

# Resource Summary Report

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## MENDELSOFT

RRID:SCR\_013177

Type: Tool

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### Proper Citation

MENDELSOFT (RRID:SCR\_013177)

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### Resource Information

**URL:** <http://www7.inra.fr/mia/T/MendelSoft/>

**Proper Citation:** MENDELSOFT (RRID:SCR\_013177)

**Description:** Software application for identifying all Mendelian inconsistencies in complex pedigree data with thousand of individuals, including many loops and several errors. Can also infer missing genotypes. (entry from Genetic Analysis Software)

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

**Keywords:** gene, genetic, genomic, c++, unix, linux/cygwin, ms-windows, macos

**Funding:**

**Resource Name:** MENDELSOFT

**Resource ID:** SCR\_013177

**Alternate IDs:** nlx\_154474

**Record Creation Time:** 20220129T080314+0000

**Record Last Update:** 20250416T063638+0000

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### Ratings and Alerts

No rating or validation information has been found for MENDELSOFT.

No alerts have been found for MENDELSOFT.

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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.