

Resource Summary Report

Generated by [SPARC SAWG](#) on Sep 14, 2026

SNPFILE

RRID:SCR_009402

Type: Tool

Proper Citation

SNPFILE (RRID:SCR_009402)

Resource Information

URL: <http://www.daimi.au.dk/%7Emailund/SNPFile/>

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Description: Software library and API for manipulating large SNP datasets with associated meta-data, such as marker names, marker locations, individuals' phenotypes, etc. in an I/O efficient binary file format. In its core, SNPFile assumes very little about the metadata associated with markers and individuals, but leaves this up to application program protocols. (entry from Genetic Analysis Software)

Resource Type: software application, software library, software resource, software toolkit

Keywords: gene, genetic, genomic, c++, linux, unix

Resource Name: SNPFILE

Resource ID: SCR_009402

Alternate IDs: nlx_154641

Record Creation Time: 20220129T080252+0000

Record Last Update: 20260912T200014+0000

Ratings and Alerts

No rating or validation information has been found for SNPFILE.

No alerts have been found for SNPFILE.

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 [SPARC SAWG](#) .

Nielsen J, et al. (2008) SNPFile--a software library and file format for large scale association mapping and population genetics studies. BMC bioinformatics, 9, 526.