

# Resource Summary Report

Generated by [NIF](#) on Aug 14, 2026

## MINSAGE

RRID:SCR\_009293

Type: Tool

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### Proper Citation

MINSAGE (RRID:SCR\_009293)

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### Resource Information

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

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**Description:** Software application to calculate the sample size of genotypes minimally required to ensure that all alleles with a specified frequency at one locus are detected with a given confidence (entry from Genetic Analysis Software)

**Abbreviations:** MINSAGE

**Synonyms:** MINimal SAmple size for GEnotypes

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

**Keywords:** gene, genetic, genomic, ms-windows

**Resource Name:** MINSAGE

**Resource ID:** SCR\_009293

**Alternate IDs:** nlx\_154484

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

**Record Last Update:** 20260813T054954+0000

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### Ratings and Alerts

No rating or validation information has been found for MINSAGE.

No alerts have been found for MINSAGE.

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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.