## **Resource Summary Report**

Generated by RRID on May 13, 2025

# **EMI**

RRID:SCR\_001171 Type: Tool

**Proper Citation** 

EMI (RRID:SCR\_001171)

#### **Resource Information**

URL: http://cs.au.dk/~qianyuxx/EMI/

Proper Citation: EMI (RRID:SCR\_001171)

**Description:** Clustering software which uses pairwise identity-by-descent (IBD) segments to infer multiple-haplotype IBD clusters. It expands clusters from seed haplotypes by adding qualified neighbors and extends clusters across sliding windows in the genome.

Abbreviations: EMI

Synonyms: Efficient Multiple-IBD

Resource Type: software resource

Defining Citation: PMID:24363374

Keywords: clustering, identity-by-descent

Funding:

Availability: GNU General Public License

Resource Name: EMI

Resource ID: SCR\_001171

Alternate IDs: OMICS\_02186

**Record Creation Time:** 20220129T080206+0000

Record Last Update: 20250420T014022+0000

## **Ratings and Alerts**

No rating or validation information has been found for EMI.

No alerts have been found for EMI.

Data and Source Information

Source: SciCrunch Registry

## **Usage and Citation Metrics**

We have not found any literature mentions for this resource.