Resource Summary Report

Generated by NIF on Sep 3, 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: analysis service resource, data analysis service, production service resource, service resource

Keywords: gene

Resource Name: HMMgene

Resource ID: SCR_011933

Alternate IDs: OMICS_01488

Record Creation Time: 20220129T080307+0000

Record Last Update: 20260829T183008+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 NIF .

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.