

Resource Summary Report

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

Phenocarta

RRID:SCR_016273

Type: Tool

Proper Citation

Phenocarta (RRID:SCR_016273)

Resource Information

URL: <https://gemma.msl.ubc.ca/phenotypes.html>

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Description: Database that consolidates information on genes and phenotypes across multiple resources and allows tracking and exploring of the associations. Part of Gemma, a web site, database and a set of tools for the meta-analysis, re-use and sharing of genomics data.

Synonyms: Phenocarta

Resource Type: data analysis software, data or information resource, data processing software, database, software application, software resource

Defining Citation: [PMID:22782548](#)

Keywords: phenotype, meta, analysis, genome, gene, expression

Availability: Free, Freely available

Resource Name: Phenocarta

Resource ID: SCR_016273

Alternate URLs: <http://pavlab.msl.ubc.ca/software-and-resources/>

License: Apache 2.0 License

License URLs: <https://pavlidislab.github.io/Gemma/#terms-and-conditions>

Record Creation Time: 20220129T080329+0000

Record Last Update: 20260905T132806+0000

Ratings and Alerts

No rating or validation information has been found for Phenocarta.

No alerts have been found for Phenocarta.

Data and Source Information

Source: [SciCrunch Registry](#)

Usage and Citation Metrics

We found 2 mentions in open access literature.

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

Na D, et al. (2023) A multi-layered network model identifies Akt1 as a common modulator of neurodegeneration. Molecular systems biology, 19(12), e11801.

Ratajczak F, et al. (2023) Speos: an ensemble graph representation learning framework to predict core gene candidates for complex diseases. Nature communications, 14(1), 7206.