

# Resource Summary Report

Generated by [ASWG](#) on Aug 2, 2026

## NCBI Genome Workbench

RRID:SCR\_011794

Type: Tool

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### Proper Citation

NCBI Genome Workbench (RRID:SCR\_011794)

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### 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](https://bio.tools/ncbi_genome_workbench)

**Record Creation Time:** 20220129T080306+0000

**Record Last Update:** 20260801T190418+0000

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### Ratings and Alerts

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

No alerts have been found for NCBI Genome Workbench.

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### 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 [ASWG](#) .

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.

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

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

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.