

Resource Summary Report

Generated by [dkNET](#) on Apr 17, 2025

MENDELISOFT

RRID:SCR_013177

Type: Tool

Proper Citation

MENDELISOFT (RRID:SCR_013177)

Resource Information

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

Proper Citation: MENDELISOFT (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: MENDELISOFT

Resource ID: SCR_013177

Alternate IDs: nlx_154474

Record Creation Time: 20220129T080314+0000

Record Last Update: 20250416T063638+0000

Ratings and Alerts

No rating or validation information has been found for MENDELISOFT.

No alerts have been found for MENDELISOFT.

Data and Source Information

Source: [SciCrunch Registry](#)

Usage and Citation Metrics

We have not found any literature mentions for this resource.