

Resource Summary Report

Generated by [RRID](#) on Aug 15, 2026

[mhcflurry](#)

RRID:SCR_028577

Type: Tool

Proper Citation

mhcflurry (RRID:SCR_028577)

Resource Information

URL: <https://github.com/openvax/mhcflurry>

Proper Citation: mhcflurry (RRID:SCR_028577)

Description: Software MHC I ligand prediction package with competitive accuracy and documented implementation. Used for peptide-MHC I binding affinity prediction.

Resource Type: software toolkit, source code, software resource

Keywords: MHC I ligand prediction, peptide-MHC I binding affinity prediction, peptide-MHC I, binding affinity prediction,

Availability: Free, Available for download, Freely available

Resource Name: mhcflurry

Resource ID: SCR_028577

License: Apache-2.0 license

Record Creation Time: 20260619T043006+0000

Record Last Update: 20260815T183324+0000

Ratings and Alerts

No rating or validation information has been found for mhcflurry.

No alerts have been found for mhcflurry.

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](#) .

Juanes-Velasco P, et al. (2024) Crucial Parameters for Immuno-peptidome Characterization: A Systematic Evaluation. International journal of molecular sciences, 25(17).

Zeng H, et al. (2019) DeepLigand: accurate prediction of MHC class I ligands using peptide embedding. Bioinformatics (Oxford, England), 35(14), i278.