

Resource Summary Report

Generated by SPARC SAWG on Aug 9, 2026

MAPMAKER/SIBS

RRID:SCR_008012

Type: Tool

Proper Citation

MAPMAKER/SIBS (RRID:SCR_008012)

Resource Information

URL: <https://github.com/gaow/genetic-analysis-software/blob/master/pages/MAPMAKER%26SIBS.md>

Proper Citation: MAPMAKER/SIBS (RRID:SCR_008012)

Description: THIS RESOURCE IS NO LONGER IN SERVICE, documented September 22, 2016. Data analysis software for complete multipoint analysis.

Abbreviations: MAPMAKER/SIBS

Synonyms: GENEHUNTER

Resource Type: software resource, software application

Keywords: gene, genetic, genomic, c, unix

Availability: THIS RESOURCE IS NO LONGER IN SERVICE

Resource Name: MAPMAKER/SIBS

Resource ID: SCR_008012

Alternate IDs: nlx_154465

Alternate URLs:

https://dsgweb.wustl.edu/aldi/software/manuals/mapmaker_sibs/mapmaker_sibs.pdf

Old URLs: <ftp://ftp-genome.wi.mit.edu/distribution/software/sibs>

Record Creation Time: 20220129T080245+0000

Record Last Update: 20260808T190500+0000

Ratings and Alerts

No rating or validation information has been found for MAPMAKER/SIBS.

No alerts have been found for MAPMAKER/SIBS.

Data and Source Information

Source: [SciCrunch Registry](#)

Usage and Citation Metrics

We found 6 mentions in open access literature.

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

Liu AY, et al. (2014) Genome-wide linkage and regional association study of obesity-related phenotypes: the GenSalt study. *Obesity* (Silver Spring, Md.), 22(2), 545.

Li C, et al. (2014) A gene-based analysis of variants in the serum/glucocorticoid regulated kinase (SGK) genes with blood pressure responses to sodium intake: the GenSalt Study. *PloS one*, 9(5), e98432.

Margaritte-Jeannin P, et al. (2007) On the choice of linkage statistics. *BMC proceedings*, 1 Suppl 1(Suppl 1), S102.

Pankratz N, et al. (2007) A two-stage classification approach identifies seven susceptibility genes for a simulated complex disease. *BMC proceedings*, 1 Suppl 1(Suppl 1), S30.

Wapenaar MC, et al. (2004) The interferon gamma gene in celiac disease: augmented expression correlates with tissue damage but no evidence for genetic susceptibility. *Journal of autoimmunity*, 23(2), 183.

Norman RA, et al. (1998) Autosomal genomic scan for loci linked to obesity and energy metabolism in Pima Indians. *American journal of human genetics*, 62(3), 659.