

# Resource Summary Report

Generated by PRECISE-TBI on Sep 8, 2026

## SILCLOD

RRID:SCR\_009383

Type: Tool

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### Proper Citation

SILCLOD (RRID:SCR\_009383)

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### Resource Information

**URL:** <http://imbs-luebeck.de/imbs/de/node/34>

**Proper Citation:** SILCLOD (RRID:SCR\_009383)

**Description:** Software application to calculate nominal significance levels and critical LOD scores depending on the length of the investigated region, number of chromosomes, and the cross-over rate. The global significance level as well as the precision of the calculation have to be specified. (entry from Genetic Analysis Software)

**Synonyms:** Significance Levels and Critical LODs

**Resource Type:** software application, software resource

**Keywords:** gene, genetic, genomic

**Resource Name:** SILCLOD

**Resource ID:** SCR\_009383

**Alternate IDs:** nlx\_154619

**Old URLs:** <http://www.imbs.uni-luebeck.de/pub/silcLOD/index.html>

**Record Creation Time:** 20220129T080252+0000

**Record Last Update:** 20260905T133251+0000

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### Ratings and Alerts

No rating or validation information has been found for SILCLOD.

No alerts have been found for SILCLOD.

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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.