

Resource Summary Report

Generated by T1D on Sep 5, 2026

Human Phenotype Ontology Annotations

RRID:SCR_006219

Type: Tool

Proper Citation

Human Phenotype Ontology Annotations (RRID:SCR_006219)

Resource Information

URL: <http://www.human-phenotype-ontology.org>

Proper Citation: Human Phenotype Ontology Annotations (RRID:SCR_006219)

Description: Downloadable dataset of Human Phenotype Ontology (HPO) annotations. Provides standardized vocabulary of phenotypic abnormalities encountered in human disease.

Abbreviations: HPO Annotations

Resource Type: data set, data or information resource

Defining Citation: [PMID:20412080](https://pubmed.ncbi.nlm.nih.gov/20412080/)

Keywords: phenotype, genetics, hereditary disease, human disease, adult human, ontology, annotation, disease, clinical

Availability: Free, Freely available

Resource Name: Human Phenotype Ontology Annotations

Resource ID: SCR_006219

Alternate IDs: nlx_151835

Old URLs: <http://www.human-phenotype-ontology.org/contao/index.php/downloads.html>

Record Creation Time: 20220129T080235+0000

Record Last Update: 20260903T234828+0000

Ratings and Alerts

No rating or validation information has been found for Human Phenotype Ontology Annotations.

No alerts have been found for Human Phenotype Ontology Annotations.

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 [T1D](#) .

Oellrich A, et al. (2014) Using association rule mining to determine promising secondary phenotyping hypotheses. Bioinformatics (Oxford, England), 30(12), i52.

Tabach Y, et al. (2013) Human disease locus discovery and mapping to molecular pathways through phylogenetic profiling. Molecular systems biology, 9, 692.