

Annotation comparison Vetea Jacot

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Chapter 1

Namespace Index

1.1 Namespace List

Here is a list of all documented namespaces with brief descriptions:

CDScompare	
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Chapter 2

Namespace Documentation

2.1 CDScompare Namespace Reference

Created on Thu May 2 11:57:29 2024.

Functions

- `main()`

Variables

- `script_dir = os.path.join(os.path.dirname(os.path.abspath(__file__)), "python_util")`

2.1.1 Detailed Description

Created on Thu May 2 11:57:29 2024.

This script is used to compute the similarities between two or more structural annotations of a same genome.

Author

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It expects as input the paths to the annotation files (in GFF format), displays the computed similarities between all annotation pairs, and creates a results CSV file detailing the loci comparisons between the annotations.

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