

# Resource Summary Report

Generated by [NIF](#) on Sep 3, 2026

## 2SNP

RRID:SCR\_009038

Type: Tool

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### Proper Citation

2SNP (RRID:SCR\_009038)

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### 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:** 20260904T000428+0000

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### Ratings and Alerts

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

No alerts have been found for 2SNP.

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## Data and Source Information

**Source:** [SciCrunch Registry](#)

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## Usage and Citation Metrics

We found 2 mentions in open access literature.

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

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.