Resource Summary Report

Generated by dkNET on May 20, 2025

Genome Reviews

RRID:SCR_007685
Type: Tool

Proper Citation

Genome Reviews (RRID:SCR_007685)

Resource Information

URL: http://www.ebi.ac.uk/GenomeReviews/

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Description: THIS RESOURCE IS NO LONGER IN SERVICE, documented April 24, 2017. The Genome Reviews database provides an up-to-date, standardized and comprehensively annotated view of the genomic sequence of organisms with completely deciphered genomes. Currently, Genome Reviews contains the genomes of archaea, bacteria, bacteriophages and selected eukaryota. Genome Reviews is available as a MySQL relational database, or a flat file format derived from that in the EMBL Nucleotide Sequence Database. An Ensembl-style browser is now available for Genome Reviews, providing a zoomable graphical view of all chromosomes and plasmids represented in the database. The location and structure of all genes is shown and the distribution of features throughout the sequence is displayed.

Synonyms: Genome Reviews Database

Resource Type: database, data or information resource

Defining Citation: PMID:16901215, PMID:15608201

Keywords: complete genome sequence, genome sequence, nucleotides, gold standard, bio.tools

Funding: European Union FELICS contract 021902 (RII3);
European Union TEMBLOR contract QLRI- CT-2001000015

Availability: THIS RESOURCE IS NO LONGER IN SERVICE

Resource Name: Genome Reviews

Resource ID: SCR_007685

Alternate IDs: nif-0000-02899, biotools:genomereviews

Alternate URLs: https://bio.tools/genomereviews

Record Creation Time: 20220129T080243+0000

Record Last Update: 20250519T204730+0000

Ratings and Alerts

No rating or validation information has been found for Genome Reviews.

No alerts have been found for Genome Reviews.

Data and Source Information

Source: SciCrunch Registry

Usage and Citation Metrics

We found 4 mentions in open access literature.

Listed below are recent publications. The full list is available at dkNET.

de Jong A, et al. (2013) The transcriptional and gene regulatory network of Lactococcus lactis MG1363 during growth in milk. PloS one, 8(1), e53085.

Sanna-Cherchi S, et al. (2012) Copy-number disorders are a common cause of congenital kidney malformations. American journal of human genetics, 91(6), 987.

Merkl R, et al. (2009) GO4genome: a prokaryotic phylogeny based on genome organization. Journal of molecular evolution, 68(5), 550.

Galperin MY, et al. (2005) The Molecular Biology Database Collection: 2005 update. Nucleic acids research, 33(Database issue), D5.