

Resource Summary Report

Generated by SPARC SAWG on Sep 17, 2026

SNPiR

RRID:SCR_000557

Type: Tool

Proper Citation

SNPiR (RRID:SCR_000557)

Resource Information

URL: <http://lilab.stanford.edu/SNPiR/>

Proper Citation: SNPiR (RRID:SCR_000557)

Description: Software for reliable Identification of Genomic Variants Using RNA-seq Data.

Abbreviations: SNPiR

Synonyms: SNPiR: Reliable Identification of Genomic Variants Using RNA-seq Data

Resource Type: software resource

Keywords: genomic variant, rna-seq

Availability: THIS RESOURCE IS NO LONGER IN SERVICE

Resource Name: SNPiR

Resource ID: SCR_000557

Alternate IDs: OMICS_01362

Record Creation Time: 20220129T080202+0000

Record Last Update: 20260912T195511+0000

Ratings and Alerts

No rating or validation information has been found for SNPiR.

No alerts have been found for SNPiR.

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](#) .

Mei??ner T, et al. (2015) OncoRep: an n-of-1 reporting tool to support genome-guided treatment for breast cancer patients using RNA-sequencing. BMC medical genomics, 8, 24.