Resource Summary Report

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HAPLOTYPE ESTIMATION

RRID:SCR_009231 Type: Tool

Proper Citation

HAPLOTYPE ESTIMATION (RRID:SCR_009231)

Resource Information

URL: http://www.bioinf.mdc-berlin.de/projects/hap/

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Description: Haplotype estimation service available in two variants: one for genetic data from unrelated probands (e.g., case/control studies) and one for core family data (entry from Genetic Analysis Software)

Abbreviations: HAPLOTYPE ESTIMATION

Resource Type: software resource, software application

Keywords: gene, genetic, genomic

Funding:

Resource Name: HAPLOTYPE ESTIMATION

Resource ID: SCR_009231

Alternate IDs: nlx_154388

Record Creation Time: 20220129T080251+0000

Record Last Update: 20250429T055316+0000

Ratings and Alerts

No rating or validation information has been found for HAPLOTYPE ESTIMATION.

No alerts have been found for HAPLOTYPE ESTIMATION.

Data and Source Information

Source: <u>SciCrunch Registry</u>

Usage and Citation Metrics

We have not found any literature mentions for this resource.