# **Resource Summary Report**

Generated by dkNET on May 24, 2025

# **HSLPred**

RRID:SCR\_011972

Type: Tool

### **Proper Citation**

HSLPred (RRID:SCR\_011972)

#### Resource Information

URL: http://www.imtech.res.in/raghava/hslpred/

**Proper Citation:** HSLPred (RRID:SCR\_011972)

**Description:** A support vector machine (SVM)-based method for the prediction of 4 major subcellular localization (cytoplasm, mitochondrial, nuclear and plasma membrane) of human proteins using various features such as i) amino acid composition, ii) dipeptide composition and iii) evolutionary information of proteins.

Abbreviations: HSLPred

**Synonyms:** HSLPred - A SVM-based Method for Subcellular Localization of Human

**Proteins** 

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

analysis service resource

**Defining Citation: PMID:15647269** 

**Keywords:** subcellular localization, protein, support vector machine, bio.tools

**Funding:** 

Availability: Acknowledgement requested

Resource Name: HSLPred

Resource ID: SCR\_011972

Alternate IDs: biotools:hslpred, OMICS\_01622

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

**Record Creation Time:** 20220129T080307+0000

Record Last Update: 20250524T060425+0000

## Ratings and Alerts

No rating or validation information has been found for HSLPred.

No alerts have been found for HSLPred.

#### **Data and Source Information**

Source: SciCrunch Registry

## **Usage and Citation Metrics**

We have not found any literature mentions for this resource.