

Resource Summary Report

Generated by [SPARC SAWG](#) on Jul 28, 2026

[npstat](#)

RRID:SCR_012128

Type: Tool

Proper Citation

npstat (RRID:SCR_012128)

Resource Information

URL: <http://code.google.com/p/npstat/>

Proper Citation: npstat (RRID:SCR_012128)

Description: Software that implements some population genetics tests and estimators that can be applied to pooled sequences from Next Generation Sequencing experiments.

Resource Type: software resource

Defining Citation: [PMID:24102736](#)

Keywords: standalone software

Resource Name: npstat

Resource ID: SCR_012128

Alternate IDs: OMICS_05826

Record Creation Time: 20220129T080308+0000

Record Last Update: 20260725T190732+0000

Ratings and Alerts

No rating or validation information has been found for npstat.

No alerts have been found for npstat.

Data and Source Information

Source: [SciCrunch Registry](#)

Usage and Citation Metrics

We found 3 mentions in open access literature.

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

Ergon ??, et al. (2022) Identification of loci controlling timing of stem elongation in red clover using genotyping by sequencing of pooled phenotypic extremes. Molecular genetics and genomics : MGG, 297(6), 1587.

Garlovsky MD, et al. (2020) Within-population sperm competition intensity does not predict asymmetry in conpopulation sperm precedence. Philosophical transactions of the Royal Society of London. Series B, Biological sciences, 375(1813), 20200071.

Khan S, et al. (2020) A Whole Genome Re-Sequencing Based GWA Analysis Reveals Candidate Genes Associated with Ivermectin Resistance in Haemonchus contortus. Genes, 11(4).