

# Resource Summary Report

Generated by [dkNET](#) on Apr 17, 2025

## SEGPATH

RRID:SCR\_009052

Type: Tool

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### Proper Citation

SEGPATH (RRID:SCR\_009052)

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### 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

**Funding:**

**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:** 20250416T063535+0000

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## Ratings and Alerts

No rating or validation information has been found for SEGPATH.

No alerts have been found for SEGPATH.

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## Data and Source Information

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

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## Usage and Citation Metrics

We found 2 mentions in open access literature.

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

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.