

Resource Summary Report

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

SIBERROR

RRID:SCR_009380

Type: Tool

Proper Citation

SIBERROR (RRID:SCR_009380)

Resource Information

URL: <http://gaow.github.io/genetic-analysis-software/s-1.html#siberror>

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Description: Software application that identifies pedigree errors in sibship data. Examples include half siblings, unrelated individuals, identical twins, and parental exclusions. The test statistic is based on the summation of the number of alleles shared by a pair of relatives for a large number of markers and the number of alleles and allele frequencies for those markers. (entry from Genetic Analysis Software)

Synonyms: SibError

Resource Type: software application, software resource

Keywords: gene, genetic, genomic, c, unix

Resource Name: SIBERROR

Resource ID: SCR_009380

Alternate IDs: nlx_154615

Record Creation Time: 20220129T080252+0000

Record Last Update: 20260805T044221+0000

Ratings and Alerts

No rating or validation information has been found for SIBERROR.

No alerts have been found for SIBERROR.

Data and Source Information

Source: [SciCrunch Registry](#)

Usage and Citation Metrics

We have not found any literature mentions for this resource.