

Resource Summary Report

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Lhasa Limited

RRID:SCR_003775

Type: Tool

Proper Citation

Lhasa Limited (RRID:SCR_003775)

Resource Information

URL: <http://www.lhasalimited.org/>

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Description: A not-for-profit membership organization and educational charity that facilitates collaborative data sharing projects in the pharmaceutical, cosmetics and chemistry-related industries specializing in the development of expert computer systems for toxicity and metabolism prediction. They provide a number of extensive and continually updated knowledge bases and the software needed to interrogate them. Its charitable aims include the sponsorship of activities that advance scientific knowledge and understanding and they regularly support computational chemistry events and initiatives that are of interest to Lhasa Limited members and the wider scientific community. All applications for sponsorship will be considered on their individual merits.

Abbreviations: Lhasa

Synonyms: Lhasa Ltd.

Resource Type: software resource, funding resource, data or information resource, portal, community building portal

Keywords: data sharing, pharmaceutical, cosmetic, chemistry, toxicity prediction, metabolism, toxicity, metabolism prediction, prediction, database, in silico, computational chemistry, computer system

Availability: Membership required, For sale

Resource Name: Lhasa Limited

Resource ID: SCR_003775

Alternate IDs: nlx_158058

Record Creation Time: 20220129T080220+0000

Ratings and Alerts

No rating or validation information has been found for Lhasa Limited.

No alerts have been found for Lhasa Limited.

Data and Source Information

Source: [SciCrunch Registry](#)

Usage and Citation Metrics

We found 12 mentions in open access literature.

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

Gopalswamy M, et al. (2025) Identification of non-charged 7.44 analogs interacting with the NHR2 domain of RUNX1-ETO with improved antiproliferative effect in RUNX-ETO positive cells. Scientific reports, 15(1), 17720.

Sviben M, et al. (2025) Resveratrol-Based Carbamates as Selective Butyrylcholinesterase Inhibitors: Design, Synthesis, Computational Study and Biometal Complexation Capability. Molecules (Basel, Switzerland), 30(2).

Dablander M, et al. (2024) Sort & Slice: a simple and superior alternative to hash-based folding for extended-connectivity fingerprints. Journal of cheminformatics, 16(1), 135.

Graham JC, et al. (2022) An Evaluation of the Occupational Health Hazards of Peptide Couplers. Chemical research in toxicology, 35(6), 1011.

Alov P, et al. (2022) In Silico Identification of Multi-Target Ligands as Promising Hit Compounds for Neurodegenerative Diseases Drug Development. International journal of molecular sciences, 23(21).

, et al. (2021) Scientific Guidance for the preparation of applications on smoke flavouring primary products. EFSA journal. European Food Safety Authority, 19(3), e06435.

Rim KT, et al. (2020) In silico prediction of toxicity and its applications for chemicals at work. Toxicology and environmental health sciences, 12(3), 191.

Kar S, et al. (2018) Impact of Pharmaceuticals on the Environment: Risk Assessment Using QSAR Modeling Approach. Methods in molecular biology (Clifton, N.J.), 1800, 395.

Dos Santos CBR, et al. (2018) Oil from the fruits of Pterodon emarginatus Vog.: A traditional anti-inflammatory. Study combining in vivo and in silico. Journal of ethnopharmacology, 222, 107.

Clark RD, et al. (2018) Predicting mammalian metabolism and toxicity of pesticides in silico. Pest management science, 74(9), 1992.

Hsin KY, et al. (2011) EDULISS: a small-molecule database with data-mining and pharmacophore searching capabilities. Nucleic acids research, 39(Database issue), D1042.

Fitzsimmons PN, et al. (2007) A compilation of in vitro rate and affinity values for xenobiotic biotransformation in fish, measured under physiological conditions. Comparative biochemistry and physiology. Toxicology & pharmacology : CBP, 145(4), 485.