

Resource Summary Report

Generated by [nidm-terms](#) on Sep 23, 2026

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: THIS RESOURCE IS NO LONGER IN SERVICE. Documented on July 31,2025. A haplotype inference program.

Abbreviations: EM-DECODER

Resource Type: software application, software resource

Keywords: gene, genetic, genomic

Availability: THIS RESOURCE IS NO LONGER IN SERVICE

Resource Name: EM-DECODER

Resource ID: SCR_000023

Alternate IDs: nlx_154297

Record Creation Time: 20220129T080159+0000

Record Last Update: 20260919T195740+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 [nidm-terms](#) .

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