

Resource Summary Report

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MAP/MAP+/MAP+H/MAP2000

RRID:SCR_009272

Type: Tool

Proper Citation

MAP/MAP+/MAP+H/MAP2000 (RRID:SCR_009272)

Resource Information

URL: <http://cedar.genetics.soton.ac.uk/pub/PROGRAMS/map>

Proper Citation: MAP/MAP+/MAP+H/MAP2000 (RRID:SCR_009272)

Description: Software application for multiple pairwise linkage analysis under interference (entry from Genetic Analysis Software)

Abbreviations: MAP/MAP+/MAP+H/MAP2000

Synonyms: (MAP+H is the radiation Hybrid module of MAP+)

Resource Type: software application, software resource

Keywords: gene, genetic, genomic, fortran, unix, sunos

Resource Name: MAP/MAP+/MAP+H/MAP2000

Resource ID: SCR_009272

Alternate IDs: nlx_154453

Record Creation Time: 20220129T080252+0000

Record Last Update: 20260905T133248+0000

Ratings and Alerts

No rating or validation information has been found for MAP/MAP+/MAP+H/MAP2000.

No alerts have been found for MAP/MAP+/MAP+H/MAP2000.

Data and Source Information

Source: [SciCrunch Registry](#)

Usage and Citation Metrics

We have not found any literature mentions for this resource.