

Resource Summary Report

Generated by [RRID](#) on Aug 2, 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: 20260801T190334+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 [RRID](#) .

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.

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

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.