

Resource Summary Report

Generated by [ASWG](#) on Aug 4, 2026

GENEHUNTER-PLUS

RRID:SCR_009193

Type: Tool

Proper Citation

GENEHUNTER-PLUS (RRID:SCR_009193)

Resource Information

URL: <http://galton.uchicago.edu/genehunterplus>

Proper Citation: GENEHUNTER-PLUS (RRID:SCR_009193)

Description: Software application that is a modification of the GENEHUNTER software package which produces output files containing the null and conditional distributions of the test statistic (in nullprobs.dat and probs.dat, respectively). These files can then be used as input to the ASM program which incorporates the allele sharing modeling for lodscores and likelihood ratio tests as developed by Kong and Cox (1997). (entry from Genetic Analysis Software)

Abbreviations: GENEHUNTER-PLUS

Synonyms: GENEHUNTER-MODSCORE, GENEHUNTER-TWOLOCUS, ALLEGRO, GENEHUNTER, GENEHUNTER-IMPRINTING

Resource Type: software resource, software application

Keywords: gene, genetic, genomic, c

Resource Name: GENEHUNTER-PLUS

Resource ID: SCR_009193

Alternate IDs: nlx_154334

Record Creation Time: 20220129T080251+0000

Record Last Update: 20260801T191057+0000

Ratings and Alerts

No rating or validation information has been found for GENEHUNTER-PLUS.

No alerts have been found for GENEHUNTER-PLUS.

Data and Source Information

Source: [SciCrunch Registry](#)

Usage and Citation Metrics

We found 1 mentions in open access literature.

Listed below are recent publications. The full list is available at [ASWG](#) .

Simpson CL, et al. (2011) Dissecting the genetic heterogeneity of myopia susceptibility in an Ashkenazi Jewish population using ordered subset analysis. Molecular vision, 17, 1641.