MINIMAC
RRID:SCR_009292
Type: Tool

Proper Citation
MINIMAC (RRID:SCR_009292)

Resource Information

URL: http://genome.sph.umich.edu/wiki/Minimac

Description: Software application that is a low memory, computationally efficient implementation of the MaCH algorithm for genotype imputation. It is designed to work on phased genotypes and can handle very large reference panels with hundreds or thousands of haplotypes. The name has two parts. The first, mini, refers to the modest amount of computational resources it requires. The second, mac, is short hand for MaCH, our widely used algorithm for genotype imputation. (entry from Genetic Analysis Software)

Resource Name: MINIMAC

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Resource Type: Resource, software resource, software application

Keywords: gene, genetic, genomic

Resource ID: SCR_009292

Website Status: Last checked up

Alternate IDs: nlx_154483

Abbreviations: MINIMAC

Mentions Count: 85

Ratings and Alerts
No rating or validation information has been found for MINIMAC.

No alerts have been found for MINIMAC.

### Data and Source Information

**Source:** SciCrunch Registry

### Usage and Citation Metrics

We found 85 mentions in open access literature.

**Listed below are recent publications.** The full list is available at [FDI Lab - SciCrunch Infrastructure](FDI Lab - SciCrunch Infrastructure).


