

Resource Summary Report

Generated by T1D on Aug 2, 2026

SASQUANT

RRID:SCR_013122

Type: Tool

Proper Citation

SASQUANT (RRID:SCR_013122)

Resource Information

URL: <http://cuke.hort.ncsu.edu/cucurbit/wehner/software.html>

Proper Citation: SASQUANT (RRID:SCR_013122)

Description: THIS RESOURCE IS NO LONGER IN SERVICE. Documented on May 24,2023. SAS software program to estimate genetic effects and heritabilities of quantitative traits in breeding populations consisting of six related generations (entry from Genetic Analysis Software)

Resource Type: software resource, software application

Keywords: gene, genetic, genomic, sas, bio.tools

Availability: THIS RESOURCE IS NO LONGER IN SERVICE

Resource Name: SASQUANT

Resource ID: SCR_013122

Alternate IDs: nlx_154610, biotools:sasquant

Alternate URLs: <https://bio.tools/sasquant>

Record Creation Time: 20220129T080314+0000

Record Last Update: 20260801T191103+0000

Ratings and Alerts

No rating or validation information has been found for SASQUANT.

No alerts have been found for SASQUANT.

Data and Source Information

Source: [SciCrunch Registry](#)

Usage and Citation Metrics

We found 4 mentions in open access literature.

Listed below are recent publications. The full list is available at [T1D](#) .

Khan MMH, et al. (2021) AMMI and GGE biplot analysis for yield performance and stability assessment of selected Bambara groundnut (*Vigna subterranea* L. Verdc.) genotypes under the multi-environmental trials (METs). Scientific reports, 11(1), 22791.

González AM, et al. (2017) Dissection of Resistance Genes to *Pseudomonas syringae* pv. *phaseolicola* in UI3 Common Bean Cultivar. International journal of molecular sciences, 18(12).

Dia M, et al. (2016) Stability of fruit quality traits in diverse watermelon cultivars tested in multiple environments. Horticulture research, 3, 16066.

Baettig OM, et al. (2016) Comprehensive characterization of ligand-induced plasticity changes in a dimeric enzyme. The FEBS journal, 283(16), 3029.