

Resource Summary Report

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CUSP

RRID:SCR_006373

Type: Tool

Proper Citation

CUSP (RRID:SCR_006373)

Resource Information

URL: <http://irvinginstitute.columbia.edu/cusp/cgi-bin/ww2ui.cgi/>

Proper Citation: CUSP (RRID:SCR_006373)

Description: Web-based, open access scientific networking system providing one-stop shopping for investigators seeking collaborators at Columbia University Medical Center in New York City. Contact information, grants, and publications are integrated to facilitate searches by topic or person.

Abbreviations: CUSP

Synonyms: Columbia University Scientific Profiles

Resource Type: community building portal, data or information resource, database, people resource, portal

Keywords: networking system, collaborator, grant, publication, network, organization, FASEB list

Availability: Open access

Resource Name: CUSP

Resource ID: SCR_006373

Alternate IDs: nlx_152161

Record Creation Time: 20220129T080235+0000

Record Last Update: 20260829T182236+0000

Ratings and Alerts

No rating or validation information has been found for CUSP.

No alerts have been found for CUSP.

Data and Source Information

Source: [SciCrunch Registry](#)

Usage and Citation Metrics

We found 58 mentions in open access literature.

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

Wang Y, et al. (2026) Dissecting the evolutionary forces shaping codon usage bias in SFTSV from eastern China: Insights from a previously underrepresented region. *Virulence*, 17(1), 2621490.

Jia S, et al. (2026) Multipartite mitochondrial genome evolution in *Halophila* seagrasses: repeat-driven structural plasticity and selection signatures of marine adaptation. *BMC plant biology*, 26(1).

Sha S, et al. (2026) Functional characterization of BbCHI3 in regulating flavonoid biosynthesis and abiotic stress tolerance in *Blumea balsamifera*. *BMC plant biology*, 26(1).

Li R, et al. (2025) Characterization of the complete chloroplast genome and comparative analysis of the phylogeny and codon usage bias of three Yunnan wild rice species. *Frontiers in plant science*, 16, 1555104.

Li D, et al. (2025) Assembly and comparative analysis of the first complete mitochondrial genome of *Camellia sinensis* var. *assamica* 'Hainan Dayezhong', endemic to Hainan Province, China. *BMC plant biology*, 25(1), 1304.

Mo ZM, et al. (2025) Chloroplast genome analysis and phylogenetic position of *Polygonatum sinopubescens* and comparison with related species. *PloS one*, 20(12), e0338103.

Mudiyanselage SD, et al. (2025) A global high resolution coastline database from satellite imagery. *Scientific data*, 12(1), 812.

Zhu W, et al. (2025) Comparative genomics and phylogenetic analysis of mitochondrial genomes of *Neocinnamomum*. *BMC plant biology*, 25(1), 289.

Cheng X, et al. (2024) Unveiling the conserved nature of *Heliconia* chloroplast genomes: insights from the assembly and analysis of four complete chloroplast genomes. *Frontiers in plant science*, 15, 1535549.

Farzanegan F, et al. (2024) Evaluating malocclusion patterns in children with autism spectrum disorder using the index of complexity, outcome and need: a cross-sectional study. *BMC oral health*, 24(1), 759.

Wang L, et al. (2024) Comparative analysis of the mitochondrial genomes of four *Dendrobium* species (Orchidaceae) reveals heterogeneity in structure, synteny, intercellular gene transfer, and RNA editing. *Frontiers in plant science*, 15, 1429545.

Huang YB, et al. (2024) *Salvia guidongensis* sp. nov.: unraveling a critical evolutionary link in East Asian *Salvia* from Central China integrating morphology, phylogeny, and plastid genomics. *Frontiers in plant science*, 15, 1332443.

Latif A, et al. (2024) Implementation and long-term efficacy of a multifaceted intervention to reduce central line-associated bloodstream infections in intensive care units of a low-middle-income country. *American journal of infection control*, 52(7), 819.

Elawsya ME, et al. (2024) Two-year clinical performance of dual- and light-cure bulk-fill resin composites in Class ?? restorations: a randomized clinical trial. *Clinical oral investigations*, 28(2), 138.

Ye J, et al. (2024) Analysis of chloroplast genome structure and phylogeny of the traditional medicinal of *Ardisia crispa* (Myrsinaceae). *Scientific reports*, 14(1), 19045.

Shi N, et al. (2024) Analysis of codon usage patterns in complete plastomes of four medicinal *Polygonatum* species (Asparagaceae). *Frontiers in genetics*, 15, 1401013.

Sari NP, et al. (2023) Is the association between mothers' autistic traits and childhood autistic traits moderated by maternal pre-pregnancy body mass index? *Molecular autism*, 14(1), 46.

Hu H, et al. (2023) Insights into the phylogenetic relationships and species boundaries of the *Myricaria squamosa* complex (Tamaricaceae) based on the complete chloroplast genome. *PeerJ*, 11, e16642.

Wang Y, et al. (2023) Comparative analysis of codon usage patterns in chloroplast genomes of ten *Epimedium* species. *BMC genomic data*, 24(1), 3.

Liu A, et al. (2023) Application of comprehensive unit-based safety program model to improve chemotherapy-induced nausea and vomiting in patients with ovarian cancer: a retrospective study. *Journal of ovarian research*, 16(1), 143.