Bio2BEL PubChem Documentation

Release 0.0.1-dev

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Bio2BEL is a project aimed at integrating biological databases and other structured knowledge sources.

Because they come from a wide variety of schemata, this package provides tools for converting them into the unified framework of Biological Expression Language.

This package provides guidelines, tutorials, and tools for making standardized `bio2bel` packages as well as a unifying framework for integrating them.

CHAPTER 1

Installation

1.1 Easiest

Download the latest stable code from PyPI with:

```
$ python3 -m pip install bio2bel
```

1.2 Get the Latest

Download the most recent code from GitHub with:

```
$ python3 -m pip install git+https://github.com/bio2bel/bio2bel.git
```

1.3 For Developers

Clone the repository from GitHub and install in editable mode with:

```
$ git clone https://github.com/bio2bel/bio2bel.git
$ cd bio2bel
$ python3 -m pip install -e .
```

CHAPTER 2

Testing

Bio2BEL is tested with Python3 on Linux using Travis CI.

CHAPTER 3

Table of Contents

CHAPTER 4

Indices and tables

- genindex
- modindex
- search

Python Module Index

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bio2bel, ??

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