

# Resource Summary Report

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## CLC Genomics Workbench

RRID:SCR\_011853

Type: Tool

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### Proper Citation

CLC Genomics Workbench (RRID:SCR\_011853)

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### Resource Information

**URL:** <http://www.clcbio.com/products/clc-genomics-workbench/>

**Proper Citation:** CLC Genomics Workbench (RRID:SCR\_011853)

**Description:** Commercially available software for visualization and analysis of next generation sequencing data. Used for viewing, exploring, and sharing of NGS analysis results. Complete toolkit for genomics, transcriptomics, epigenomics, and metagenomics in one program.

**Resource Type:** data analysis software, data processing software, data visualization software, software application, software resource

**Keywords:** ngs, next, generation, sequencing, gene, rna, visualisation, analysis

**Availability:** Restricted

**Resource Name:** CLC Genomics Workbench

**Resource ID:** SCR\_011853

**Alternate IDs:** SCR\_016245, OMICS\_01124

**License:** Commercial license

**Record Creation Time:** 20220129T080307+0000

**Record Last Update:** 20260821T193920+0000

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### Ratings and Alerts

No rating or validation information has been found for CLC Genomics Workbench.

No alerts have been found for CLC Genomics Workbench.

## Data and Source Information

**Source:** [SciCrunch Registry](#)

## Usage and Citation Metrics

We found 187 mentions in open access literature.

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

Molotkova A, et al. (2026) ROME, an Ancient Gene with a Novel Function in Vertebrates, Is a Key Modulator of Embryonal Development and Cancer Metastasis. *Cancer research communications*, 6(3), 477.

Telford MS, et al. (2026) Antibacterial microcins self-assemble to promote biofilm formation in *Acinetobacter baumannii*. *Cell reports*, 45(4), 117156.

Bjornson S, et al. (2026) Stepwise evolutionary transitions in Cladophorales plastid genomes reveal origins of hairpin chromosomes. *Current biology : CB*, 36(7), 1659.

Wang G, et al. (2026) VGLL4 promotes hepatocellular carcinoma progression via the Wnt/ $\beta$ -catenin pathway. *Tissue & cell*, 100, 103391.

Casey-Moore MC, et al. (2025) Genomic Analysis of Rotavirus G8P[8] Strains Detected in the United States Through Active Surveillance, 2016-2017. *Viruses*, 17(9).

Franck L, et al. (2025) Elevated hexosylceramides in Parkinson's disease cause gene upregulations in neurons mimicking responses to pathogens. *NPJ Parkinson's disease*, 11(1), 268.

Mehta I, et al. (2025) Control of pili synthesis and putrescine homeostasis in *Escherichia coli*. *eLife*, 13.

Diolintzi A, et al. (2025) Novel Hyperplastic Expansion of White Adipose Tissue Underlies the Metabolically Healthy Obese Phenotype of Male LFABP Null Mice. *Cells*, 14(11).

Turner LA, et al. (2025) Differences in gene expression between high and low tolerance rainbow trout (*Oncorhynchus mykiss*) to acute thermal stress. *PloS one*, 20(1), e0312694.

Schmidt KG, et al. (2025) Coronavirus replicase epitopes induce cross-reactive CD8 T cell responses in SARS-CoV-2-naïve people with HIV-1. *iScience*, 28(3), 111949.

Colgan DJ, et al. (2024) The Families of Non-LTR Transposable Elements within Neritimorpha and Other Gastropoda. *Genes*, 15(6).

Cheng J, et al. (2024) Myeloid cells coordinately induce glioma cell-intrinsic and cell-extrinsic pathways for chemoresistance via GP130 signaling. *Cell reports. Medicine*, 5(8), 101658.

Wang Y, et al. (2024) The Mohawk homeobox gene represents a marker and osteo-inhibitory factor in calvarial suture osteoprogenitor cells. *Cell death & disease*, 15(6), 420.

Itoh M, et al. (2024) Molecular alterations associated with pathophysiology in liver-specific ZO-1 and ZO-2 knockout mice. *Cell structure and function*, 49(2), 83.

Sprenger M, et al. (2024) Small RNAs direct attack and defense mechanisms in a quorum sensing phage and its host. *Cell host & microbe*, 32(5), 727.

Kim SY, et al. (2024) Antibacterial action, proteolytic immunity, and in vivo activity of a *Vibrio cholerae* microcin. *Cell host & microbe*, 32(11), 1959.

Reeves KD, et al. (2024) Mapping the geographical distribution of the mucosa-associated gut microbiome in GI-symptomatic children with autism spectrum disorder. *American journal of physiology. Gastrointestinal and liver physiology*, 327(2), G217.

Onuma K, et al. (2024) Bardoxolone methyl prevents metabolic dysfunction-associated steatohepatitis by inhibiting macrophage infiltration. *British journal of pharmacology*, 181(15), 2545.

Kitt EM, et al. (2023) Genotypic investigation of a rotavirus cluster at a quaternary-care pediatric hospital. *Infection control and hospital epidemiology*, 44(10), 1680.

Chapagain P, et al. (2023) Dual RNA-Seq of *Flavobacterium psychrophilum* and Its Outer Membrane Vesicles Distinguishes Genes Associated with Susceptibility to Bacterial Cold-Water Disease in Rainbow Trout (*Oncorhynchus mykiss*). *Pathogens (Basel, Switzerland)*, 12(3).