

Resource Summary Report

Generated by [RRID](#) on Aug 12, 2026

[findCPcli](#)

RRID:SCR_023391

Type: Tool

Proper Citation

findCPcli (RRID:SCR_023391)

Resource Information

URL: <https://github.com/findCP/findCPcli>

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Description: Software command line Python package for computation of chokepoint reactions in genome scale metabolic models. Used to compute chokepoints by taking into account both topology and dynamic information of network. Can compute and remove dead-end metabolites, find essential reactions and update flux bounds of reactions according to results of Flux Variability Analysis.

Resource Type: software toolkit, software resource

Defining Citation: [DOI:10.1007/978-3-030-60327-4_6](https://doi.org/10.1007/978-3-030-60327-4_6)

Keywords: compute chokepoints, chokepoint reactions, genome scale metabolic models, network topology and dynamic information,

Availability: Free, Available for download, Freely available

Resource Name: findCPcli

Resource ID: SCR_023391

Alternate URLs: <https://findcpcli.readthedocs.io/en/latest/>

License: GNU GPL v3.0

Record Creation Time: 20230321T180026+0000

Record Last Update: 20260810T040837+0000

Ratings and Alerts

No rating or validation information has been found for findCPcli.

No alerts have been found for findCPcli.

Data and Source Information

Source: [SciCrunch Registry](#)

Usage and Citation Metrics

We found 1 mentions in open access literature.

Listed below are recent publications. The full list is available at [RRID](#) .

Bannerman BP, et al. (2023) Mycobacterial metabolic model development for drug target identification. GigaByte (Hong Kong, China), 2023, gigabyte80.