

Resource Summary Report

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BioMart MartView

RRID:SCR_010714

Type: Tool

Proper Citation

BioMart MartView (RRID:SCR_010714)

Resource Information

URL: <http://www.biomart.org/biomart/martview/ddb9ce1ad275cde372c968d13fa11f5f>

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Description: A web server interface of BioMart software and provides a unified view over disparate data sources that enable bioscientists to retrieve data from one or multiple sources in a simple and efficient way. This MartView web server features seamless data federation making cross querying of data sources in a user friendly and unified way. Data sources include major biomolecular sequence, pathway and annotation databases such as Ensembl, Uniprot, Reactome, HGNC, Wormbase, etc. The web server not only provides access through a web interface, it also supports programmatic access through a Perl API as well as RESTful and SOAP oriented web services.

Synonyms: MartView, BioMart Central Portal

Resource Type: data set, web service, software resource, data access protocol, data or information resource

Defining Citation: [PMID:19420058](#), [PMID:19144180](#)

Keywords: gold standard

Funding: Ontario Institute for Cancer Research ;
Wellcome Trust ;
EMBL ;
European Union FP6 contract LHSG-CT-2004-512092

Availability: Free, Public

Resource Name: BioMart MartView

Resource ID: SCR_010714

Alternate IDs: nlx_89178

Record Creation Time: 20220129T080300+0000

Record Last Update: 20260728T043350+0000

Ratings and Alerts

No rating or validation information has been found for BioMart MartView.

No alerts have been found for BioMart MartView.

Data and Source Information

Source: [SciCrunch Registry](#)

Usage and Citation Metrics

We found 10 mentions in open access literature.

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

Kalef-Ezra E, et al. (2024) Single-cell somatic copy number variants in brain using different amplification methods and reference genomes. Communications biology, 7(1), 1288.

Sen D, et al. (2023) Metabolic regulation of CTCF expression and chromatin association dictates starvation response in mice and flies. iScience, 26(7), 107128.

Howard-McCombe J, et al. (2023) Genetic swamping of the critically endangered Scottish wildcat was recent and accelerated by disease. Current biology : CB, 33(21), 4761.

Tavares M, et al. (2022) JAZF1-SUZ12 dysregulates PRC2 function and gene expression during cell differentiation. Cell reports, 39(9), 110889.

Skalska L, et al. (2021) Nascent RNA antagonizes the interaction of a set of regulatory proteins with chromatin. Molecular cell, 81(14), 2944.

Li J, et al. (2021) KDM6B-dependent chromatin remodeling underpins effective virus-specific CD8+ T cell differentiation. Cell reports, 34(11), 108839.

Pouyet F, et al. (2017) Recombination, meiotic expression and human codon usage. eLife, 6.

Sako H, et al. (2016) Genome-Wide Analysis of Acute Endurance Exercise-Induced Translational Regulation in Mouse Skeletal Muscle. PloS one, 11(2), e0148311.

Hangauer MJ, et al. (2013) Pervasive transcription of the human genome produces thousands of previously unidentified long intergenic noncoding RNAs. PLoS genetics, 9(6), e1003569.

Ghosh S, et al. (2012) Analysis of polymorphisms and haplotype structure of the human thymidylate synthase genetic region: a tool for pharmacogenetic studies. PloS one, 7(4),

e34426.