

Resource Summary Report

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SnapGene Viewer

RRID:SCR_015053

Type: Tool

Proper Citation

SnapGene Viewer (RRID:SCR_015053)

Resource Information

URL: http://www.snapgene.com/products/snapgene_viewer/

Proper Citation: SnapGene Viewer (RRID:SCR_015053)

Description: Visualization software that allows users to create, browse, and share annotated DNA sequence files from SnapGene.

Resource Type: annotation tool

Keywords: plasmid map, plasmid visualization, plasmid creation

Funding:

Availability: Available for download

Resource Name: SnapGene Viewer

Resource ID: SCR_015053

Record Creation Time: 20220129T080323+0000

Record Last Update: 20250525T031404+0000

Ratings and Alerts

No rating or validation information has been found for SnapGene Viewer.

No alerts have been found for SnapGene Viewer.

Data and Source Information

Source: [SciCrunch Registry](#)

Usage and Citation Metrics

We found 47 mentions in open access literature.

Listed below are recent publications. The full list is available at [FDI Lab - SciCrunch.org](#).

Hernández-Orozco AA, et al. (2024) Frequency of rs1051338 and rs116928232 Variants in Individuals from Northwest Mexico. *Journal of clinical laboratory analysis*, 38(13-14), e25083.

Pavlichenko VV, et al. (2024) Simplified Method for Agrobacterium-Mediated Genetic Transformation of *Populus x berolinensis* K. Koch. *Methods and protocols*, 7(1).

Buckingham AB, et al. (2024) Optimization of a lentivirus-mediated gene therapy targeting HIV-1 RNA to eliminate HIV-1-infected cells. *Molecular therapy. Nucleic acids*, 35(4), 102341.

Wang L, et al. (2024) A Novel Splicing Mutation Leading to Wiskott-Aldrich Syndrome from a Family. *International journal of genomics*, 2024, 2277956.

Ren CY, et al. (2024) Engineered CBEs based on *Macaca fascicularis* A3A with improved properties for precise genome editing. *Cell reports*, 43(3), 113878.

Nejabat S, et al. (2024) Immunoinformatics approach: Developing a multi-epitope vaccine with novel carriers targeting monkeypox virus. *FASEB journal : official publication of the Federation of American Societies for Experimental Biology*, 38(24), e70257.

Wisniewski DJ, et al. (2023) Flotillin-2 regulates epidermal growth factor receptor activation, degradation by Cbl-mediated ubiquitination, and cancer growth. *The Journal of biological chemistry*, 299(1), 102766.

Xedzro C, et al. (2023) Emergence of colistin-resistant *Enterobacter cloacae* and *Raoultella ornithinolytica* carrying the phosphoethanolamine transferase gene, mcr-9, derived from vegetables in Japan. *Microbiology spectrum*, 11(6), e0106323.

Kleinova R, et al. (2023) The ADAR1 editome reveals drivers of editing-specificity for ADAR1-isoforms. *Nucleic acids research*, 51(9), 4191.

Yang J, et al. (2023) Development of highly accurate digital PCR method and reference material for monkeypox virus detection. *Analytical and bioanalytical chemistry*, 415(7), 1333.

Kapoor S, et al. (2023) PP2A-B55SUR-6 promotes nuclear envelope breakdown in *C. elegans* embryos. *Cell reports*, 42(12), 113495.

Batu Öztürk A, et al. (2023) Conditioned media of mouse macrophages modulates neuronal

dynamics in mouse hippocampal cells. *International immunopharmacology*, 114, 109548.

Rashid FM, et al. (2023) The environmentally-regulated interplay between local three-dimensional chromatin organisation and transcription of proVWX in *E. coli*. *Nature communications*, 14(1), 7478.

Ajay AK, et al. (2022) Deletion of STAT3 from Foxd1 cell population protects mice from kidney fibrosis by inhibiting pericytes trans-differentiation and migration. *Cell reports*, 38(10), 110473.

Sousos N, et al. (2022) In utero origin of myelofibrosis presenting in adult monozygotic twins. *Nature medicine*, 28(6), 1207.

Jin WB, et al. (2022) Genetic manipulation of gut microbes enables single-gene interrogation in a complex microbiome. *Cell*, 185(3), 547.

Riva C, et al. (2022) A natural transdifferentiation event involving mitosis is empowered by integrating signaling inputs with conserved plasticity factors. *Cell reports*, 40(12), 111365.

Xie W, et al. (2022) A high-concentrate diet provokes inflammatory responses by downregulating Forkhead box protein A2 (FOXA2) through epigenetic modifications in the liver of dairy cows. *Gene*, 837, 146703.

Fang J, et al. (2022) Functional interactomes of the Ebola virus polymerase identified by proximity proteomics in the context of viral replication. *Cell reports*, 38(12), 110544.

Acs-Szabo L, et al. (2021) Genome Comparisons of the Fission Yeasts Reveal Ancient Collinear Loci Maintained by Natural Selection. *Journal of fungi (Basel, Switzerland)*, 7(10).