

Resource Summary Report

Generated by RRID on Aug 4, 2026

SEGPATH

RRID:SCR_009052

Type: Tool

Proper Citation

SEGPATH (RRID:SCR_009052)

Resource Information

URL: <https://github.com/gaow/genetic-analysis-software/blob/master/pages/SEGPATH.md>

Proper Citation: SEGPATH (RRID:SCR_009052)

Description: THIS RESOURCE IS NO LONGER IN SERVICE, documented September 29, 2016. Software for segregation and pathway analysis.

Abbreviations: SEGPATH

Synonyms: SEGregation analysis and PATH analysis

Resource Type: software resource, software application

Keywords: gene, genetic, genomic

Availability: THIS RESOURCE IS NO LONGER IN SERVICE

Resource Name: SEGPATH

Resource ID: SCR_009052

Alternate IDs: nlx_154030

Old URLs: <http://www.biostat.wustl.edu/~mike/segpath/>

Record Creation Time: 20220129T080250+0000

Record Last Update: 20260801T191043+0000

Ratings and Alerts

No rating or validation information has been found for SEGPATH.

No alerts have been found for SEGPATH.

Data and Source Information

Source: [SciCrunch Registry](#)

Usage and Citation Metrics

We found 2 mentions in open access literature.

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

Kraja AT, et al. (2012) Obesity-insulin targeted genes in the 3p26-25 region in human studies and LG/J and SM/J mice. Metabolism: clinical and experimental, 61(8), 1129.

Almasy L, et al. (2005) Software for quantitative trait analysis. Human genomics, 2(3), 191.