
PathMe-Viewer Documentation

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PathMe Viewer.

A plugin for PathMe that allows to explore pathway knowledge.

INSTALLATION

1. `pathme_viewer` can be installed with the following commands:

```
$ python3 -m pip install git+https://github.com/ComPath/PathMe-Viewer.git@master
```

2. or in editable mode with:

```
$ git clone https://github.com/ComPath/PathMe-Viewer.git
$ cd PathMe-Viewer
$ python3 -m pip install -e .
```


DATABASE

The web application requires to load the pathways from the databases in the BEL. Thus, it is required to the following command to load the database (note that the first time it runs might take a couple of hours).

```
python3 -m pathme_viewer manage load_database
```

In order to check the status of the database, you can run:

```
python3 -m pathme_viewer manage summarize
```

The content of the database can be erased by running:

```
python3 -m pathme_viewer manage drop
```


DEPLOYMENT

Once the desired pathway databases are loaded, you can deploy the web application by running:

```
python3 -m pip install pathme_viewer web
```

Note that the database runs by default in the following port: <http://0.0.0.0:5000/>. The Flask host and port can be modified by changing the default parameters (run: “python3 -m pathme_viewer web -help” for more info).

COMMAND LINE INTERFACE

PathMe-Viewer commands.

4.1 pathme_viewer

PathMe

```
pathme_viewer [OPTIONS] COMMAND [ARGS]...
```

4.1.1 manage

Manage the database.

```
pathme_viewer manage [OPTIONS] COMMAND [ARGS]...
```

Options

-c, --connection <connection>
Cache connection. Defaults to `sqlite:///home/docs/.bio2bel/bio2bel.db`

drop

Delete all database entries

```
pathme_viewer manage drop [OPTIONS]
```

Options

-v, --debug
Turn on debugging.

-y, --yes

-c, --connection <connection>
Cache connection. Defaults to `sqlite:///home/docs/.bio2bel/bio2bel.db`

export-to-tsv

Export pathways to tsv

```
pathme_viewer manage export-to-tsv [OPTIONS]
```

Options

-c, --connection <connection>
Defaults to sqlite:///home/docs/.bio2bel/bio2bel.db

-a, --all

load

Load Pathways

```
pathme_viewer manage load [OPTIONS]
```

Options

-c, --connection <connection>
Cache connection. Defaults to sqlite:///home/docs/.bio2bel/bio2bel.db

-kp, --kegg_path <kegg_path>
KEGG data folder. Defaults to /home/docs/.pathme/kegg

-rp, --reactome_path <reactome_path>
Reactome data folder. Defaults to /home/docs/.pathme/reactome

-wp, --wikipathways_path <wikipathways_path>
WikiPathways data folder. Defaults to /home/docs/.pathme/wikipathways

-f, --flatten <flatten>
Flat complexes/composites. Defaults to False

-y, --yes
Skip confirmation

summarize

Summarizes Entries in Database

```
pathme_viewer manage summarize [OPTIONS]
```

Options

-c, --connection <connection>
Defaults to sqlite:///home/docs/.bio2bel/bio2bel.db

4.1.2 web

Run web service.

```
pathme_viewer web [OPTIONS]
```

Options

--host <host>
Flask host. Defaults to 0.0.0.0

--port <port>
Flask port. Defaults to 5000

--template <template>
Defaults to “./templates”

--static <static>
Defaults to “./static”

-v, --verbose

DATABASE MODELS

PathMe-Viewer database models.

PathMe models.

```
class pathme_viewer.models.Pathway(**kwargs)
```

Represents a pathway network in BEL format harmonized by ComPath

A simple constructor that allows initialization from kwargs.

Sets attributes on the constructed instance using the names and values in `kwargs`.

Only keys that are present as attributes of the instance's class are allowed. These could be, for example, any mapped columns or relationships.

name

Pathway name

resource_name

Database of origin

pathway_id

Pathway identifier in database of origin

number_of_nodes

Number of nodes

number_of_edges

Number of edges

version

Version of the BEL file

authors

Authors of the underlying BEL file

contact

Contact email from the underlying pathway

description

Descriptive text from the pathway

pybel_version

Version of PyBEL

blob

A pickled version of this pathway

display_name

Return pathway name.

pathway_tuple

Return pathway name.

as_bel ()

Get this network and loads it into a BELGraph.

Return type pybel.BELGraph

DATABASE MANAGER

PathMe-Viewer database manager.

This module contains the PathMe database manager.

```
class pathme_viewer.manager.Manager (engine, session)
```

Database manager.

Init PathMe manager.

```
create_all (check_first=True)
```

Create tables for PathMe.

```
drop_all (check_first=True)
```

Drop all tables for PathMe.

```
count_pathways ()
```

Count the pathways in the database.

Return type `int`

```
count_pathways_by_resource ()
```

Count the pathways in the database grouping by resource.

Return type `int`

```
get_all_pathways ()
```

Get all pathways in the database.

Return type `list[Pathway]`

```
get_all_pathway_graphs ()
```

Get all pathway graphs.

Return type `list[pybel.BELGraph]`

```
get_pathway_by_id (pathway_id, resource_name)
```

Get pathway by canonical identifier.

Parameters

- **pathway_id** (*str*) – pathway identifier
- **resource_name** (*str*) – name of the database

Return type `Optional[Pathway]`

```
get_pathway_by_name (pathway_name, resource_name)
```

Get pathway by name.

Parameters

- **pathway_name** (*str*) – pathway identifier
- **resource_name** (*str*) – name of the database

Return type Optional[*Pathway*]

get_pathways_from_resource (*resource_name*)

Get pathways from a given database.

Parameters **resource_name** (*str*) – name of the database

Return type Optional[list[*Pathway*]]

create_pathway (*pathway_dict*)

Create pathway.

Parameters **pathway_dict** (*dict*) – pathway identifier

Return type *Pathway*

delete_pathway (*pathway_id*, *resource_name*)

Delete a pathway.

Parameters

- **pathway_id** (*str*) – pathway identifier
- **resource_name** (*str*) – name of the database

Return type *bool*

delete_all_pathways ()

Delete all the pathways.

delete_pathways_from_resource (*resource_name*)

Delete pathways from a given database.

Parameters **resource_name** (*str*) – name of the database

Return type *bool*

get_or_create_pathway (*pathway_dict*)

Get or create pathway.

Parameters **pathway_dict** (*dict*) – pathway info

Return type *Pathway*

query_pathway_by_name (*query*, *limit=None*)

Return all pathways having the query in their names.

Parameters

- **query** (*str*) – query string
- **limit** (*Optional[int]*) – limit result query

Return type list[*Pathway*]

query_pathway_by_name_and_resource (*query*, *resource_name*, *limit=None*)

Return all pathways having the query in their names.

Parameters

- **query** (*str*) – query string
- **resource_name** (*str*) – database name
- **limit** (*Optional[int]*) – limit result query

Return type `list[Pathway]`

- genindex

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