

Resource Summary Report

Generated by [dkNET](#) on Sep 4, 2026

JenPep

RRID:SCR_008191

Type: Tool

Proper Citation

JenPep (RRID:SCR_008191)

Resource Information

URL: <http://www.jenner.ac.uk/>

Proper Citation: JenPep (RRID:SCR_008191)

Description: JenPep is a database of quantitative binding data for immunological protein-peptide interactions. It is a system which allows speedy access to this binding data through simple on-line interfaces and effective search mechanisms.

Synonyms: JenPep

Resource Type: data or information resource, database

Resource Name: JenPep

Resource ID: SCR_008191

Alternate IDs: nif-0000-21229

Old URLs: <http://www.jenner.ac.uk/Jenpep/>

Record Creation Time: 20220129T080246+0000

Record Last Update: 20260904T000238+0000

Ratings and Alerts

No rating or validation information has been found for JenPep.

No alerts have been found for JenPep.

Data and Source Information

Source: [SciCrunch Registry](#)

Usage and Citation Metrics

We found 12 mentions in open access literature.

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

Bou Zerdan M, et al. (2021) Mechanisms of Immunotoxicity: Stressors and Evaluators. International journal of molecular sciences, 22(15).

Lawton G, et al. (2020) Vaccines: hope vs reality. New scientist (1971), 248(3309), 8.

Zhang Y, et al. (2018) Compositional and Functional Differences in the Human Gut Microbiome Correlate with Clinical Outcome following Infection with Wild-Type Salmonella enterica Serovar Typhi. mBio, 9(3).

Karkhah A, et al. (2016) A potent multivalent vaccine for modulation of immune system in atherosclerosis: an in silico approach. Clinical and experimental vaccine research, 5(1), 50.

Reeve R, et al. (2016) Tracking the Antigenic Evolution of Foot-and-Mouth Disease Virus. PloS one, 11(7), e0159360.

Alharbi NK, et al. (2015) Investigation of IRES Insertion into the Genome of Recombinant MVA as a Translation Enhancer in the Context of Transcript Decapping. PloS one, 10(5), e0127978.

Alharbi NK, et al. (2015) Deletion of Fifteen Open Reading Frames from Modified Vaccinia Virus Ankara Fails to Improve Immunogenicity. PloS one, 10(6), e0128626.

Biswas S, et al. (2014) Assessment of humoral immune responses to blood-stage malaria antigens following ChAd63-MVA immunization, controlled human malaria infection and natural exposure. PloS one, 9(9), e107903.

Jan M, et al. (2010) Inflammatory and autoimmune reactions in atherosclerosis and vaccine design informatics. Journal of biomedicine & biotechnology, 2010, 459798.

Lundegaard C, et al. (2007) Modeling the adaptive immune system: predictions and simulations. Bioinformatics (Oxford, England), 23(24), 3265.

Toseland CP, et al. (2006) PPD v1.0--an integrated, web-accessible database of experimentally determined protein pKa values. Nucleic acids research, 34(Database issue), D199.

Doytchinova IA, et al. (2003) Proteomics in Vaccinology and Immunobiology: An Informatics Perspective of the Immunone. Journal of biomedicine & biotechnology, 2003(5), 267.