

Resource Summary Report

Generated by [ASWG](#) on Aug 30, 2026

EpiGRAPH

RRID:SCR_004326

Type: Tool

Proper Citation

EpiGRAPH (RRID:SCR_004326)

Resource Information

URL: <http://epigraph.mpi-inf.mpg.de/WebGRAPH/>

Proper Citation: EpiGRAPH (RRID:SCR_004326)

Description: A software for genome and epigenome analysis., THIS RESOURCE IS NO LONGER IN SERVICE. Documented on September 16,2025.

Abbreviations: EpiGRAPH

Resource Type: software resource

Keywords: bio.tools

Availability: THIS RESOURCE IS NO LONGER IN SERVICE

Resource Name: EpiGRAPH

Resource ID: SCR_004326

Alternate IDs: OMICS_00633, biotools:epigraph

Alternate URLs: <https://bio.tools/epigraph>

Record Creation Time: 20220129T080224+0000

Record Last Update: 20260829T182151+0000

Ratings and Alerts

No rating or validation information has been found for EpiGRAPH.

No alerts have been found for EpiGRAPH.

Data and Source Information

Usage and Citation Metrics

We found 18 mentions in open access literature.

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

Techasen A, et al. (2025) Identification and characterization of a target antigen recognized by the monoclonal antibody against *Opisthorchis viverrini*. PloS one, 20(5), e0324137.

Subiza-Pérez M, et al. (2025) Examining Predictors of Environmental Health Knowledge and Competences in a Sample of Catalan Paediatricians. Health science reports, 8(3), e70569.

Petro-Turnquist EM, et al. (2025) Epitope-optimized vaccine elicits enduring immunity against swine influenza A virus. Nature communications, 16(1), 4046.

Petro-Turnquist E, et al. (2024) Multivalent Epigraph Hemagglutinin Vaccine Protects against Influenza B Virus in Mice. Pathogens (Basel, Switzerland), 13(2).

Petro-Turnquist E, et al. (2024) Swine influenza A virus: challenges and novel vaccine strategies. Frontiers in cellular and infection microbiology, 14, 1336013.

Ugwu C, et al. (2022) Humoral and cellular immune responses to Lassa fever virus in Lassa fever survivors and their exposed contacts in Southern Nigeria. Scientific reports, 12(1), 22330.

Garcia-Bates TM, et al. (2021) Dendritic cells focus CTL responses toward highly conserved and topologically important HIV-1 epitopes. EBioMedicine, 63, 103175.

Singh DE, et al. (2021) Simulation of COVID-19 Propagation Scenarios in the Madrid Metropolitan Area. Frontiers in public health, 9, 636023.

Gras A, et al. (2021) Use of suggestive seizure manipulation methods in the investigation of patients with possible psychogenic nonepileptic seizures-An international ILAE survey. Epilepsia open, 6(3), 472.

Guzmán-Merino M, et al. (2021) Assessing population-sampling strategies for reducing the COVID-19 incidence. Computers in biology and medicine, 139, 104938.

Troya MI, et al. (2020) Covid-19 Estimating the burden of symptomatic disease in the community and the impact of public health measures on physical, mental and social wellbeing: a study protocol. HRB open research, 3, 49.

Singh DE, et al. (2020) Evaluating the impact of the weather conditions on the influenza propagation. BMC infectious diseases, 20(1), 265.

Xiao X, et al. (2017) Molecular mechanisms underlying noncoding risk variations in psychiatric genetic studies. Molecular psychiatry, 22(4), 497.

Theiler J, et al. (2016) Epigraph: A Vaccine Design Tool Applied to an HIV Therapeutic Vaccine and a Pan-Filovirus Vaccine. *Scientific reports*, 6, 33987.

Jeffries AR, et al. (2016) Erasure and reestablishment of random allelic expression imbalance after epigenetic reprogramming. *RNA* (New York, N.Y.), 22(10), 1620.

Cao K, et al. (2013) High-resolution mapping of h1 linker histone variants in embryonic stem cells. *PLoS genetics*, 9(4), e1003417.

Edwards SL, et al. (2013) Beyond GWASs: illuminating the dark road from association to function. *American journal of human genetics*, 93(5), 779.

Lim SJ, et al. (2010) Computational Epigenetics: the new scientific paradigm. *Bioinformatics*, 4(7), 331.