

# Resource Summary Report

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## MGA-MAPF2

RRID:SCR\_008108

Type: Tool

### Proper Citation

MGA-MAPF2 (RRID:SCR\_008108)

### Resource Information

**URL:** <http://mga.bionet.nsc.ru/soft/mgamapf2/>

**Proper Citation:** MGA-MAPF2 (RRID:SCR\_008108)

**Description:** Software application that map QTLs in F-2 intercross in model organisms (entry from Genetic Analysis Software)

**Abbreviations:** MGA-MAPF2

**Resource Type:** software application, software resource

**Keywords:** gene, genetic, genomic, fortran, ms-dos, ms-windows, (95/98/nt)

**Resource Name:** MGA-MAPF2

**Resource ID:** SCR\_008108

**Alternate IDs:** nlx\_154478

**Record Creation Time:** 20220129T080245+0000

**Record Last Update:** 20260829T183100+0000

### Ratings and Alerts

No rating or validation information has been found for MGA-MAPF2.

No alerts have been found for MGA-MAPF2.

### Data and Source Information

**Source:** [SciCrunch Registry](#)

## Usage and Citation Metrics

We have not found any literature mentions for this resource.