

Resource Summary Report

Generated by [nidm-terms](#) on Jul 28, 2026

EAGLET

RRID:SCR_009166

Type: Tool

Proper Citation

EAGLET (RRID:SCR_009166)

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: 20260727T040453+0000

Ratings and Alerts

No rating or validation information has been found for EAGLET.

No alerts have been found for EAGLET.

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 [nidm-terms](#) .

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.