
jbiot Documentation

Release

KongDeju

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This is the documentation of **jbiot**.

Note: This is the main page of your project's [Sphinx](#) documentation. It is formatted in [reStructuredText](#). Add additional pages by creating `rst`-files in `docs` and adding them to the `toctree` below. Use then [references](#) in order to link them from this page, e.g. [authors](#) and [Changelog](#).

It is also possible to refer to the documentation of other Python packages with the [Python domain syntax](#). By default you can reference the documentation of [Sphinx](#), [Python](#), [NumPy](#), [SciPy](#), [matplotlib](#), [Pandas](#), [Scikit-Learn](#). You can add more by extending the `intersphinx_mapping` in your `Sphinx`'s `conf.py`.

The pretty useful extension [autodoc](#) is activated by default and lets you include documentation from docstrings. Docstrings can be written in [Google](#) (recommended!), [NumPy](#) and [classical](#) style.

CHAPTER 1

Contents

1.1 License

Copyright 2017 KongDeju

1.2 Developers

- KongDeju <kongdeju@gene.ac>

1.3 Changelog

1.3.1 Version 0.1

- Feature A added
- FIX: nasty bug #1729 fixed
- add your changes here!

1.4 jbiot

1.4.1 jbiot package

Subpackages

jbiot.subjobs package

Submodules

jbiot.subjobs.cmdPara module

jbiot.subjobs.cmdStep module

jbiot.subjobs.cs module

jbiot.subjobs.csub module

jbiot.subjobs.lsub module

jbiot.subjobs.msub module

Module contents

Submodules

jbiot.active module

jbiot.arrange module

jbiot.dict2yaml module

jbiot.filterTable module

jbiot.jbiotWorker module

jbiot.logrun module

jbiot.logrun2 module

jbiot.md2html module

jbiot.render module

jbiot.skeleton module

jbiot.smartFqs module

Module contents

CHAPTER 2

Indices and tables

- `genindex`
- `modindex`
- `search`