Resource Summary Report

Generated by RRID on Aug 4, 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 SERVCE, 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: 20260801T191053+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 RRID .

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.