0': 0.837429,
      '1': 0.162571
    },
    'value_list': ['Full-size luxury car']
  },
]
```



```
    ...  
]
```

download (*args, **kwargs)

Download the classifier

Parameters **filepath** – the filepath where store the classifier

exports ()

List all exports

feature (*args, **kwargs)

Information about feature name

It returns level, weight, discretization attributes.

Parameters **name** – feature name

```
>>> classifier.feature('Car_Type')  
{  
  'level': 0.103459,  
  'maximum_a_posteriori': True,  
  'name': 'Car_Type',  
  'nb_parts': 4,  
  'parts': [  
    {  
      'coverage': 0.0248497,  
      'frequency': 529,  
      'target_distribution': {  
        '0': 0.837429,  
        '1': 0.162571  
      },  
      'value_list': ['Full-size luxury car']  
    },  
    ...  
  ],  
  'weight': 0.832425  
}
```

feature_set (*args, **kwargs)

Return the underlying feature set

Note: This feature set could be used to change type, unused some features

features (*args, **kwargs)

List all the features used by the current classifier

```
>>> classifier.features()  
[  
  {  
    'level': 0.103459,  
    'maximum_a_posteriori': True,  
    'name': 'Car_Type',  
    'nb_parts': 4,  
    'parts': [  
      {  
        'coverage': 0.0248497,  
        'frequency': 529,  

```

```
        'target_distribution': {
            '0': 0.837429,
            '1': 0.162571
        },
        'value_list': ['Full-size luxury car']
    },
    ...
],
'weight': 0.832425
},
...
]
```

improve (*name=None, tags=[], nb_aggregates=None, maximum_features=None*)

Create a new classifier

Parameters

- **name** – (*optional*) classifier name
- **tags** – (*optional*) list of project tag
- **nb_aggregates** – (*optional*) number of aggregates generated for the new classifier
- **maximum_features** – (*optional*) maximal number of features used by the new classifier

Raises **MangroveError** – if the number of requested aggregates is provided and it is smaller than `.nb_aggregates()`

level (**args, **kwargs*)

Return the level of the feature named *name*

Parameters **name** – feature name

The level indicates the correlation between the feature and the outcome

nb_aggregates ()

Return the number of aggregates

outcome ()

Outcome field predicted by the current classifier

set_unused (**args, **kwargs*)

Set feature *name* unused

Parameters **name** – feature name

update_name (*new_name*)

Update the classifier name

Parameters **new_name** – new classifier name

weight (**args, **kwargs*)

Return the weight of the feature named *name*

Parameters **name** – feature name

The weight indicates how the feature discriminates more than others relevant features (with level > 0)

6.3.1 Assessment

class mangrove_surface.wrapper.classifier_evaluation_report.**ClassifierEvaluationReportWrapper**
Classifier Evaluation Report resource

ACC ()
Accuracy

Note:

This method has some alias:

- ACC
-

AUC (*args, **kwargs)
Area under curve

DOR ()
Diagnostic odds ratio

Note:

This method has some alias:

- DOR
-

F1_score (outcome_modality=None)
F1 score

Parameters **outcome_modality** – (optional) the modality

FDR (outcome_modality)
False discovery rate

Parameters **outcome_modality** – (optional) the modality

Note:

This method has some alias:

- FDR
-

FNR (outcome_modality=None)
False negative rate

Parameters **outcome_modality** – (optional) the modality

Note:

This method has some alias:

- FNR
 - miss_rate
-

FOR (outcome_modality)
False omission rate

Parameters `outcome_modality` – (*optional*) the modality

Note:

This method has some alias:

- `FOR`
-

FPR (*outcome_modality=None*)

False positive rate

Parameters `outcome_modality` – (*optional*) the modality

Note:

This method has some alias:

- `FPR`
 - `fall_out`
-

LRm ()

Negative Likelihood ratio

Note:

This method has some alias:

- `LRp`
-

LRp ()

Positive Likelihood ratio

Note:

This method has some alias:

- `LRp`
-

NPV (*outcome_modality*)

Negative predictive value

Parameters `outcome_modality` – (*optional*) the modality

Note:

This method has some alias:

- `NPV`
-

PPV (*outcome_modality=None*)

Precision

Parameters `outcome_modality` – (*optional*) the modality

Note:**This method has some alias:**

- `positive_predictive_value`
-

SPC (*outcome_modality=None*)

True negative rate

Parameters `outcome_modality` – (*optional*) the modality

Note:**This method has some alias:**

- `TNR`
 - `specificity`
 - `SPC`
-

TNR (*outcome_modality=None*)

True negative rate

Parameters `outcome_modality` – (*optional*) the modality

Note:**This method has some alias:**

- `TNR`
 - `specificity`
 - `SPC`
-

TPR (*outcome_modality=None*)

True positive rate

Parameters `outcome_modality` – (*optional*) the modality

Note:**This method has some alias:**

- `recall`
 - `TPR`
 - `sensitivity`
 - `probability_of_detection`
-

accuracy ()

Accuracy

Note:

This method has some alias:

- ACC
-

area_under_curve (*args, **kwargs)

Area under curve

auc (*args, **kwargs)

Area under curve

confusion_matrix (*args, **kwargs)

Confusion matrix

::

```
>>> ass.confusion_matrix()
{
  'matrix': [
    [13376, 1393],
    [ 683, 4084]
  ],
  'modalities': ['N', 'Y']
}
```

diagnostic_odds_ratio()

Diagnostic odds ratio

Note:

This method has some alias:

- DOR
-

fall_out (outcome_modality=None)

False positive rate

Parameters **outcome_modality** – (*optional*) the modality

Note:

This method has some alias:

- FPR
 - fall_out
-

false_discovery_rate (outcome_modality)

False discovery rate

Parameters **outcome_modality** – (*optional*) the modality

Note:

This method has some alias:

- FDR
-

false_negative (*outcome_modality*)

Number of false negative errors of the *outcome_modality*

False negative = incorrectly rejected

Parameters *outcome_modality* – (*optional*) compute the number of incorrect rejection of the modality

false_negative_rate (*outcome_modality=None*)

False negative rate

Parameters *outcome_modality* – (*optional*) the modality

Note:

This method has some alias:

- FNR
 - miss_rate
-

false_omission_rate (*outcome_modality*)

False omission rate

Parameters *outcome_modality* – (*optional*) the modality

Note:

This method has some alias:

- FOR
-

false_positive (*outcome_modality=None*)

Number of incorrect predictions

False positive = incorrectly identified

Parameters *outcome_modality* – (*optional*) compute the number of incorrect prediction associated to this modality

Raises **KeyError** – if the *outcome_modality* does not exist

```
>>> ass.false_positive()
2076

>>> ass.false_positive('Y')
4084
```

false_positive_rate (*outcome_modality=None*)

False positive rate

Parameters *outcome_modality* – (*optional*) the modality

Note:

This method has some alias:

- FPR
- fall_out

gini()

Gini coefficient

instances (*outcome_modality=None*)

Number of instances evaluated

lift_curve (*args, **kwargs)

Lift curve over the schema

Parameters using – is classifier or optimal; by default the lift curve associated to the classifier.

miss_rate (*outcome_modality=None*)

False negative rate

Parameters outcome_modality – (*optional*) the modality

Note:

This method has some alias:

- FNR
- miss_rate

negative_likelihood_ratio()

Negative Likelihood ratio

Note:

This method has some alias:

- LRp

negative_predictive_value (*outcome_modality*)

Negative predictive value

Parameters outcome_modality – (*optional*) the modality

Note:

This method has some alias:

- NPV

positive_likelihood_ratio()

Positive Likelihood ratio

Note:

This method has some alias:

- LRp

positive_predictive_value (*outcome_modality=None*)

Precision

Parameters `outcome_modality` – (*optional*) the modality

Note:

This method has some alias:

- `positive_predictive_value`
-

precision (`outcome_modality=None`)

Precision

Parameters `outcome_modality` – (*optional*) the modality

Note:

This method has some alias:

- `positive_predictive_value`
-

prevalence ()

Prevalence

probability_of_detection (`outcome_modality=None`)

True positive rate

Parameters `outcome_modality` – (*optional*) the modality

Note:

This method has some alias:

- `recall`
 - `TPR`
 - `sensitivity`
 - `probability_of_detection`
-

recall (`outcome_modality=None`)

True positive rate

Parameters `outcome_modality` – (*optional*) the modality

Note:

This method has some alias:

- `recall`
 - `TPR`
 - `sensitivity`
 - `probability_of_detection`
-

sensitivity (`outcome_modality=None`)

True positive rate

Parameters `outcome_modality` – (*optional*) the modality

Note:

This method has some alias:

- `recall`
 - `TPR`
 - `sensitivity`
 - `probability_of_detection`
-

specificity (*outcome_modality=None*)

True negative rate

Parameters `outcome_modality` – (*optional*) the modality

Note:

This method has some alias:

- `TNR`
 - `specificity`
 - `SPC`
-

target_rate (*outcome_modality*)

Target rate of the modality `outcome_modality`

Parameters `outcome_modality` – a modality

true_negative (*outcome_modality*)

Number of true negative errors of the `outcome_modality`

True negative = correctly rejected

Parameters `outcome_modality` – (*optional*) compute the number of correct rejection of the modality

true_negative_rate (*outcome_modality=None*)

True negative rate

Parameters `outcome_modality` – (*optional*) the modality

Note:

This method has some alias:

- `TNR`
 - `specificity`
 - `SPC`
-

true_positive (*outcome_modality=None*)

Number of correct predictions

True positive = correctly identified

Parameters `outcome_modality` – (*optional*) compute the number of correct prediction associated to this modality

Raises **KeyError** – if the `outcome_modality` does not exist

```
>>> ass.true_positive()
17460

>>> ass.true_positive('Y')
4084
```

true_positive_rate (*outcome_modality=None*)

True positive rate

Parameters `outcome_modality` – (*optional*) the modality

Note:

This method has some alias:

- `recall`
 - `TPR`
 - `sensitivity`
 - `probability_of_detection`
-

6.4 Export

class `mangrove_surface.wrapper.export.ExportWrapper`

binned_variables ()

List the binned variables

classifier ()

Return the classifier used

download (*args, **kwargs)

Download the export

Parameters `filepath` – the filepath where store the classifier

instances_submitted (*args, **kwargs)

Number of instances submitted to export

push_s3 (*bucket, key*)

Push the current export to S3

Parameters

- **bucket** – the S3 bucket
- **key** – the S3 key

raw_variables ()

List the raw variables

6.5 Feature set

class mangrove_surface.wrapper.feature_set.**FeatureSetWrapper** (*feature_set_resource*,
collection)

Feature set resource

A feature set is a set of frames (one for each data set). A frame contains variables and its metadata (type, use or not).

It is used to customize data, generate aggregates and to train classifiers.

central (**args*, ***kwargs*)

Return the central frame

The central frame is the one used to train classifiers.

clone (*new_name=None*, *tags=None*)

Clone the current feature set.

fit_classifier (*name=None*, *tags=[]*, *nb_aggregates=None*, *maximum_features=None*)

Fit a new classifier

Parameters

- **name** – (*optional*) classifier name (by default the name will be the project name concatenated with the current time)
- **tags** – the classifier tags
- **nb_aggregates** – used to generates *nb_aggregates* aggregates on the central frame used to train the classifier
- **maximum_features** – used to allow at most *maximum_features* features in the new classifier

frame (**args*, ***kwargs*)

Return the frame named *name*

Parameters *name* – data (set) frame name

frames (**args*, ***kwargs*)

List all frames

generate_aggregates (**args*, ***kwargs*)

Generate a new feature set with *n* aggregates

Parameters *n* – number of aggregates requested (a non-negative integer)

is_modified (**args*, ***kwargs*)

Indicates if the current feature set has been modified

save (**args*, ***kwargs*)

Save all the modifications (change variables type, set unused, *etc.*)

Warning: If `clone = False` the method overrides the current feature set resource

Raises Exception – if `clone = False` and the current feature set is the default one.

6.5.1 Frame

class FeatureSetWrapper._Frame (dataset, change_type, fs)

features (filt=<function <lambda>>, id=False)

List features of the current frame

```
>>> fs.features()
[
  {
    'name': 'Flag_Prospect',
    'type': 'categorical',
    'use': True
  },
  {
    'name': 'LABEL',
    'type': 'continuous',
    'use': True
  },
  ...
]
```

Parameters **filt** – (optional) a function that can be used to filter features

```
>>> fs.features(filter=lambda feat: fs.is_categorical(feat))
[
  {
    'name': 'Flag_Prospect',
    'type': 'categorical',
    'use': True
  },
  ...
]
```

or:

```
>>> fs.features(filter=lambda feat: feat["name"].startswith("Foo"))
[
  {
    'name': 'FooBar',
    'type': 'categorical',
    'use': True
  },
  {
    'name': 'FooFoo',
    'type': 'continuous',
    'use': False
  },
  ...
]
```

is_categorical (variable)

Indicates if the feature variable is categorical or not

:param variable:: feature name

is_central()
Indicates if the frame is central

is_change_type_allowed()
Indicate if the frame allows to change feature type
It is forbidden to change type of peripheral frame features if there is aggregates in the central frame

is_continuous(*variable*)
Indicates if the feature *variable* is continuous or not
:param variable:: feature name

is_modified()
Indicates if the frame has been modified

is_used(*variable*)
Return if the feature is used or not

modalities(*name*)
List modalities of the feature name
Parameters *name* – feature name

set_categorical(*variable*)
Change the type of the feature *variable* to categorical
Parameters *variable* – the feature name
Raises **MangroveChangeForbidden** – if change type is not allowed

set_continuous(*variable*)
Change the type of the feature *variable* to continuous
Parameters *variable* – the feature name
Raises **MangroveChangeForbidden** – if change type is not allowed

set_unused(*variable*)
Set unused the feature *variable*
Parameters *variable* – the feature name

set_used(*variable*)
Set used the feature *variable*
Parameters *variable* – the feature name

type(*variable*)
Return the type of the feature *variable* The type could be `categorical` or `continuous` (other types can be provided like `timestamps` but they are not managed)
Parameters *variable* – the feature

6.6 Collection

class `mangrove_surface.wrapper.collection.CollectionWrapper`
Collection
A collection is a set of dataset schemas which are similars.
create_schema(*type_schm*, *schema*, *name=None*, *check=True*)
Create a new schema into the current collection

Parameters

- **type_schm** – train, test or export
- **schema** – a python dictionary recording datasets like this

```
{
  "tags": ["dataset", "tag"],
  "datasets": [
    {
      "name": "Dataset Name",
      "filepath": "/path/to/dataset.csv",
      "tags": ["optional", "tags"],
      "central": True | False,
      "keys": ["index"], # optional if there is only
                        # one dataset
      "separator": ",", # could be `|`, `,`, `;` or ``
    }, ...
  ]
}
```

schema (*name*)

Return schema named name

Parameters **name** – the name of the requested schema

schemas ()

List all schemas of the current collection

6.7 Logger

This logger is configured to log SDK behaviors.

Each api request and answer are logged at level DEBUG and each api resource created is logged at level INFO. Warnings and errors are logged at the expected level: WARNING and ERROR.

If you meet some unexpected behavior or bugs then please use the following lines to store in a file the behavior:

```
>>> import os.path as path
>>> path_logfile = path.join( # similar to /home/my_nickname/.mang-sdk.log
...     path.expanduser('~'),
...     '.mang-sdk.log'
... )
>>> from mangrove.logger import logger, logging
>>> from logging.handlers import RotatingFileHandler
>>> file_handler = RotatingFileHandler(
...     path_to_your_logfile, 'a', 1000000, 1
... )
>>> file_handler.setLevel(logging.DEBUG)
>>> logger.addHandler(file_handler)
```

Run your script/code (with the unexpected behavior) and send it to our support (support@mangrove.ai).

6.8 Indices and tables

- [genindex](#)

- [modindex](#)
- [search](#)

m

- mangrove_surface, [13](#)
- mangrove_surface.logger, [35](#)
- mangrove_surface.wrapper.classifier, [19](#)
- mangrove_surface.wrapper.classifier_evaluation_report,
 [23](#)
- mangrove_surface.wrapper.collection, [34](#)
- mangrove_surface.wrapper.export, [31](#)
- mangrove_surface.wrapper.feature_set,
 [32](#)
- mangrove_surface.wrapper.project, [18](#)

A

ACC() (mangrove_surface.wrapper.classifier_evaluation_report.ClassifierEvaluationReportWrapper
 method), 23
 accuracy() (mangrove_surface.wrapper.classifier_evaluation_report.ClassifierEvaluationReportWrapper
 method), 25
 add_schema() (mangrove_surface.wrapper.classifier.ClassifierWrapper
 method), 19
 add_schema_and_export() (mangrove_surface.wrapper.classifier.ClassifierWrapper
 method), 19
 area_under_curve() (mangrove_surface.wrapper.classifier_evaluation_report.ClassifierEvaluationReportWrapper
 method), 26
 AUC() (mangrove_surface.wrapper.classifier_evaluation_report.ClassifierEvaluationReportWrapper
 method), 23
 auc() (mangrove_surface.wrapper.classifier_evaluation_report.ClassifierEvaluationReportWrapper
 method), 26
 collection() (mangrove_surface.wrapper.project.ProjectWrapper
 method), 18
 collections() (mangrove_surface.wrapper.project.ProjectWrapper
 method), 18
 CollectionWrapper (class in mangrove_surface.wrapper.collection), 34
 compatible_schemas() (mangrove_surface.wrapper.classifier.ClassifierWrapper
 method), 20
 compute_assessments() (mangrove_surface.wrapper.classifier.ClassifierWrapper
 method), 20
 compute_export() (mangrove_surface.wrapper.classifier_evaluation_report.ClassifierEvaluationReportWrapper
 method), 20
 confusion_matrix() (mangrove_surface.wrapper.classifier_evaluation_report.ClassifierEvaluationReportWrapper
 method), 26

B

binned_variables() (mangrove_surface.wrapper.export.ExportWrapper
 method), 31
 create_collection() (mangrove_surface.wrapper.project.ProjectWrapper
 method), 18
 create_project() (mangrove_surface.SurfaceClient
 method), 14
 create_schema() (mangrove_surface.wrapper.collection.CollectionWrapper
 method), 34

C

central() (mangrove_surface.wrapper.feature_set.FeatureSetWrapper
 method), 32
 classifier() (mangrove_surface.wrapper.export.ExportWrapper
 method), 31
 classifier() (mangrove_surface.wrapper.project.ProjectWrapper
 method), 18
 ClassifierEvaluationReportWrapper (class in mangrove_surface.wrapper.classifier_evaluation_report), 23
 classifiers() (mangrove_surface.wrapper.project.ProjectWrapper
 method), 18
 ClassifierWrapper (class in mangrove_surface.wrapper.classifier), 19
 clone() (mangrove_surface.wrapper.feature_set.FeatureSetWrapper
 method), 32
 create_token() (mangrove_surface.SurfaceClient._Admin
 method), 15
 create_user() (mangrove_surface.SurfaceClient._Admin
 method), 15

D

default_feature_set() (mangrove_surface.wrapper.project.ProjectWrapper
 method), 18
 delete_users() (mangrove_surface.SurfaceClient._Admin
 method), 15
 description() (mangrove_surface.wrapper.project.ProjectWrapper
 method), 18
 diagnostic_odds_ratio() (mangrove_surface.wrapper.classifier_evaluation_report.ClassifierEvaluationReportWrapper
 method), 26

method), 26
 discretization_attribute() (mangrove_surface.wrapper.classifier.ClassifierWrapper method), 23
 DOR() (mangrove_surface.wrapper.classifier_evaluation_report.ClassifierEvaluationReportWrapper method), 23
 download() (mangrove_surface.wrapper.classifier.ClassifierWrapper method), 21
 download() (mangrove_surface.wrapper.export.ExportWrapper method), 31
 FNR() (mangrove_surface.wrapper.classifier_evaluation_report.ClassifierEvaluationReportWrapper method), 23
 FOR() (mangrove_surface.wrapper.classifier_evaluation_report.ClassifierEvaluationReportWrapper method), 23
 FPR() (mangrove_surface.wrapper.classifier_evaluation_report.ClassifierEvaluationReportWrapper method), 24
 feature() (mangrove_surface.wrapper.classifier.ClassifierWrapper method), 21
 features() (mangrove_surface.wrapper.feature_set.FeatureSetWrapper method), 32
 features() (mangrove_surface.wrapper.feature_set.FeatureSetWrapper method), 32

E

exports() (mangrove_surface.wrapper.classifier.ClassifierWrapper method), 21
 ExportWrapper (class in mangrove_surface.wrapper.export), 31
 aggregate_aggregates() (mangrove_surface.wrapper.feature_set.FeatureSetWrapper method), 32
 gini() (mangrove_surface.wrapper.classifier_evaluation_report.ClassifierEvaluationReportWrapper method), 28

G

F

F1_score() (mangrove_surface.wrapper.classifier_evaluation_report.ClassifierEvaluationReportWrapper method), 23
 fall_out() (mangrove_surface.wrapper.classifier_evaluation_report.ClassifierEvaluationReportWrapper method), 26
 false_discovery_rate() (mangrove_surface.wrapper.classifier_evaluation_report.ClassifierEvaluationReportWrapper method), 26
 false_negative() (mangrove_surface.wrapper.classifier_evaluation_report.ClassifierEvaluationReportWrapper method), 26
 false_negative_rate() (mangrove_surface.wrapper.classifier_evaluation_report.ClassifierEvaluationReportWrapper method), 27
 false_omission_rate() (mangrove_surface.wrapper.classifier_evaluation_report.ClassifierEvaluationReportWrapper method), 27
 false_positive() (mangrove_surface.wrapper.classifier_evaluation_report.ClassifierEvaluationReportWrapper method), 27
 false_positive_rate() (mangrove_surface.wrapper.classifier_evaluation_report.ClassifierEvaluationReportWrapper method), 27
 FDR() (mangrove_surface.wrapper.classifier_evaluation_report.ClassifierEvaluationReportWrapper method), 23
 feature() (mangrove_surface.wrapper.classifier.ClassifierWrapper method), 21
 feature_set() (mangrove_surface.wrapper.classifier.ClassifierWrapper method), 21
 features() (mangrove_surface.wrapper.classifier.ClassifierWrapper method), 21
 features() (mangrove_surface.wrapper.feature_set.FeatureSetWrapper method), 33
 FeatureSetWrapper (class in mangrove_surface.wrapper.feature_set), 32
 FeatureSetWrapper.Frame (class in mangrove_surface.wrapper.feature_set), 33
 fit_classifier() (mangrove_surface.wrapper.feature_set.FeatureSetWrapper method), 32
 improve() (mangrove_surface.wrapper.classifier.ClassifierWrapper method), 23
 instances() (mangrove_surface.wrapper.classifier_evaluation_report.ClassifierEvaluationReportWrapper method), 28
 is_categorical() (mangrove_surface.wrapper.feature_set.FeatureSetWrapper method), 33
 is_change_type_allowed() (mangrove_surface.wrapper.feature_set.FeatureSetWrapper.Frame method), 34
 is_continuous() (mangrove_surface.wrapper.feature_set.FeatureSetWrapper method), 34
 is_modified() (mangrove_surface.wrapper.feature_set.FeatureSetWrapper method), 33
 is_modified() (mangrove_surface.wrapper.feature_set.FeatureSetWrapper.Frame method), 31
 is_used() (mangrove_surface.wrapper.feature_set.FeatureSetWrapper.Frame method), 34
 level() (mangrove_surface.wrapper.classifier.ClassifierWrapper method), 22
 license_information() (mangrove_surface.SurfaceClient.Admin method), 15
 lift_curve() (mangrove_surface.wrapper.classifier_evaluation_report.ClassifierEvaluationReportWrapper method), 28
 LRm() (mangrove_surface.wrapper.classifier_evaluation_report.ClassifierEvaluationReportWrapper method), 24
 log_loss() (mangrove_surface.wrapper.classifier_evaluation_report.ClassifierEvaluationReportWrapper method), 24

M

[mangrove_surface](#) (module), 13
[mangrove_surface.logger](#) (module), 35
[mangrove_surface.wrapper.classifier](#) (module), 19
[mangrove_surface.wrapper.classifier_evaluation_report](#) (module), 23
[mangrove_surface.wrapper.collection](#) (module), 34
[mangrove_surface.wrapper.export](#) (module), 31
[mangrove_surface.wrapper.feature_set](#) (module), 32
[mangrove_surface.wrapper.project](#) (module), 18
[miss_rate\(\)](#) ([mangrove_surface.wrapper.classifier_evaluation_report.ClassifierEvaluationReportWrapper](#) method), 28
[modalities\(\)](#) ([mangrove_surface.wrapper.feature_set.FeatureSetWrapper](#) method), 34

N

[nb_aggregates\(\)](#) ([mangrove_surface.wrapper.classifier.ClassifierWrapper](#) method), 22
[negative_likelihood_ratio\(\)](#) ([mangrove_surface.wrapper.classifier_evaluation_report.ClassifierEvaluationReportWrapper](#) method), 28
[negative_predictive_value\(\)](#) ([mangrove_surface.wrapper.classifier_evaluation_report.ClassifierEvaluationReportWrapper](#) method), 28
[new_license\(\)](#) ([mangrove_surface.SurfaceClient._Admin](#) method), 16
[NPV\(\)](#) ([mangrove_surface.wrapper.classifier_evaluation_report.ClassifierEvaluationReportWrapper](#) method), 24

O

[outcome\(\)](#) ([mangrove_surface.wrapper.classifier.ClassifierWrapper](#) method), 22

P

[positive_likelihood_ratio\(\)](#) ([mangrove_surface.wrapper.classifier_evaluation_report.ClassifierEvaluationReportWrapper](#) method), 28
[positive_predictive_value\(\)](#) ([mangrove_surface.wrapper.classifier_evaluation_report.ClassifierEvaluationReportWrapper](#) method), 28
[PPV\(\)](#) ([mangrove_surface.wrapper.classifier_evaluation_report.ClassifierEvaluationReportWrapper](#) method), 24
[precision\(\)](#) ([mangrove_surface.wrapper.classifier_evaluation_report.ClassifierEvaluationReportWrapper](#) method), 29
[prevalence\(\)](#) ([mangrove_surface.wrapper.classifier_evaluation_report.ClassifierEvaluationReportWrapper](#) method), 29
[probability_of_detection\(\)](#) ([mangrove_surface.wrapper.classifier_evaluation_report.ClassifierEvaluationReportWrapper](#) method), 29
[project\(\)](#) ([mangrove_surface.SurfaceClient](#) method), 15
[projects\(\)](#) ([mangrove_surface.SurfaceClient](#) method), 15
[ProjectWrapper](#) (class in [mangrove_surface.wrapper.project](#)), 18

[push_s3\(\)](#) ([mangrove_surface.wrapper.export.ExportWrapper](#) method), 31

R

[raw_variables\(\)](#) ([mangrove_surface.wrapper.export.ExportWrapper](#) method), 31
[recall\(\)](#) ([mangrove_surface.wrapper.classifier_evaluation_report.ClassifierEvaluationReportWrapper](#) method), 29

S

[schemas\(\)](#) ([mangrove_surface.wrapper.collection.CollectionWrapper](#) method), 32
[schemas\(\)](#) ([mangrove_surface.wrapper.project.ProjectWrapper](#) method), 18
[sensitivity\(\)](#) ([mangrove_surface.wrapper.classifier_evaluation_report.ClassifierEvaluationReportWrapper](#) method), 29
[set_categorical\(\)](#) ([mangrove_surface.wrapper.feature_set.FeatureSetWrapper](#) method), 34
[set_continuous\(\)](#) ([mangrove_surface.wrapper.feature_set.FeatureSetWrapper._Frame](#) method), 34
[set_unused\(\)](#) ([mangrove_surface.wrapper.classifier.ClassifierWrapper](#) method), 22
[set_unused\(\)](#) ([mangrove_surface.wrapper.feature_set.FeatureSetWrapper._Frame](#) method), 34
[set_used\(\)](#) ([mangrove_surface.wrapper.feature_set.FeatureSetWrapper._Frame](#) method), 34
[SPC\(\)](#) ([mangrove_surface.wrapper.classifier_evaluation_report.ClassifierEvaluationReportWrapper](#) method), 25
[specificity\(\)](#) ([mangrove_surface.wrapper.classifier_evaluation_report.ClassifierEvaluationReportWrapper](#) method), 30
[SurfaceClient](#) (class in [mangrove_surface](#)), 13
[SurfaceClient._Admin](#) (class in [mangrove_surface](#)), 15

T

[tags\(\)](#) ([mangrove_surface.wrapper.project.ProjectWrapper](#) method), 18
[target_rate\(\)](#) ([mangrove_surface.wrapper.classifier_evaluation_report.ClassifierEvaluationReportWrapper](#) method), 30
[TNR\(\)](#) ([mangrove_surface.wrapper.classifier_evaluation_report.ClassifierEvaluationReportWrapper](#) method), 25
[tokens\(\)](#) ([mangrove_surface.SurfaceClient._Admin](#) method), 17
[TPR\(\)](#) ([mangrove_surface.wrapper.classifier_evaluation_report.ClassifierEvaluationReportWrapper](#) method), 25
[true_negative\(\)](#) ([mangrove_surface.wrapper.classifier_evaluation_report.ClassifierEvaluationReportWrapper](#) method), 30
[true_negative_rate\(\)](#) ([mangrove_surface.wrapper.classifier_evaluation_report.ClassifierEvaluationReportWrapper](#) method), 30

`true_positive()` (mangrove_surface.wrapper.classifier_evaluation_report.ClassifierEvaluationReportWrapper method), [30](#)

`true_positive_rate()` (mangrove_surface.wrapper.classifier_evaluation_report.ClassifierEvaluationReportWrapper method), [31](#)

`type()` (mangrove_surface.wrapper.feature_set.FeatureSetWrapper._Frame method), [34](#)

U

`update_description()` (mangrove_surface.wrapper.project.ProjectWrapper method), [18](#)

`update_name()` (mangrove_surface.wrapper.classifier.ClassifierWrapper method), [22](#)

`update_name()` (mangrove_surface.wrapper.project.ProjectWrapper method), [18](#)

`update_tags()` (mangrove_surface.wrapper.project.ProjectWrapper method), [19](#)

`users()` (mangrove_surface.SurfaceClient._Admin method), [17](#)

V

`versions()` (mangrove_surface.SurfaceClient._Admin method), [17](#)

W

`weight()` (mangrove_surface.wrapper.classifier.ClassifierWrapper method), [22](#)