

Resource Summary Report

Generated by [RRID](#) on Sep 10, 2026

PEDHUNTER

RRID:SCR_002031

Type: Tool

Proper Citation

PEDHUNTER (RRID:SCR_002031)

Resource Information

URL: <http://www.ncbi.nlm.nih.gov/CBBresearch/Schaffer/pedhunter.html>

Proper Citation: PEDHUNTER (RRID:SCR_002031)

Description: Software package that facilitates creation and verification of pedigrees within large genealogies. The pedigrees are produced as files in LINKAGE format ready for linkage analysis and for drawing with a variety of drawing programs, such as PEDDRAW and cranefoot.

Abbreviations: PedHunter

Resource Type: software application, software resource

Defining Citation: [PMID:20433770](#), [PMID:9521925](#)

Keywords: gene, genetic, genomic, genealogy, pedigree, bio.tools

Availability: THIS RESOURCE IS NO LONGER IN SERVICE

Resource Name: PEDHUNTER

Resource ID: SCR_002031

Alternate IDs: biotools:pedhunter, OMICS_00211, nlx_154518

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

Record Creation Time: 20220129T080211+0000

Record Last Update: 20260905T133225+0000

Ratings and Alerts

No rating or validation information has been found for PEDHUNTER.

No alerts have been found for PEDHUNTER.

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 [RRID](#) .

Finke K, et al. (2021) Ancestral haplotype reconstruction in endogamous populations using identity-by-descent. PLoS computational biology, 17(2), e1008638.

Lee WJ, et al. (2010) PedHunter 2.0 and its usage to characterize the founder structure of the Old Order Amish of Lancaster County. BMC medical genetics, 11, 68.