**PredictSNP**

**RRID:** SCR_006327  
**Type:** Tool

**Proper Citation**

PredictSNP (RRID:SCR_006327)

**Resource Information**

**URL:** http://loschmidt.chemi.muni.cz/predictsnp/

**Proper Citation:** PredictSNP (RRID:SCR_006327)

**Description:** Consensus classifier tool that combines six of the top performing tools for the prediction of the effects of mutation on protein function. The obtained results are provided together with annotations extracted from the Protein Mutant Database and the UniProt database. A stand-alone version is also available.

**Abbreviations:** PredictSNP

**Synonyms:** PredictSNP - Consensus classifier for prediction of disease-related mutations

**Resource Type:** software resource, analysis service resource, service resource, production service resource, data analysis service

**Defining Citation:** PMID:24453961

**Keywords:** single nucleotide polymorphism, classifier, prediction, mutation, protein function, FASEB list

**Availability:** Free for academic use

**Resource Name:** PredictSNP

**Resource ID:** SCR_006327

**Alternate IDs:** OMICS_02218

Ratings and Alerts
No rating or validation information has been found for PredictSNP.

No alerts have been found for PredictSNP.

Data and Source Information

Source: SciCrunch Registry

Usage and Citation Metrics

We found 101 mentions in open access literature.

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


Kumar S, et al. (2022) Omicron (BA.1) and sub-variants (BA.1.1, BA.2, and BA.3) of SARS-CoV-2 spike infectivity and pathogenicity: A comparative sequence and structural-based...

Singh AK, et al. (2022) Contrasting Distribution of SARS-CoV-2 Lineages across Multiple Rounds of Pandemic Waves in West Bengal, the Gateway of East and North-East States of India. Microbiology spectrum, 10(4), e0091422.


Peres KC, et al. (2021) Clinical utility of TGFB1 and its receptors (TGFR1 and TGFR2) in thyroid nodules: evaluation based on single nucleotide polymorphisms and mRNA analysis. Archives of endocrinology and metabolism, 65(2), 172.


Bug DS, et al. (2021) Towards Understanding the Pathogenicity of DROSHA Mutations in Oncohematology. Cells, 10(9).