## **Resource Summary Report**

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# **GENEHUNTER-PLUS**

RRID:SCR\_009193 Type: Tool

**Proper Citation** 

GENEHUNTER-PLUS (RRID:SCR\_009193)

#### **Resource Information**

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

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**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

Funding:

**Resource Name: GENEHUNTER-PLUS** 

Resource ID: SCR\_009193

Alternate IDs: nlx\_154334

**Record Creation Time:** 20220129T080251+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: <u>SciCrunch Registry</u>

#### **Usage and Citation Metrics**

We found 1 mentions in open access literature.

Listed below are recent publications. The full list is available at <u>dkNET</u>.

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.