

Resource Summary Report

Generated by [dkNET](#) on Aug 8, 2026

[mlgt](#)

RRID:SCR_001211

Type: Tool

Proper Citation

mlgt (RRID:SCR_001211)

Resource Information

URL: <http://cran.r-project.org/web/packages/mlgt/index.html>

Proper Citation: mlgt (RRID:SCR_001211)

Description: Software for processing and analysis of high throughput (Roche 454) sequences generated from multiple loci and multiple biological samples. Sequences are assigned to their locus and sample of origin, aligned and trimmed. Where possible, genotypes are called and variants mapped to known alleles.

Abbreviations: mlgt

Synonyms: Multi-Locus Geno-Typing, mlgt: Multi-Locus Geno-Typing

Resource Type: software resource, software application, data processing software

Keywords: roche, windows, os x, genotype, variant, allele, high throughput sequencing, locus, bio.tools

Availability: THIS RESOURCE IS NO LONGER IN SERVICE

Resource Name: mlgt

Resource ID: SCR_001211

Alternate IDs: BioTools:mlgt, OMICS_02131, biotools:mlgt

Alternate URLs: <https://bio.tools/mlgt>, <https://bio.tools/mlgt>, <https://bio.tools/mlgt>

Record Creation Time: 20220129T080206+0000

Record Last Update: 20260808T190307+0000

Ratings and Alerts

No rating or validation information has been found for mlgt.

No alerts have been found for mlgt.

Data and Source Information

Source: [SciCrunch Registry](#)

Usage and Citation Metrics

We have not found any literature mentions for this resource.