

Resource Summary Report

Generated by [Kravitz](#) on Sep 10, 2026

Cybase

RRID:SCR_012925

Type: Tool

Proper Citation

Cybase (RRID:SCR_012925)

Resource Information

URL: <http://www.cybase.org.au/>

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Description: Cybase is dedicated to the study of a fascinating new class of proteins that possess a cyclic backbone in which the N and C termini have been joined with a conventional amide bond. These recently characterized molecules have now been found in organisms from all kingdoms of life and given the current rate of discovery the number of sequences could soon number in the hundreds. Research in our lab is aimed at further characterizing cyclic proteins and adapting them for commercial and medicinal use. In particular we work on a class of cyclic protein named the cyclotides. These proteins are found in the plants of the Rubiaceae and Violaceae and our specific goals include: determining the role that cyclotides play in plants, discovering the mechanism of action of the wide range of biological activities displayed by the cyclotides (including anti-HIV, anti-bacterial and insecticidal activity), characterising the genetics of the cyclotides and further discovery of novel cyclotides.

Synonyms: Cybase

Resource Type: data or information resource, database

Keywords: cyclic protein

Resource Name: Cybase

Resource ID: SCR_012925

Alternate IDs: nif-0000-02710

Old URLs: <http://research1t.imb.uq.edu.au/cybase/>

Record Creation Time: 20220129T080313+0000

Record Last Update: 20260905T133203+0000

Ratings and Alerts

No rating or validation information has been found for Cybase.

No alerts have been found for Cybase.

Data and Source Information

Source: [SciCrunch Registry](#)

Usage and Citation Metrics

We found 38 mentions in open access literature.

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

Hellinger R, et al. (2026) Tropical psychotria plants are a rich source for peptide inhibitors of human prolyl oligopeptidase. *Natural products and bioprospecting*, 16(1).

Sankar S, et al. (2026) Computational prediction of cyclotides from *Viola odorata* as potential inhibitors against the neuraminidase of *Streptococcus pneumoniae*. *Molecular diversity*, 30(1), 991.

Karaca Ocak Ö, et al. (2026) Computational screening of AI-derived cyclotides as putative VEGFR2 binders for wound-site angiogenesis. *Scientific reports*, 16(1).

Deegala S, et al. (2025) Transgenic Innovation: Harnessing Cyclotides as Next Generation Pesticides. *ACS omega*, 10(7), 6323.

Castillo-Mendieta K, et al. (2024) Peptide hemolytic activity analysis using visual data mining of similarity-based complex networks. *NPJ systems biology and applications*, 10(1), 115.

Badaczewska-Dawid A, et al. (2024) Structure prediction of linear and cyclic peptides using CABS-flex. *Briefings in bioinformatics*, 25(2).

Tomašević N, et al. (2024) Discovery and development of macrocyclic peptide modulators of the cannabinoid 2 receptor. *The Journal of biological chemistry*, 300(6), 107330.

Taghizadeh MS, et al. (2024) Unveiling the insecticidal efficiency of *Viola ignobilis* against *Macrosiphum rosae* and *Agonoscyta pistaciae*: From chemical composition to cytotoxicity analysis. *Heliyon*, 10(23), e40636.

Sychta K, et al. (2024) The involvement of cyclotides in the heavy metal tolerance of *Viola* spp. *Scientific reports*, 14(1), 19306.

Lian Y, et al. (2024) Characterization and evaluation of cytotoxic and antimicrobial activities of cyclotides from *Viola japonica*. *Scientific reports*, 14(1), 9733.

Hussain M, et al. (2024) Computational modeling of cyclotides as antimicrobial agents against *Neisseria gonorrhoeae* PorB porin protein: integration of docking, immune, and

molecular dynamics simulations. *Frontiers in chemistry*, 12, 1493165.

Karami Y, et al. (2023) Exploring a Structural Data Mining Approach to Design Linkers for Head-to-Tail Peptide Cyclization. *Journal of chemical information and modeling*, 63(20), 6436.

Rajendran S, et al. (2023) Screening for Cyclotides in Sri Lankan Medicinal Plants: Discovery, Characterization, and Bioactivity Screening of Cyclotides from *Geophila repens*. *Journal of natural products*, 86(1), 52.

Ramazi S, et al. (2022) A review on antimicrobial peptides databases and the computational tools. *Database : the journal of biological databases and curation*, 2022.

Du Q, et al. (2022) Mutagenesis of bracelet cyclotide hyen D reveals functionally and structurally critical residues for membrane binding and cytotoxicity. *The Journal of biological chemistry*, 298(4), 101822.

Slazak B, et al. (2022) The Influence of Plant Stress Hormones and Biotic Elicitors on Cyclotide Production in *Viola uliginosa* Cell Suspension Cultures. *Plants (Basel, Switzerland)*, 11(14).

Huang YH, et al. (2022) Protocols for measuring the stability and cytotoxicity of cyclotides. *Methods in enzymology*, 663, 19.

Pinto MEF, et al. (2021) Cyclotides from Brazilian *Palicourea sessilis* and Their Effects on Human Lymphocytes. *Journal of natural products*, 84(1), 81.

Manniello MD, et al. (2021) Insect antimicrobial peptides: potential weapons to counteract the antibiotic resistance. *Cellular and molecular life sciences : CMLS*, 78(9), 4259.

Slazak B, et al. (2021) Cyclotide host-defense tailored for species and environments in violets from the Canary Islands. *Scientific reports*, 11(1), 12452.