

Resource Summary Report

Generated by [PRECISE-TBI](#) on Jul 31, 2026

HMMgene

RRID:SCR_011933

Type: Tool

Proper Citation

HMMgene (RRID:SCR_011933)

Resource Information

URL: <http://www.cbs.dtu.dk/services/HMMgene/>

Proper Citation: HMMgene (RRID:SCR_011933)

Description: Data analysis service for prediction of vertebrate and C. elegans genes.

Abbreviations: HMMgene

Resource Type: service resource, production service resource, analysis service resource, data analysis service

Keywords: gene

Resource Name: HMMgene

Resource ID: SCR_011933

Alternate IDs: OMICS_01488

Record Creation Time: 20220129T080307+0000

Record Last Update: 20260731T042839+0000

Ratings and Alerts

No rating or validation information has been found for HMMgene.

No alerts have been found for HMMgene.

Data and Source Information

Source: [SciCrunch Registry](#)

Usage and Citation Metrics

We found 11 mentions in open access literature.

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

Leijon M, et al. (2025) Characterization of Esocid Herpesvirus 1 (EsHV1) from Europe. *Pathogens* (Basel, Switzerland), 14(1).

Leijon M, et al. (2025) Genome Characterisation of Esocid Herpesvirus 1 (EsHV-1). *Viruses*, 17(10).

Schwartz JC, et al. (2019) The Structure, Evolution, and Gene Expression Within the Caprine Leukocyte Receptor Complex. *Frontiers in immunology*, 10, 2302.

Schwartz JC, et al. (2017) The evolution of the natural killer complex; a comparison between mammals using new high-quality genome assemblies and targeted annotation. *Immunogenetics*, 69(4), 255.

Burns AR, et al. (2017) The novel nematicide wact-86 interacts with aldicarb to kill nematodes. *PLoS neglected tropical diseases*, 11(4), e0005502.

Combi R, et al. (2009) Clinical and genetic familial study of a large cohort of Italian children with idiopathic epilepsy. *Brain research bulletin*, 79(2), 89.

Kashyap L, et al. (2007) Computational and molecular characterization of multiple isoforms of lfe-2 gene in nematode *C. elegans*. *Bioinformation*, 2(1), 17.

Bondurand N, et al. (2007) Deletions at the SOX10 gene locus cause Waardenburg syndrome types 2 and 4. *American journal of human genetics*, 81(6), 1169.

Ro S, et al. (2006) Template switching within exons 3 and 4 of KV11.1 (HERG) gives rise to a 5' truncated cDNA. *Biochemical and biophysical research communications*, 345(4), 1342.

Wandstrat AE, et al. (2004) Association of extensive polymorphisms in the SLAM/CD2 gene cluster with murine lupus. *Immunity*, 21(6), 769.

Wang Z, et al. (2004) A brief review of computational gene prediction methods. *Genomics, proteomics & bioinformatics*, 2(4), 216.