

Resource Summary Report

Generated by [dkNET](#) on Aug 14, 2026

GRONLOD

RRID:SCR_009049

Type: Tool

Proper Citation

GRONLOD (RRID:SCR_009049)

Resource Information

URL: <https://github.com/gaow/genetic-analysis-software/blob/master/pages/GRONLOD.md>

Proper Citation: GRONLOD (RRID:SCR_009049)

Description: Conversion programs from LINKAGE files are available. The program uses peeling and can employ nested conditioning. There is an automatic peeling program that will unravel (multiple) loops. Alleles do not need to be recoded, so real allele sizes can be used. Genotype probabilities for a chosen person can be calculated for purposes of genetic risk calculation. Later versions include one for calculations using linked markers and mutations and mosaicism, made by Martin van der Meulen. A symbolic versions will generate the formula to compute the pedigree likelihood. (entry from Genetic Analysis Software)

Abbreviations: GRONLOD

Synonyms: GRONingen university LOR score calculation

Resource Type: software resource, software application

Keywords: gene, genetic, genomic, prolog, ms-windows, (3.1/95/nt)

Resource Name: GRONLOD

Resource ID: SCR_009049

Alternate IDs: nlx_154024

Record Creation Time: 20220129T080250+0000

Record Last Update: 20260813T054950+0000

Ratings and Alerts

No rating or validation information has been found for GRONLOD.

No alerts have been found for GRONLOD.

Data and Source Information

Source: [SciCrunch Registry](#)

Usage and Citation Metrics

We have not found any literature mentions for this resource.