

Resource Summary Report

Generated by [nidm-terms](#) on Jul 29, 2026

2SNP

RRID:SCR_009038

Type: Tool

Proper Citation

2SNP (RRID:SCR_009038)

Resource Information

URL: <https://github.com/gaow/genetic-analysis-software/blob/master/pages/2SNP.md>

Proper Citation: 2SNP (RRID:SCR_009038)

Description: THIS RESOURCE IS NO LONGER IN SERVICE, documented September 22, 2016. An algorithm resource for scalable phasing method for trios and unrelated individuals.

Abbreviations: 2SNP

Resource Type: software application, software resource

Defining Citation: [PMID:18451440](#)

Keywords: gene, genetic, genomic

Availability: THIS RESOURCE IS NO LONGER IN SERVICE

Resource Name: 2SNP

Resource ID: SCR_009038

Alternate IDs: nlx_154003

Alternate URLs: <https://omictools.com/2snp-tool>

Old URLs: <http://alla.cs.gsu.edu/~software/2SNP/>

Record Creation Time: 20220129T080250+0000

Record Last Update: 20260727T040443+0000

Ratings and Alerts

No rating or validation information has been found for 2SNP.

No alerts have been found for 2SNP.

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 [nidm-terms](#) .

Xu Y, et al. (2012) WinHAP: an efficient haplotype phasing algorithm based on scalable sliding windows. PloS one, 7(8), e43163.

He Y, et al. (2011) Accelerating haplotype-based genome-wide association study using perfect phylogeny and phase-known reference data. PloS one, 6(7), e22097.