

Resource Summary Report

Generated by [RRID](#) on Aug 11, 2026

EpiJen

RRID:SCR_026689

Type: Tool

Proper Citation

EpiJen (RRID:SCR_026689)

Resource Information

URL: <https://www.ddg-pharmfac.net/epijen/EpiJen/EpiJen.htm>

Proper Citation: EpiJen (RRID:SCR_026689)

Description: Web server for multistep T cell epitope prediction. Also considers proteasome cleavage and TAP binding, therefore it is able to mimic the MHC binding mechanism in real life. At present only HLA class I alleles are included in the prediction.

Resource Type: web application, software resource

Defining Citation: [PMID:16533401](#)

Keywords: multistep T cell epitope prediction, epitope prediction, proteasome cleavage, TAP binding, HLA class I alleles,

Funding: GlaxoSmithKline ;
UK Department of Health

Availability: Free, Freely available

Resource Name: EpiJen

Resource ID: SCR_026689

Record Creation Time: 20250401T060330+0000

Record Last Update: 20260808T190751+0000

Ratings and Alerts

No rating or validation information has been found for EpiJen.

No alerts have been found for EpiJen.

Data and Source Information

Source: [SciCrunch Registry](#)

Usage and Citation Metrics

We found 6 mentions in open access literature.

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

Fathi S, et al. (2026) Systems Vaccinology-Integrated Proteomics to Develop a Novel Prophylactic and Therapeutic Vaccine Against Human Papillomavirus 16. *Bioinformatics and biology insights*, 20, 11779322261437247.

Lara-Ramirez D, et al. (2025) In silico design of Ebola virus Glycoprotein antigenic peptides as vaccine candidates. *PloS one*, 20(3), e0319496.

Wei Y, et al. (2025) Application of Humanized MHC Transgenic Mice in the Screening of HLA-Restricted T Cell Epitopes for Influenza Vaccines. *Vaccines*, 13(3).

Shafiee F, et al. (2016) Expression and purification of truncated diphtheria toxin, DT386, in *Escherichia coli*: An attempt for production of a new vaccine against diphtheria. *Research in pharmaceutical sciences*, 11(5), 428.

Idrees S, et al. (2013) HCV Envelope protein 2 sequence comparison of Pakistani isolate and In-silico prediction of conserved epitopes for vaccine development. *Journal of translational medicine*, 11, 105.

Seyed N, et al. (2011) In silico analysis of six known *Leishmania* major antigens and in vitro evaluation of specific epitopes eliciting HLA-A2 restricted CD8 T cell response. *PLoS neglected tropical diseases*, 5(9), e1295.