WSsas - Web Service for the SAS tool

RRID:SCR_007051
Type: Tool

Proper Citation

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Resource Information

URL: http://www.ebi.ac.uk/thornton-srv/databases/WSsas/

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Description: SAS is a tool for applying structural information to a given protein sequence. It uses FASTA to scan a given protein sequence against all the proteins of known 3D structure in the Protein Data Bank and provides functional residue annotation based on data from the Catalytic Site Atlas and PDBsum. The web service is aimed to facilitate the use of the SAS tool when having a huge number of queries. Currently, the web service provides annotation for binding sites (to ligand, metal or nucleic acid), catalytic residues and amino acids related to protein-protein interactions.

Abbreviations: WSsas

Synonyms: Web Service for the SAS tool

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

Keywords: gold standard, bio.tools

Resource Name: WSsas - Web Service for the SAS tool

Resource ID: SCR_007051

Alternate IDs: nlx_18182, biotools:wssas

Alternate URLs: https://bio.tools/wssas

Ratings and Alerts
No rating or validation information has been found for WSsas - Web Service for the SAS tool.

No alerts have been found for WSsas - Web Service for the SAS tool.

Data and Source Information

**Source:** SciCrunch Registry

Usage and Citation Metrics

We have not found any literature mentions for this resource.