Resource Summary Report

Generated by RRID on Sep 19, 2026

NGlycPred

RRID:SCR_016610
Type: Tool

Proper Citation

NGlycPred (RRID:SCR_016610)

Resource Information

URL: https://bioinformatics.niaid.nih.gov/nglycpred/

Proper Citation: NGlycPred (RRID:SCR_016610)

Description: Web based server to discriminate between glycosylated and non-glycosylated N-X-(T/S) sequons by incorporating structural information and patterns.

Abbreviations: NGlycPred

Synonyms: N linked Glycosylation Prediction server, N linked GLYCosylation PREDiction server

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

Defining Citation: DOI:10.1093/bioinformatics/bts426

Keywords: predict, classify, glycosylated, nonglycosylate, asparagine, sequons, structure, pattern, information

Funding: Intramural Research Program NIAID

Availability: Free, Freely available

Resource Name: NGlycPred

Resource ID: SCR_016610

Record Creation Time: 20220129T080331+0000

Record Last Update: 20260912T195842+0000

Ratings and Alerts

No rating or validation information has been found for NGlycPred.

No alerts have been found for NGlycPred.

Data and Source Information

Source: SciCrunch Registry

Usage and Citation Metrics

We found 1 mentions in open access literature.

Listed below are recent publications. The full list is available at RRID .

Cereghino C, et al. (2023) The E2 glycoprotein holds key residues for Mayaro virus adaptation to the urban Aedes aegypti mosquito. PLoS pathogens, 19(4), e1010491.