

Resource Summary Report

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Glide

RRID:SCR_000187

Type: Tool

Proper Citation

Glide (RRID:SCR_000187)

Resource Information

URL: <https://www.schrodinger.com/glide>

Proper Citation: Glide (RRID:SCR_000187)

Description: Software package which approximates a complete search of the conformational, orientational, and positional space of the ligand in a given receptor. Used in drug development for predicting protein ligand binding modes and ranking ligands via high throughput virtual screening.

Abbreviations: Glide

Resource Type: simulation software, software resource, software application

Defining Citation: [PMID:18428795](#)

Keywords: ligand, receptor, docking, computation, virtual, screening, drug, discovery

Availability: Restricted

Resource Name: Glide

Resource ID: SCR_000187

Alternate IDs: OMICS_01601

Record Creation Time: 20220129T080200+0000

Record Last Update: 20260819T044009+0000

Ratings and Alerts

No rating or validation information has been found for Glide.

No alerts have been found for Glide.

Data and Source Information

Source: [SciCrunch Registry](#)

Usage and Citation Metrics

We found 17 mentions in open access literature.

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

Feher A, et al. (2026) A kinetic trap mechanism underlies hHv1 inhibition by NZ-58. The Journal of general physiology, 158(4).

Pyun WY, et al. (2026) Nutrient Availability Dictates Cancer Metabolism-Based Therapeutic Responses to Nononcology Drugs. Cancer research, 86(5), 1194.

Motso A, et al. (2025) GRK-biased adrenergic agonists for the treatment of type 2 diabetes and obesity. Cell.

Puleo N, et al. (2025) Identification of a TNIK-CDK9 Axis as a Targetable Strategy for Platinum-Resistant Ovarian Cancer. Molecular cancer therapeutics, 24(4), 639.

Pedersen CN, et al. (2024) Cryo-EM structure of the dopamine transporter with a novel atypical non-competitive inhibitor bound to the orthosteric site. Journal of neurochemistry, 168(9), 2043.

Xu G, et al. (2024) Proteomic Profiling of Serum Extracellular Vesicles Identifies Diagnostic Signatures and Therapeutic Targets in Breast Cancer. Cancer research, 84(19), 3267.

Brown E, et al. (2024) Inhibitors of the small membrane (M) protein viroporin prevent Zika virus infection. eLife, 13.

Schneider C, et al. (2024) A Novel AMPK Inhibitor Sensitizes Pancreatic Cancer Cells to Ferroptosis Induction. Advanced science (Weinheim, Baden-Wurttemberg, Germany), 11(31), e2307695.

Masone A, et al. (2023) A tetracationic porphyrin with dual anti-prion activity. iScience, 26(9), 107480.

Lettl C, et al. (2023) Selective killing of the human gastric pathogen Helicobacter pylori by mitochondrial respiratory complex I inhibitors. Cell chemical biology, 30(5), 499.

Wang Y, et al. (2022) Scutellarein attenuates atopic dermatitis by selectively inhibiting transient receptor potential vanilloid 3 channels. British journal of pharmacology, 179(20), 4792.

Guo X, et al. (2021) Structure and mechanism of a phage-encoded SAM lyase revises catalytic function of enzyme family. eLife, 10.

Palomares B, et al. (2020) Δ^9 -Tetrahydrocannabinolic acid alleviates collagen-induced arthritis: Role of PPAR γ and CB1 receptors. *British journal of pharmacology*, 177(17), 4034.

Reyna DE, et al. (2017) Direct Activation of BAX by BTSA1 Overcomes Apoptosis Resistance in Acute Myeloid Leukemia. *Cancer cell*, 32(4), 490.

Min DK, et al. (2015) In silico Screening of Chemical Libraries to Develop Inhibitors That Hamper the Interaction of PCSK9 with the LDL Receptor. *Yonsei medical journal*, 56(5), 1251.

Borhade SR, et al. (2014) Inhibition of Insulin-Regulated Amino peptidase (IRAP) by Arylsulfonamides. *ChemistryOpen*, 3(6), 256.

Repasky MP, et al. (2007) Flexible ligand docking with Glide. *Current protocols in bioinformatics*, Chapter 8.