# Resource Summary Report

Generated by dkNET on May 19, 2025

## 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

**Funding:**

**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:** 20250420T014559+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 dkNET.


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.