

Resource Summary Report

Generated by [NIF](#) on Aug 7, 2026

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

Availability: THIS RESOURCE IS NO LONGER IN SERVICE

Resource Name: EMI

Resource ID: SCR_001171

Alternate IDs: OMICS_02186

Record Creation Time: 20220129T080206+0000

Record Last Update: 20260801T190133+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.