Resource Summary Report

Generated by NIF on Apr 9, 2025

STRENDA

RRID:SCR_017422
Type: Tool

Proper Citation

STRENDA (RRID:SCR_017422)

Resource Information

URL: https://www.beilstein-strenda-db.org/strenda/

Proper Citation: STRENDA (RRID:SCR_017422)

Description: Storage and search platform supported by Beilstein-Institut that incorporates STRENDA Guidelines. For authors who prepare manuscript containing functional enzymology data, STRENDA DB provides means to ensure that data sets are complete and valid before submitting them to journal.

Abbreviations: STRENDA

Synonyms: , Standards for Reporting Enzymology Data, Beilstein-Institut, STRENDA

Resource Type: database, data or information resource, service resource, storage service resource, data repository

Keywords: Storage, Beilstein Institut, functional, enzymology, data, complition, validation, dataset, guideline, standard, reporting

Funding:

Availability: Restricted

Resource Name: STRENDA

Resource ID: SCR_017422

Alternate IDs: DOI:10.22011, DOI:10.17616/R3536N, DOI:10.25504/FAIRsharing.ekj9zx

Alternate URLs: http://www.strenda-db.org/, https://doi.org/10.17616/R3536N,

https://doi.org/10.17616/r3536n, https://doi.org/10.22011/, https://dx.doi.org/10.22011/, https://fairsharing.org/10.25504/FAIRsharing.ekj9zx

Record Creation Time: 20220129T080335+0000

Record Last Update: 20250409T061507+0000

Ratings and Alerts

No rating or validation information has been found for STRENDA .

No alerts have been found for STRENDA .

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 NIF.

Washington EJ, et al. (2024) Structures of trehalose-6-phosphate synthase, Tps1, from the fungal pathogen Cryptococcus neoformans: A target for antifungals. Proceedings of the National Academy of Sciences of the United States of America, 121(32), e2314087121.

McDonald AG, et al. (2022) Parameter Reliability and Understanding Enzyme Function. Molecules (Basel, Switzerland), 27(1).

Pinto MF, et al. (2021) interferENZY: A Web-Based Tool for Enzymatic Assay Validation and Standardized Kinetic Analysis. Journal of molecular biology, 433(11), 166613.