

# Resource Summary Report

Generated by SPARC SAWG on Aug 8, 2026

## SIBLINK

RRID:SCR\_009381

Type: Tool

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### Proper Citation

SIBLINK (RRID:SCR\_009381)

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### Resource Information

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

**Proper Citation:** SIBLINK (RRID:SCR\_009381)

**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 resource, software application

**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:** 20260808T042657+0000

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### Ratings and Alerts

No rating or validation information has been found for SIBLINK.

No alerts have been found for SIBLINK.

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## Data and Source Information

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

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## Usage and Citation Metrics

We have not found any literature mentions for this resource.