

Resource Summary Report

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STRENDA

RRID:SCR_017422

Type: Tool

Proper Citation

STREND A (RRID:SCR_017422)

Resource Information

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

Proper Citation: STREND A (RRID:SCR_017422)

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

Abbreviations: STREND A

Synonyms: , Standards for Reporting Enzymology Data, Beilstein-Institut, STREND A

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

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

Availability: Restricted

Resource Name: STREND A

Resource ID: SCR_017422

Alternate IDs: DOI:10.22011, DOI:10.17616/R3536N, DOI:10.25504/FAIRsharing.ekj9zx, r3d100012329

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>, <https://doi.org/10.17616/R3536N>

Record Creation Time: 20220129T080335+0000

Record Last Update: 20260729T044435+0000

Ratings and Alerts

No rating or validation information has been found for STREND A .

No alerts have been found for STREND A .

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 [SPARC SAWG](#) .

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.