

# Resource Summary Report

Generated by SPARC SAWG on Sep 14, 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 application, software resource

**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:** 20260912T200241+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 [SPARC SAWG](#) .

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.