

Resource Summary Report

Generated by [FDI Lab - SciCrunch.org](#) on Apr 15, 2025

[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

Funding:

Resource Name: npstat

Resource ID: SCR_012128

Alternate IDs: OMICS_05826

Record Creation Time: 20220129T080308+0000

Record Last Update: 20250410T070230+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 [FDI Lab - SciCrunch.org](#).

Ergon Å, 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.

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).

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.