

# Resource Summary Report

Generated by [NIF](#) on Sep 2, 2026

## EAGLET

RRID:SCR\_009166

Type: Tool

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### Proper Citation

EAGLET (RRID:SCR\_009166)

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### Resource Information

**URL:** <http://www.columbia.edu/~ws2267/SOFT/EAGLET/eaglet.html>

**Proper Citation:** EAGLET (RRID:SCR\_009166)

**Description:** THIS RESOURCE IS NO LONGER IN SERVICE, documented September 22, 2016. Software package that provides a number of improved statistics for detecting linkage and estimating trait location. EAGLET uses multiple subsamples of dense SNP data to detect linkage with increased power, and to construct sharp 95% confidence intervals for the true trait location.

**Abbreviations:** EAGLET

**Synonyms:** Efficient Analysis of Genetic Linkage: Testing and Estimation

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

**Keywords:** gene, genetic, genomic, perl, c, linux, macos

**Availability:** THIS RESOURCE IS NO LONGER IN SERVICE

**Resource Name:** EAGLET

**Resource ID:** SCR\_009166

**Alternate IDs:** nlx\_154288

**Record Creation Time:** 20220129T080251+0000

**Record Last Update:** 20260829T183107+0000

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### Ratings and Alerts

No rating or validation information has been found for EAGLET.

No alerts have been found for EAGLET.

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## Data and Source Information

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

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## Usage and Citation Metrics

We found 1 mentions in open access literature.

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

Stewart WC, et al. (2014) Next-generation linkage and association methods applied to hypertension: a multifaceted approach to the analysis of sequence data. BMC proceedings, 8(Suppl 1 Genetic Analysis Workshop 18Vanessa Olmo), S111.