

# Resource Summary Report

Generated by [kravitz2](#) on Sep 4, 2026

## GWASELECT

RRID:SCR\_013303

Type: Tool

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### Proper Citation

GWASELECT (RRID:SCR\_013303)

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### Resource Information

**URL:** <http://dlin.web.unc.edu/software/gwaselect-2/>

**Proper Citation:** GWASELECT (RRID:SCR\_013303)

**Description:** Software application that implements a novel variable selection method for GWAS data and is able to handle more than half million SNPs. Extensive simulation studies and real data analysis show that this method enjoys high power and low false discovery rate compared to existing variable selection methods. The variables selected by GWASelect can be readily placed into a logistic regression model for disease prediction. The current release is designed for binary outcome under the additive mode of inheritance. (entry from Genetic Analysis Software)

**Abbreviations:** GWASELECT

**Resource Type:** software application, software resource

**Keywords:** gene, genetic, genomic

**Resource Name:** GWASELECT

**Resource ID:** SCR\_013303

**Alternate IDs:** nlx\_154370

**Record Creation Time:** 20220129T080315+0000

**Record Last Update:** 20260904T000445+0000

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### Ratings and Alerts

No rating or validation information has been found for GWASELECT.

No alerts have been found for GWASELECT.

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## Data and Source Information

**Source:** [SciCrunch Registry](#)

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## Usage and Citation Metrics

We have not found any literature mentions for this resource.