

# Resource Summary Report

Generated by [SPARC SAWG](#) on Aug 9, 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.