

Resource Summary Report

Generated by SPARC SAWG on Sep 17, 2026

NCBI Genome Workbench

RRID:SCR_011794

Type: Tool

Proper Citation

NCBI Genome Workbench (RRID:SCR_011794)

Resource Information

URL: <http://www.ncbi.nlm.nih.gov/tools/gbench/>

Proper Citation: NCBI Genome Workbench (RRID:SCR_011794)

Description: An integrated application for viewing and analyzing sequence data.

Abbreviations: Genome Workbench

Resource Type: software resource

Keywords: bio.tools

Resource Name: NCBI Genome Workbench

Resource ID: SCR_011794

Alternate IDs: OMICS_00920, biotools:ncbi_genome_workbench

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

Record Creation Time: 20220129T080306+0000

Record Last Update: 20260912T195739+0000

Ratings and Alerts

No rating or validation information has been found for NCBI Genome Workbench.

No alerts have been found for NCBI Genome Workbench.

Data and Source Information

Source: [SciCrunch Registry](#)

Usage and Citation Metrics

We found 11 mentions in open access literature.

Listed below are recent publications. The full list is available at [SPARC SAWG](#) .

Futas J, et al. (2024) Comparative genomics of the Natural Killer Complex in carnivores. *Frontiers in immunology*, 15, 1459122.

Ferguson KB, et al. (2021) Jekyll or Hyde? The genome (and more) of *Nesidiocoris tenuis*, a zoophytophagous predatory bug that is both a biological control agent and a pest. *Insect molecular biology*, 30(2), 188.

Rajput B, et al. (2019) RefSeq curation and annotation of stop codon recoding in vertebrates. *Nucleic acids research*, 47(2), 594.

Logacheva MD, et al. (2016) Comparative analysis of plastid genomes of non-photosynthetic Ericaceae and their photosynthetic relatives. *Scientific reports*, 6, 30042.

Lv M, et al. (2016) A Family of CS?? Defensins and Defensin-Like Peptides from the Migratory Locust, *Locusta migratoria*, and Their Expression Dynamics during Mycosis and Nosemosis. *PloS one*, 11(8), e0161585.

O'Leary NA, et al. (2016) Reference sequence (RefSeq) database at NCBI: current status, taxonomic expansion, and functional annotation. *Nucleic acids research*, 44(D1), D733.

Pavlopoulos GA, et al. (2015) Visualizing genome and systems biology: technologies, tools, implementation techniques and trends, past, present and future. *GigaScience*, 4, 38.

Rajput B, et al. (2015) RefSeq curation and annotation of antizyme and antizyme inhibitor genes in vertebrates. *Nucleic acids research*, 43(15), 7270.

Warren WC, et al. (2015) The genome of the vervet (*Chlorocebus aethiops sabaeus*). *Genome research*, 25(12), 1921.

Brown JC, et al. (2014) The role of DNA repair in herpesvirus pathogenesis. *Genomics*, 104(4), 287.

Di Cesare A, et al. (2013) Aquaculture can promote the presence and spread of antibiotic-resistant Enterococci in marine sediments. *PloS one*, 8(4), e62838.