

Resource Summary Report

Generated by PRECISE-TBI on Aug 1, 2026

FINESSE

RRID:SCR_009100

Type: Tool

Proper Citation

FINESSE (RRID:SCR_009100)

Resource Information

URL: http://www.molecular-haplotype.org/finesse/finesse_intro.htm

Proper Citation: FINESSE (RRID:SCR_009100)

Description: THIS RESOURCE IS NO LONGER IN SERVICE, documented September 22, 2016. Software application to incorporate regressive models into the VITESSE likelihood engine.

Abbreviations: FINESSE

Synonyms: VITESSE

Resource Type: software resource, software application

Keywords: gene, genetic, genomic, linux, unix, solaris

Availability: THIS RESOURCE IS NO LONGER IN SERVICE

Resource Name: FINESSE

Resource ID: SCR_009100

Alternate IDs: nlx_154193

Record Creation Time: 20220129T080251+0000

Record Last Update: 20260731T042810+0000

Ratings and Alerts

No rating or validation information has been found for FINESSE.

No alerts have been found for FINESSE.

Data and Source Information

Source: [SciCrunch Registry](#)

Usage and Citation Metrics

We have not found any literature mentions for this resource.