

Resource Summary Report

Generated by [PRECISE-TBI](#) on Sep 8, 2026

Tablet

RRID:SCR_000017

Type: Tool

Proper Citation

Tablet (RRID:SCR_000017)

Resource Information

URL: <http://bioinf.scri.ac.uk/tablet/>

Proper Citation: Tablet (RRID:SCR_000017)

Description: A lightweight, high-performance graphical viewer for next generation sequence assemblies and alignments.

Abbreviations: Tablet

Synonyms: Tablet - Next Generation Sequence Assembly Visualization

Resource Type: software resource

Defining Citation: [PMID:22445902](#)

Keywords: next generation sequence, assembly, alignment

Availability: Free, Available for download, Freely available

Resource Name: Tablet

Resource ID: SCR_000017

Alternate IDs: OMICS_00896

Record Creation Time: 20220129T080159+0000

Record Last Update: 20260905T132408+0000

Ratings and Alerts

No rating or validation information has been found for Tablet.

No alerts have been found for Tablet.

Data and Source Information

Source: [SciCrunch Registry](#)

Usage and Citation Metrics

We found 8 mentions in open access literature.

Listed below are recent publications. The full list is available at [PRECISE-TBI](#) .

Sotcheff S, et al. (2023) ViReMa: a virus recombination mapper of next-generation sequencing data characterizes diverse recombinant viral nucleic acids. GigaScience, 12.

Jaworski E, et al. (2021) Tiled-ClickSeq for targeted sequencing of complete coronavirus genomes with simultaneous capture of RNA recombination and minority variants. eLife, 10.

Holmes A, et al. (2016) Mechanistic signatures of HPV insertions in cervical carcinomas. NPJ genomic medicine, 1, 16004.

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

de Brevern AG, et al. (2015) Trends in IT Innovation to Build a Next Generation Bioinformatics Solution to Manage and Analyse Biological Big Data Produced by NGS Technologies. BioMed research international, 2015, 904541.

Valach M, et al. (2014) Trans-splicing and RNA editing of LSU rRNA in Diplonema mitochondria. Nucleic acids research, 42(4), 2660.

Cazalla D, et al. (2011) A primate herpesvirus uses the integrator complex to generate viral microRNAs. Molecular cell, 43(6), 982.

Barchi L, et al. (2011) Identification of SNP and SSR markers in eggplant using RAD tag sequencing. BMC genomics, 12, 304.