Bio2BEL WikiPathways Documentation

Release 0.2.3

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Contents:

Manager

Database Manager and useful query methods.

Command Line Interface

The command line interface allows you to communicate with the package and perform basic functions such as:

- Populate the database: `python3 -m bio2bel_wikipathways populate`. By default the database is reset every time is populated. However, another optional parameter "–reset-db=False", allows you to avoid the reset. More logging can be activated by added "-vv" or "-v" as an argument.

- Drop the database: `python3 -m bio2bel_wikipathways drop`. More logging can be activated by added "-vv" or "-v" as an argument.

- Export gene sets as an excel file: `python3 -m bio2bel_wikipathways export`.

Models

Database models.

Constants

This module contains all the constants used in this package.

Parser

This module contains all methods that allow to parse the requests to the WikiPathways API

CHAPTER 6

Web

This module contains the web application to explore the database

CHAPTER 7

Indices and tables

- genindex
- modindex
- search