

Resource Summary Report

Generated by [nidm-terms](#) on Jul 28, 2026

INSEGT

RRID:SCR_013126

Type: Tool

Proper Citation

INSEGT (RRID:SCR_013126)

Resource Information

URL: <http://www.stat.washington.edu/thompson/Genepi/InSegT.shtml>

Proper Citation: INSEGT (RRID:SCR_013126)

Description: Software application that constructs feasible haplotype configurations and the corresponding segregation types on pedigrees. the haplotype configuration minimizes recombinations on the pedigree. (entry from Genetic Analysis Software)

Synonyms: INference of SEGregation Types PANGAEA

Resource Type: software application, software resource

Keywords: gene, genetic, genomic, c++, (the standard template library, stl, is used)

Resource Name: INSEGT

Resource ID: SCR_013126

Alternate IDs: nlx_154412

Record Creation Time: 20220129T080314+0000

Record Last Update: 20260727T040550+0000

Ratings and Alerts

No rating or validation information has been found for INSEGT.

No alerts have been found for INSEGT.

Data and Source Information

Source: [SciCrunch Registry](#)

Usage and Citation Metrics

We have not found any literature mentions for this resource.