

# Resource Summary Report

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## HAPMIXMAP

RRID:SCR\_006066

Type: Tool

### Proper Citation

HAPMIXMAP (RRID:SCR\_006066)

### Resource Information

**URL:** [http://www.homepages.ed.ac.uk/pmckeigu/hapmixmap/hapmixmap\\_manual.html](http://www.homepages.ed.ac.uk/pmckeigu/hapmixmap/hapmixmap_manual.html)

**Proper Citation:** HAPMIXMAP (RRID:SCR\_006066)

**Description:** Software application for modelling extended haplotypes in genetic association studies, similar to the FASTPHASE program. (entry from Genetic Analysis Software)

**Abbreviations:** HAPMIXMAP

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

**Keywords:** gene, genetic, genomic

**Resource Name:** HAPMIXMAP

**Resource ID:** SCR\_006066

**Alternate IDs:** nlx\_154391

**Record Creation Time:** 20220129T080234+0000

**Record Last Update:** 20260810T040421+0000

### Ratings and Alerts

No rating or validation information has been found for HAPMIXMAP.

No alerts have been found for HAPMIXMAP.

### Data and Source Information

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

## Usage and Citation Metrics

We have not found any literature mentions for this resource.