

Resource Summary Report

Generated by [RRID](#) on Sep 10, 2026

SIBLINK

RRID:SCR_009381

Type: Tool

Proper Citation

SIBLINK (RRID:SCR_009381)

Resource Information

URL: <http://dmpi.duke.edu/siblink-v-30>

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Description: Software application that allows the user to perform multipoint linkage analysis based on estimated IBD sharing between affected sibpairs. IBD sharing is inferred from IBS status, given marker genotypes, frequencies, and locations. Resulting LOD scores are maximized across a grid of possible disease locations and IBD sharing vectors. (entry from Genetic Analysis Software)

Resource Type: software application, software resource

Keywords: gene, genetic, genomic, c++, unix, solaris, linux, bio.tools

Resource Name: SIBLINK

Resource ID: SCR_009381

Alternate IDs: biotools:siblink, nlx_154616

Alternate URLs: <https://bio.tools/siblink>

Old URLs: <http://wwwchg.duhs.duke.edu/software/siblink.html>

Record Creation Time: 20220129T080252+0000

Record Last Update: 20260905T133251+0000

Ratings and Alerts

No rating or validation information has been found for SIBLINK.

No alerts have been found for SIBLINK.

Data and Source Information

Source: [SciCrunch Registry](#)

Usage and Citation Metrics

We have not found any literature mentions for this resource.