

Resource Summary Report

Generated by FDI Lab - SciCrunch.org on Apr 8, 2025

Ensembl Genome Browser

RRID:SCR_013367

Type: Tool

Proper Citation

Ensembl Genome Browser (RRID:SCR_013367)

Resource Information

URL: <http://Mar2008.archive.ensembl.org>

Proper Citation: Ensembl Genome Browser (RRID:SCR_013367)

Description: THIS RESOURCE IS NO LONGER IN SERVICE. Documented on March 17,2022. Genome databases for vertebrates and other eukaryotic species. Analysis and annotation maintained on current data.Distribution of analysis to other bioinformatics laboratories. Ensembl concentrates on vertebrate genomes, but other groups have adapted system for use with plant and fungal genomes (see Powered by Ensembl list on website).

Resource Type: data or information resource, database

Funding: European Bioinformatics Institute ;
Wellcome Trust Sanger Institute

Availability: THIS RESOURCE IS NO LONGER IN SERVICE.

Resource Name: Ensembl Genome Browser

Resource ID: SCR_013367

Alternate IDs: nif-0000-30518

Alternate URLs: <http://uswest.ensembl.org/index.html>

Record Creation Time: 20220129T080315+0000

Record Last Update: 20250404T061005+0000

Ratings and Alerts

No rating or validation information has been found for Ensembl Genome Browser.

No alerts have been found for Ensembl Genome Browser.

Data and Source Information

Source: [SciCrunch Registry](#)

Usage and Citation Metrics

We found 1165 mentions in open access literature.

Listed below are recent publications. The full list is available at [FDI Lab - SciCrunch.org](#).

De Paolis V, et al. (2025) An antisense-long-noncoding-RNA modulates p75NTR expression levels during neuronal polarization. *iScience*, 28(1), 111566.

Lee SY, et al. (2025) Exploring the importance of predicted camel NRAP exon 4 for environmental adaptation using a mouse model. *Animal genetics*, 56(1), e13490.

Trastulli G, et al. (2025) Sample Tracking Tool: A Comprehensive Approach Based on OpenArray Technology and R Scripting for Genomic Sample Monitoring. *Diagnostics (Basel, Switzerland)*, 15(2).

Lim B, et al. (2025) Cell deconvolution-based integrated time-series network of whole blood transcriptome reveals systemic antiviral activities and cell-specific immunological changes against PRRSV infection. *Veterinary research*, 56(1), 19.

Raffle J, et al. (2025) Identification of novel genetic variants associated with feline cardiomyopathy using targeted next-generation sequencing. *Scientific reports*, 15(1), 3871.

Deebani A, et al. (2025) Knockdown of zebrafish tmem242 enhances the production of ROS that signals to increase f9a expression resulting in DIC-like condition. *Scientific reports*, 15(1), 3058.

Rolfs LA, et al. (2025) myh9b is a critical non-muscle myosin II encoding gene that interacts with myh9a and myh10 during zebrafish development in both compensatory and redundant pathways. *G3 (Bethesda, Md.)*, 15(1).

Bibi H, et al. (2025) Molecular and computational analysis of a novel pathogenic variant in emopamil-binding protein (EBP) involved in cholesterol biosynthetic pathway causing a rare male EBP disorder with neurologic defects (MEND syndrome). *Molecular biology reports*, 52(1), 101.

Vermani L, et al. (2025) A Haplotype GWAS in Syndromic Familial Colorectal Cancer. *International journal of molecular sciences*, 26(2).

Abuzahra M, et al. (2025) A novel p.127Val>Ile single nucleotide polymorphism in the MTNR1A gene and its relation to litter size in Thin-tailed Indonesian ewes. *Animal bioscience*, 38(2), 209.

Kontou D, et al. (2025) Adaptation in a keystone grazer under novel predation pressure. *Proceedings. Biological sciences*, 292(2039), 20241935.

Kasitipradit K, et al. (2025) Sex-specific effects of prenatal bisphenol A exposure on transcriptome-interactome profiles of autism candidate genes in neural stem cells from offspring hippocampus. *Scientific reports*, 15(1), 2882.

DeMeis JD, et al. (2025) Long G4-rich enhancers target promoters via a G4 DNA-based mechanism. *Nucleic acids research*, 53(2).

Harvey A, et al. (2025) Uncovering possible silent acquired long QT syndrome using exercise stress testing in long-term pediatric acute lymphoblastic leukemia survivors. *International journal of cancer*, 156(2), 403.

Morillo-Bernal J, et al. (2024) HuR (ELAVL1) Stabilizes SOX9 mRNA and Promotes Migration and Invasion in Breast Cancer Cells. *Cancers*, 16(2).

Hauck JS, et al. (2024) Heat shock factor 1 directly regulates transsulfuration pathway to promote prostate cancer proliferation and survival. *Communications biology*, 7(1), 9.

Duijvelaar E, et al. (2024) Longitudinal plasma proteomics reveals biomarkers of alveolar-capillary barrier disruption in critically ill COVID-19 patients. *Nature communications*, 15(1), 744.

Wanniarachchi DV, et al. (2024) The evaluation of transcription factor binding site prediction tools in human and Arabidopsis genomes. *BMC bioinformatics*, 25(1), 371.

Missailidis D, et al. (2024) A blood-based mRNA signature distinguishes people with Long COVID from recovered individuals. *Frontiers in immunology*, 15, 1450853.

Gholami M, et al. (2024) Genetic Variants and Haplotype Structures in the CASC Gene Family to Predict Cancer Risk: A Bioinformatics Study. *Health science reports*, 7(12), e70228.