

Resource Summary Report

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Chado

RRID:SCR_024073

Type: Tool

Proper Citation

Chado (RRID:SCR_024073)

Resource Information

URL: <http://gmod.org/wiki/Chado>

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Description: Relational database schema that underlies many GMOD installations. It is capable of representing many of the general classes of data frequently encountered in modern biology such as sequence, sequence comparisons, phenotypes, genotypes, ontologies, publications, and phylogeny. It has been designed to handle complex representations of biological knowledge and should be considered one of the most sophisticated relational schemas currently available in molecular biology. The price of this capability is that the new user must spend some time becoming familiar with its fundamentals.

Synonyms: libchado-perl

Resource Type: software application, software resource, data management software

Keywords: database schema, GMOD installations,

Availability: Free, Available for download, Freely available,

Resource Name: Chado

Resource ID: SCR_024073

Alternate URLs: <https://sources.debian.org/src/libchado-perl/>

License: GNU Free Documentation License

Record Creation Time: 20230824T050211+0000

Record Last Update: 20260729T044603+0000

Ratings and Alerts

No rating or validation information has been found for Chado.

No alerts have been found for Chado.

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 [kravitz2](#) .

Mansueto L, et al. (2024) Building a community-driven bioinformatics platform to facilitate Cannabis sativa multi-omics research. GigaByte (Hong Kong, China), 2024, gigabyte137.