# **Resource Summary Report**

Generated by RRID on May 19, 2025

# **EM-DECODER**

RRID:SCR\_000023

Type: Tool

### **Proper Citation**

EM-DECODER (RRID:SCR\_000023)

#### **Resource Information**

URL: http://www.people.fas.harvard.edu/~junliu/em/em.htm

**Proper Citation:** EM-DECODER (RRID:SCR\_000023)

**Description:** A haplotype inference program.

**Abbreviations:** EM-DECODER

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

Keywords: gene, genetic, genomic

Funding:

Resource Name: EM-DECODER

Resource ID: SCR\_000023

Alternate IDs: nlx\_154297

**Record Creation Time:** 20220129T080159+0000

**Record Last Update:** 20250519T204844+0000

### **Ratings and Alerts**

No rating or validation information has been found for EM-DECODER.

No alerts have been found for EM-DECODER.

## Data and Source Information

Source: SciCrunch Registry

## **Usage and Citation Metrics**

We found 1 mentions in open access literature.

Listed below are recent publications. The full list is available at RRID.

Weale ME, et al. (2004) A survey of current software for haplotype phase inference. Human genomics, 1(2), 141.