# Resource Summary Report

Generated by FDI Lab - SciCrunch.org on Apr 11, 2025

## Phenocarta

RRID:SCR_016273
Type: Tool

### Proper Citation

Phenocarta (RRID:SCR_016273)

### Resource Information

**URL:** https://gemma.msl.ubc.ca/phenotypes.html

**Proper Citation:** Phenocarta (RRID:SCR_016273)

**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 processing software, database, software application, data or information resource, software resource, data analysis software

**Defining Citation:** PMID:22782548

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

**Funding:**

**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:** 20250411T055901+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 FDI Lab - SciCrunch.org.

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.

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