

Resource Summary Report

Generated by [PRECISE-TBI](#) on Aug 2, 2026

PASSer

RRID:SCR_027790

Type: Tool

Proper Citation

PASSer (RRID:SCR_027790)

Resource Information

URL: <https://passer.smu.edu/>

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Description: Web allosteric site prediction tool. Web server designed for prediction and visualization of protein allosteric sites, which are crucial locations on proteins for regulating their function and key targets for drug development. Helps researchers identify these regulatory sites on proteins, supporting efforts in allosteric drug discovery by analyzing protein structures from Protein Data Bank.

Synonyms: Protein Allosteric Sites Server

Resource Type: software resource, data access protocol, web application, web service

Keywords: allosteric site prediction, identify regulatory sites, identify regulatory sites on proteins, proteins, allosteric drug discovery, analyzing protein structures,

Availability: Free, Freely available

Resource Name: PASSer

Resource ID: SCR_027790

Record Creation Time: 20251216T055148+0000

Record Last Update: 20260801T191433+0000

Ratings and Alerts

No rating or validation information has been found for PASSer.

No alerts have been found for PASSer.

Data and Source Information

Source: [SciCrunch Registry](#)

Usage and Citation Metrics

We have not found any literature mentions for this resource.