

# Resource Summary Report

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

## HilbertVis

RRID:SCR\_007862

Type: Tool

### Proper Citation

HilbertVis (RRID:SCR\_007862)

### Resource Information

**URL:** <http://www.ebi.ac.uk/huber-srv/hilbert/>

**Proper Citation:** HilbertVis (RRID:SCR\_007862)

**Description:** Software tool that allows to display very long data vectors in a space-efficient manner, allowing the user to visually judge the large scale structure and distribution of features simultaneously with the rough shape and intensity of individual features.

**Abbreviations:** HilbertVis

**Resource Type:** software resource

**Defining Citation:** [DOI:10.1093/bioinformatics/btp152](https://doi.org/10.1093/bioinformatics/btp152)

**Keywords:** bio.tools

**Resource Name:** HilbertVis

**Resource ID:** SCR\_007862

**Alternate IDs:** OMICS\_00627, biotools:hilbertvis

**Alternate URLs:** <https://bio.tools/hilbertvis>, <https://sources.debian.org/src/r-bioc-hilbertvis/>

**Record Creation Time:** 20220129T080244+0000

**Record Last Update:** 20260905T132615+0000

### Ratings and Alerts

No rating or validation information has been found for HilbertVis.

No alerts have been found for HilbertVis.

## Data and Source Information

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

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## Usage and Citation Metrics

We found 4 mentions in open access literature.

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

Gaggioli V, et al. (2023) Dynamic de novo heterochromatin assembly and disassembly at replication forks ensures fork stability. *Nature cell biology*, 25(7), 1017.

Reverón-Gómez N, et al. (2018) Accurate Recycling of Parental Histones Reproduces the Histone Modification Landscape during DNA Replication. *Molecular cell*, 72(2), 239.

Jansen G, et al. (2015) Evolutionary Transition from Pathogenicity to Commensalism: Global Regulator Mutations Mediate Fitness Gains through Virulence Attenuation. *Molecular biology and evolution*, 32(11), 2883.

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