

Resource Summary Report

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GenoList genome browser

RRID:SCR_007683

Type: Tool

Proper Citation

GenoList genome browser (RRID:SCR_007683)

Resource Information

URL: <http://genolist.pasteur.fr/>

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Description: GenoList is a resource for microbial genomes. It provides links to databases that give information on specific microbes. A link to the Genolist itself also incorporates many microbial genomes and search and analysis tools. Of the links provided on the GenoList homepage, species included are: Bacillus subtilis Escherichia coli Mycobacterium tuberculosis Mycobacterium leprae Mycobacterium bovis Mycobacterium ulcerans Listeria monocytogenes Legionella pneumophila Helicobacter pylori Mycoplasma pulmonis Synechocystis Anabaena Staphylococcus aureus Streptococcus pneumoniae Streptococcus agalactiae Photorhabdus luminescens Candida albicans Bacillus subtilis Escherichia coli Mycobacterium tuberculosis Mycobacterium leprae Mycobacterium bovis Mycobacterium ulcerans Listeria monocytogenes Legionella pneumophila Helicobacter pylori Mycoplasma pulmonis Synechocystis Anabaena Staphylococcus aureus Streptococcus pneumoniae Streptococcus agalactiae Photorhabdus luminescens Candida albicans

Synonyms: GenoList

Resource Type: data or information resource, database

Keywords: bacillus subtilis escherichia coli mycobacterium tuberculosis mycobacterium leprae mycobacterium bovis mycobacterium ulcerans listeria monocytogenes legionella pneumophila helicobacter pylori mycoplasma pulmonis synechocystis anabaena staphylococcus aureus streptococcus pneumoniae streptococcus agalactiae photorhabdus luminescens candida albicans

Resource Name: GenoList genome browser

Resource ID: SCR_007683

Alternate IDs: nif-0000-02895

Record Creation Time: 20220129T080243+0000

Record Last Update: 20260729T044153+0000

Ratings and Alerts

No rating or validation information has been found for GenoList genome browser.

No alerts have been found for GenoList genome browser.

Data and Source Information

Source: [SciCrunch Registry](#)

Usage and Citation Metrics

We found 323 mentions in open access literature.

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

Nakatochi M, et al. (2025) Copy number variations in RNF216 and postsynaptic membrane-associated genes are associated with bipolar disorder: a case-control study in the Japanese population. *Psychiatry and clinical neurosciences*, 79(1), 12.

Amatruda M, et al. (2025) Neuroprotective effect of neuron-specific deletion of the C16 ceramide synthetic enzymes in an animal model of multiple sclerosis. *Glia*, 73(2), 271.

Zarka KA, et al. (2024) T-DNA characterization of genetically modified 3-R-gene late blight-resistant potato events with a novel procedure utilizing the Samplix Xdrop?? enrichment technology. *Frontiers in plant science*, 15, 1330429.

Liu YN, et al. (2024) Immunosuppressive role of BDNF in therapy-induced neuroendocrine prostate cancer. *Molecular oncology*.

Steux C, et al. (2024) The Maintenance of Deleterious Variation in Wild Chinese Rhesus Macaques. *bioRxiv : the preprint server for biology*.

Wei WH, et al. (2024) Human-specific insights into candidate genes and boosted discoveries of novel loci illuminate roles of neuroglia in reading disorders. *Genes, brain, and behavior*, 23(3), e12899.

Dong J, et al. (2024) ZNF143 binds DNA and stimulates transcription initiation to activate and repress direct target genes. *bioRxiv : the preprint server for biology*.

Morales-Vicente DA, et al. (2024) The Human Developing Cerebral Cortex Is Characterized by an Elevated De Novo Expression of Long Noncoding RNAs in Excitatory Neurons. *Molecular biology and evolution*, 41(7).

Tadaka S, et al. (2024) jMorp: Japanese Multi-Omics Reference Panel update report 2023. *Nucleic acids research*, 52(D1), D622.

Yang J, et al. (2024) rNMPID: a database for riboNucleoside MonoPhosphates in DNA. *Bioinformatics advances*, 4(1), vbae063.

Steux C, et al. (2024) The Maintenance of Deleterious Variation in Wild Chinese Rhesus Macaques. *Genome biology and evolution*, 16(6).

Wei J, et al. (2024) Temporospatial hierarchy and allele-specific expression of zygotic genome activation revealed by distant interspecific urochordate hybrids. *Nature communications*, 15(1), 2395.

Tsui KH, et al. (2024) WNT1-inducible signaling pathway protein 1 activation through C-X-C motif chemokine ligand 5/C-X-C chemokine receptor type 2/leukemia inhibitory factor/leukemia inhibitory factor receptor signaling promotes immunosuppression and neuroendocrine differentiation in prostate cancer. *iScience*, 27(8), 110562.

Kilic A, et al. (2024) Vitamin D constrains inflammation by modulating the expression of key genes on Chr17q12-21.1. *eLife*, 12.

Miller CD, et al. (2024) Pan-Cancer Interrogation of B7-H3 (CD276) as an Actionable Therapeutic Target Across Human Malignancies. *Cancer research communications*, 4(5), 1369.

Fekete N, et al. (2024) Flow Cytometry-Based Assay to Detect Alpha Galactosidase Enzymatic Activity at the Cellular Level. *Cells*, 13(8).

Cao YL, et al. (2024) Wolfberry genome database: integrated genomic datasets for studying molecular biology. *Frontiers in plant science*, 15, 1310346.

Koehler AC, et al. (2024) Gene model for the ortholog of Dsor1 in *Drosophila persimilis*. *microPublication biology*, 2024.

Shi Q, et al. (2024) DRED: A Comprehensive Database of Genes Related to Repeat Expansion Diseases. *Genomics, proteomics & bioinformatics*, 22(5).

Nuttle X, et al. (2024) Parallelized engineering of mutational models using piggyBac transposon delivery of CRISPR libraries. *Cell reports methods*, 4(1), 100672.