

Resource Summary Report

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NJplot

RRID:SCR_024137

Type: Tool

Proper Citation

NJplot (RRID:SCR_024137)

Resource Information

URL: <https://doua.prabi.fr/software/njplot>

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Description: Software tool as tree drawing program to draw any phylogenetic tree expressed in Newick phylogenetic tree format (e.g., the format used by the PHYLIP package). Used for rooting the unrooted trees obtained from parsimony, distance or maximum likelihood tree-building methods.

Synonyms: njplot

Resource Type: software library, software toolkit, software resource

Keywords: tree drawing program, draw any phylogenetic tree, Newick phylogenetic tree format,

Availability: Free, Available for download, Freely available,

Resource Name: NJplot

Resource ID: SCR_024137

Alternate IDs: OMICS_04815

Alternate URLs: <https://sources.debian.org/src/njplot/>

Record Creation Time: 20230824T050212+0000

Record Last Update: 20260807T043016+0000

Ratings and Alerts

No rating or validation information has been found for NJplot.

No alerts have been found for NJplot.

Data and Source Information

Source: [SciCrunch Registry](#)

Usage and Citation Metrics

We found 2 mentions in open access literature.

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

Fujita T, et al. (2024) SoxC and MmpReg promote blastema formation in whole-body regeneration of fragmenting potworms *Enchytraeus japonensis*. *Nature communications*, 15(1), 6659.

Shulman LM, et al. (2022) Epidemiology of Type 3 Poliovirus AFP Cases in Israel between 1973 and 1988: Whole Genome Sequencing of RNA Extracted Directly from Archived Stocks to Avoid Re-Culturing Neurovirulent Wild Poliovirus. *Vaccines*, 10(12).