Resource Summary Report

Generated by RRID on Apr 11, 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: 20250410T070359+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 RRID.

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.

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).

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.

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

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.

Manzoor Y, et al. (2024) Loss of the yeast transporter Agp2 upregulates the pleiotropic drug-resistant pump Pdr5 and confers resistance to the protein synthesis inhibitor cycloheximide. PloS one, 19(5), e0303747.

Zheng W, et al. (2024) RUNX1-induced upregulation of PTGS2 enhances cell growth, migration and invasion in colorectal cancer cells. Scientific reports, 14(1), 11670.

Rajput P, et al. (2024) Differential Vulnerability and Response to Injury among Brain Cell Types Comprising the Neurovascular Unit. The Journal of neuroscience : the official journal of the Society for Neuroscience, 44(22).

Pasquini L, et al. (2024) Frontotemporal lobar degeneration targets brain regions linked to expression of recently evolved genes. Brain : a journal of neurology, 147(9), 3032.

Akinsola OM, et al. (2024) Genomic insights into adaptation and inbreeding among Sub-Saharan African cattle from pastoral and agropastoral systems. Frontiers in genetics, 15, 1430291.

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.