

Resource Summary Report

Generated by [RRID](#) on Aug 13, 2026

ExonScan Web Server

RRID:SCR_022516

Type: Tool

Proper Citation

ExonScan Web Server (RRID:SCR_022516)

Resource Information

URL: <http://hollywood.mit.edu/exonscan/>

Proper Citation: ExonScan Web Server (RRID:SCR_022516)

Description: Software framework for modeling sequence motifs based on maximum entropy principle.

Synonyms: ExonScan

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

Defining Citation: [PMID:15285897](#)

Keywords: modeling sequence motifs, maximum entropy principle

Availability: Free, Freely available

Resource Name: ExonScan Web Server

Resource ID: SCR_022516

Record Creation Time: 20220625T050140+0000

Record Last Update: 20260813T055233+0000

Ratings and Alerts

No rating or validation information has been found for ExonScan Web Server.

No alerts have been found for ExonScan Web Server.

Data and Source Information

Source: [SciCrunch Registry](#)

Usage and Citation Metrics

We found 2 mentions in open access literature.

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

Lin BC, et al. (2023) In silico methods for predicting functional synonymous variants. Genome biology, 24(1), 126.

Holcomb DD, et al. (2022) Protocol to identify host-viral protein interactions between coagulation-related proteins and their genetic variants with SARS-CoV-2 proteins. STAR protocols, 3(3), 101648.