Resource Summary Report

Generated by RRID on Apr 17, 2025

HMMvar

RRID:SCR_001177 Type: Tool

Proper Citation

HMMvar (RRID:SCR_001177)

Resource Information

URL: https://bioinformatics.cs.vt.edu/zhanglab/hmm/

Proper Citation: HMMvar (RRID:SCR_001177)

Description: Software applying a quantitative prediction method to predict the effect of genetic variation using hidden Markov models.

Abbreviations: HMMvar

Resource Type: software resource

Defining Citation: PMID:24405700

Keywords: genetic variation, insertion, deletion, indel

Funding:

Availability: Free, Public

Resource Name: HMMvar

Resource ID: SCR_001177

Alternate IDs: OMICS_02181

Record Creation Time: 20220129T080206+0000

Record Last Update: 20250410T064652+0000

Ratings and Alerts

No rating or validation information has been found for HMMvar.

No alerts have been found for HMMvar.

Data and Source Information

Source: SciCrunch Registry

Usage and Citation Metrics

We have not found any literature mentions for this resource.