

Resource Summary Report

Generated by [NIF](#) on Sep 17, 2026

APE

RRID:SCR_009122

Type: Tool

Proper Citation

APE (RRID:SCR_009122)

Resource Information

URL: <https://github.com/gaow/genetic-analysis-software/blob/master/pages/APE.md>

Proper Citation: APE (RRID:SCR_009122)

Description: THIS RESOURCE IS NO LONGER IN SERVICE, documented September 22, 2016. Software application that extends partially observed genotype data to the whole pedigree. Can be used for generating starting points for MCMC samplers and for checking that the genotype data are consistent with the pedigree structure.

Abbreviations: APE

Synonyms: Allelic Path Explorer

Resource Type: software application, software resource

Keywords: gene, genetic, genomic, c, unix

Availability: THIS RESOURCE IS NO LONGER IN SERVICE

Resource Name: APE

Resource ID: SCR_009122

Alternate IDs: nlx_154227

Old URLs: <http://www.helsinki.fi/~mpirinen/download>

Record Creation Time: 20220129T080251+0000

Record Last Update: 20260912T200240+0000

Ratings and Alerts

No rating or validation information has been found for APE.

No alerts have been found for APE.

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](#) .

Ghafari M, et al. (2021) A mechanistic evolutionary model explains the time-dependent pattern of substitution rates in viruses. Current biology : CB, 31(21), 4689.

Charvet CJ, et al. (2017) Combining diffusion magnetic resonance tractography with stereology highlights increased cross-cortical integration in primates. The Journal of comparative neurology, 525(5), 1075.