# **Resource Summary Report**

Generated by FDI Lab - SciCrunch.org on Apr 23, 2025

# **MINSAGE**

RRID:SCR\_009293

Type: Tool

## **Proper Citation**

MINSAGE (RRID:SCR\_009293)

#### **Resource Information**

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

Proper Citation: MINSAGE (RRID:SCR\_009293)

**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

**Funding:** 

Resource Name: MINSAGE

Resource ID: SCR 009293

Alternate IDs: nlx\_154484

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

**Record Last Update:** 20250421T053724+0000

### **Ratings and Alerts**

No rating or validation information has been found for MINSAGE.

No alerts have been found for MINSAGE.

## **Data and Source Information**

Source: SciCrunch Registry

# **Usage and Citation Metrics**

We have not found any literature mentions for this resource.