

Resource Summary Report

Generated by SPARC SAWG on Sep 19, 2026

PhenoFam

RRID:SCR_000640

Type: Tool

Proper Citation

PhenoFam (RRID:SCR_000640)

Resource Information

URL: <http://sourceforge.net/projects/phenofam/>

Proper Citation: PhenoFam (RRID:SCR_000640)

Description: A web-based application that performs gene set enrichment analysis (GSEA) by employing structural and functional information on families of protein domains as annotation terms.

Abbreviations: PhenoFam

Resource Type: software application, software resource

Defining Citation: [PMID:20478033](https://pubmed.ncbi.nlm.nih.gov/20478033/)

Keywords: java, javascript, gene, gene set enrichment analysis, structure, function, protein domain, bio.tools

Availability: Free, Available for download, Freely available

Resource Name: PhenoFam

Resource ID: SCR_000640

Alternate IDs: OMICS_02230, biotools:phenofam

Alternate URLs: <https://bio.tools/phenofam>

License: GNU General Public License, v3

Record Creation Time: 20220129T080202+0000

Record Last Update: 20260912T200224+0000

Ratings and Alerts

No rating or validation information has been found for PhenoFam.

No alerts have been found for PhenoFam.

Data and Source Information

Source: [SciCrunch Registry](#)

Usage and Citation Metrics

We have not found any literature mentions for this resource.