

Resource Summary Report

Generated by [RRID](#) on Aug 5, 2026

SIBSIM

RRID:SCR_009113

Type: Tool

Proper Citation

SIBSIM (RRID:SCR_009113)

Resource Information

URL: <http://www.imbs.uni-luebeck.de/pub/sibsim/index.html>

Proper Citation: SIBSIM (RRID:SCR_009113)

Description: THIS RESOURCE IS NO LONGER IN SERVICE, documented September 22, 2016. Software application that simulates either genotype and/or quantitative phenotype data in family structures in a modern, easy to use and highly scalable way. It is based on XML, completely written in C++ and published under the GNU General Public License.

Abbreviations: SIBSIM

Resource Type: software application, software resource

Keywords: gene, genetic, genomic, c++, xml, unix

Availability: THIS RESOURCE IS NO LONGER IN SERVICE

Resource Name: SIBSIM

Resource ID: SCR_009113

Alternate IDs: nlx_154212

Record Creation Time: 20220129T080251+0000

Record Last Update: 20260805T044219+0000

Ratings and Alerts

No rating or validation information has been found for SIBSIM.

No alerts have been found for SIBSIM.

Data and Source Information

Source: [SciCrunch Registry](#)

Usage and Citation Metrics

We have not found any literature mentions for this resource.