

Resource Summary Report

Generated by [dkNET](#) on Aug 26, 2026

GenePattern Notebook

RRID:SCR_015699

Type: Tool

Proper Citation

GenePattern Notebook (RRID:SCR_015699)

Resource Information

URL: <http://www.genepattern-notebook.org/>

Proper Citation: GenePattern Notebook (RRID:SCR_015699)

Description: Interactive analysis notebook environment that streamlines genomics research by interleaving text, multimedia, and executable code into unified, sharable, reproducible “research narratives.” It integrates the dynamic capabilities of notebook systems with an investigator-focused, simple interface that provides access to hundreds of genomic tools without the need to write code.

Synonyms: GenePattern Notebook environment

Resource Type: electronic laboratory notebook, software application, software resource, systems interoperability software, web application

Defining Citation: [PMID:28822753](#)

Keywords: gene, genomics research, research narrative, notebook system, analysis notebook, bio.tools

Funding: NIGMS R01-GM074024;
NCI U24-CA194107

Availability: Open Source, Free, Available for download, Account required

Resource Name: GenePattern Notebook

Resource ID: SCR_015699

Alternate IDs: biotools:GenePattern_notebook

Alternate URLs: https://bio.tools/GenePattern_notebook

License: BSD-style license

Record Creation Time: 20220129T080327+0000

Record Last Update: 20260821T194025+0000

Ratings and Alerts

No rating or validation information has been found for GenePattern Notebook .

No alerts have been found for GenePattern Notebook .

Data and Source Information

Source: [SciCrunch Registry](#)

Usage and Citation Metrics

We found 3 mentions in open access literature.

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

Chen K, et al. (2021) A phenotypically supervised single-cell analysis protocol to study within-cell-type heterogeneity of cultured mammalian cells. STAR protocols, 2(2), 100561.

Mah CK, et al. (2018) An accessible GenePattern notebook for the copy number variation analysis of Illumina Infinium DNA methylation arrays. F1000Research, 7.

Reich M, et al. (2017) The GenePattern Notebook Environment. Cell systems, 5(2), 149.