

# Resource Summary Report

Generated by NIF on Sep 5, 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](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 application, software resource

**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:** 20260904T000430+0000

### Ratings and Alerts

No rating or validation information has been found for FINESSE.

No alerts have been found for FINESSE.

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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.