

Resource Summary Report

Generated by NIF on Apr 20, 2025

ANCESTRY

RRID:SCR_009121

Type: Tool

Proper Citation

ANCESTRY (RRID:SCR_009121)

Resource Information

URL: <http://ciil.case.edu:3000>

Proper Citation: ANCESTRY (RRID:SCR_009121)

Description: THIS RESOURCE IS NO LONGER IN SERVICE, documented September 22, 2016. Software application that provides an easy to use web-based user interface to compute individual ancestry estimates using the Maximum Likelihood Estimation method. This website also allows registered users to manage genotype and marker/allele frequency files. Registration is optional.

Abbreviations: ANCESTRY

Resource Type: software application, software resource

Keywords: gene, genetic, genomic, ruby on rails, fortran, web-based

Funding:

Availability: THIS RESOURCE IS NO LONGER IN SERVICE

Resource Name: ANCESTRY

Resource ID: SCR_009121

Alternate IDs: nlx_154224

Record Creation Time: 20220129T080251+0000

Record Last Update: 20250420T015744+0000

Ratings and Alerts

No rating or validation information has been found for ANCESTRY.

No alerts have been found for ANCESTRY.

Data and Source Information

Source: [SciCrunch Registry](#)

Usage and Citation Metrics

We found 2 mentions in open access literature.

Listed below are recent publications. The full list is available at [NIF](#).

Foltz SM, et al. (2024) Somatic mutation phasing and haplotype extension using linked-reads in multiple myeloma. bioRxiv : the preprint server for biology.

Edge MD, et al. (2020) Attacks on genetic privacy via uploads to genealogical databases. eLife, 9.