Resource Summary Report

Generated by RRID on Apr 28, 2025

FINDbase Worldwide

RRID:SCR_012744
Type: Tool

Proper Citation

FINDbase Worldwide (RRID:SCR_012744)

Resource Information

URL: http://www.findbase.org

Proper Citation: FINDbase Worldwide (RRID:SCR_012744)

Description: FINDbase Worldwide is an online repository of information about the frequency of different mutations leading to inherited disorders in various populations around the globe. Frequency data about 32 disorders, 25 genes within 98 populations covering 1226 mutations is now available. 28 curators worldwide contributed to this database containing data from 37 submissions.

Synonyms: FINDbase

Resource Type: database, data or information resource

Keywords: genetic disorder, human mutation, inherited disorder, mutation pathogenesis, bio.tools

Funding:

Resource Name: FINDbase Worldwide

Resource ID: SCR_012744

Alternate IDs: biotools:findbase, nif-0000-02838

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

Record Creation Time: 20220129T080312+0000

Record Last Update: 20250428T053718+0000

Ratings and Alerts

No rating or validation information has been found for FINDbase Worldwide.

No alerts have been found for FINDbase Worldwide.

Data and Source Information

Source: SciCrunch Registry

Usage and Citation Metrics

We found 12 mentions in open access literature.

Listed below are recent publications. The full list is available at RRID.

Giardine BM, et al. (2021) Clinically relevant updates of the HbVar database of human hemoglobin variants and thalassemia mutations. Nucleic acids research, 49(D1), D1192.

Vihinen M, et al. (2018) Systematics for types and effects of DNA variations. BMC genomics, 19(1), 974.

Viennas E, et al. (2017) Expanded national database collection and data coverage in the FINDbase worldwide database for clinically relevant genomic variation allele frequencies. Nucleic acids research, 45(D1), D846.

Riso B, et al. (2017) Ethical sharing of health data in online platforms - which values should be considered? Life sciences, society and policy, 13(1), 12.

Galperin MY, et al. (2017) The 24th annual Nucleic Acids Research database issue: a look back and upcoming changes. Nucleic acids research, 45(D1), D1.

Mitropoulos K, et al. (2015) Success stories in genomic medicine from resource-limited countries. Human genomics, 9(1), 11.

Giardine B, et al. (2014) Updates of the HbVar database of human hemoglobin variants and thalassemia mutations. Nucleic acids research, 42(Database issue), D1063.

Potamias G, et al. (2014) Deciphering next-generation pharmacogenomics: an information technology perspective. Open biology, 4(7).

Piel FB, et al. (2013) Online biomedical resources for malaria-related red cell disorders. Human mutation, 34(7), 937.

Patrinos GP, et al. (2011) Recommendations for genetic variation data capture in developing countries to ensure a comprehensive worldwide data collection. Human mutation, 32(1), 2.

van Baal S, et al. (2010) ETHNOS : A versatile electronic tool for the development and curation of national genetic databases. Human genomics, 4(5), 361.

Patrinos GP, et al. (2009) A new scientific journal linked to a genetic database: towards a novel publication modality. Human genomics and proteomics : HGP, 2009.