## **Resource Summary Report**

Generated by NIF on May 3, 2025

# **FESTA**

RRID:SCR\_009081

Type: Tool

### **Proper Citation**

FESTA (RRID:SCR\_009081)

### **Resource Information**

URL: http://www.sph.umich.edu/csg/qin/FESTA/

**Proper Citation:** FESTA (RRID:SCR\_009081)

**Description:** Software application (entry from Genetic Analysis Software)

**Abbreviations:** FESTA

**Synonyms:** Fragmented Exhaustive Seearch for TAgsnps

Resource Type: software resource, software application

Keywords: gene, genetic, genomic

Funding:

Resource Name: FESTA

Resource ID: SCR\_009081

Alternate IDs: nlx\_154092

**Record Creation Time:** 20220129T080251+0000

**Record Last Update:** 20250503T060054+0000

## **Ratings and Alerts**

No rating or validation information has been found for FESTA.

#### Data and Source Information

Source: SciCrunch Registry

## **Usage and Citation Metrics**

We found 6 mentions in open access literature.

**Listed below are recent publications.** The full list is available at NIF.

Hasegawa Y, et al. (2021) The relationship between bone density and the oral function in older adults: a cross-sectional observational study. BMC geriatrics, 21(1), 591.

Rago A, et al. (2020) Sex biased expression and co-expression networks in development, using the hymenopteran Nasonia vitripennis. PLoS genetics, 16(1), e1008518.

Uwitonze A, et al. (2017) Connectivity Restoration in Wireless Sensor Networks via Space Network Coding. Sensors (Basel, Switzerland), 17(4).

Punt M, et al. (2014) Clinimetric properties of a novel feedback device for assessing gait parameters in stroke survivors. Journal of neuroengineering and rehabilitation, 11, 30.

Badke YM, et al. (2013) Methods of tagSNP selection and other variables affecting imputation accuracy in swine. BMC genetics, 14, 8.

Gualdrón Duarte JL, et al. (2013) Genotype imputation accuracy in a F2 pig population using high density and low density SNP panels. BMC genetics, 14, 38.