

Resource Summary Report

Generated by [dkNET](#) on Apr 29, 2025

NetCoffee

RRID:SCR_012095

Type: Tool

Proper Citation

NetCoffee (RRID:SCR_012095)

Resource Information

URL: <https://code.google.com/p/netcoffee/>

Proper Citation: NetCoffee (RRID:SCR_012095)

Description: A fast and accurate algorithm which allows to find a global alignment of multiple protein-protein interaction networks.

Resource Type: software resource

Defining Citation: [PMID:24336806](#)

Keywords: standalone software, bio.tools

Funding:

Availability: GNU General Public License

Resource Name: NetCoffee

Resource ID: SCR_012095

Alternate IDs: biotools:netcoffee, OMICS_05172

Alternate URLs: <https://bio.tools/netcoffee>

Record Creation Time: 20220129T080308+0000

Record Last Update: 20250420T014606+0000

Ratings and Alerts

No rating or validation information has been found for NetCoffee.

No alerts have been found for NetCoffee.

Data and Source Information

Source: [SciCrunch Registry](#)

Usage and Citation Metrics

We found 3 mentions in open access literature.

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

Chen J, et al. (2024) Comparative transcriptomic analysis provides insights into the genetic networks regulating oil differential production in oil crops. BMC biology, 22(1), 110.

Milano M, et al. (2022) Challenges and Limitations of Biological Network Analysis. Biotech (Basel (Switzerland)), 11(3).

Gu S, et al. (2021) Data-driven biological network alignment that uses topological, sequence, and functional information. BMC bioinformatics, 22(1), 34.