

# Resource Summary Report

Generated by [Kravitz](#) on Sep 10, 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](#)

---

## Usage and Citation Metrics

We found 4 mentions in open access literature.

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

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.