

Resource Summary Report

Generated by [RRID](#) on Aug 5, 2026

TREELD

RRID:SCR_009111

Type: Tool

Proper Citation

TREELD (RRID:SCR_009111)

Resource Information

URL: <http://pritch.bsd.uchicago.edu/treeld.html>

Proper Citation: TREELD (RRID:SCR_009111)

Description: Free software tool for mapping complex trait loci. TreeLD performs a multipoint LD-analysis by inferring the ancestry of a genomic region and analyzing this ancestry for signals of disease mutations. The generated likelihoods can be used to test for the presence of a disease locus and to fine-map its location, providing a point estimate and a credible region. Furthermore, the package provides a novel way of visualizing the association signal in a sample. TreeLD is designed for high-density SNP haplotypes and can be applied to case-control data, TDT trio data and quantitative trait data. (entry from Genetic Analysis Software)

Abbreviations: TREELD

Resource Type: software application, software resource

Keywords: gene, genetic, genomic, unix, solaris, linux, ms-windows, (95/98/nt/2000/xp)

Availability: Free

Resource Name: TREELD

Resource ID: SCR_009111

Alternate IDs: nlx_154210

Record Creation Time: 20220129T080251+0000

Record Last Update: 20260805T044217+0000

Ratings and Alerts

No rating or validation information has been found for TREELD.

No alerts have been found for TREELD.

Data and Source Information

Source: [SciCrunch Registry](#)

Usage and Citation Metrics

We have not found any literature mentions for this resource.