

Resource Summary Report

Generated by [Kravitz](#) on Sep 10, 2026

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 Search for TAGsnps

Resource Type: software application, software resource

Keywords: gene, genetic, genomic

Resource Name: FESTA

Resource ID: SCR_009081

Alternate IDs: nlx_154092

Record Creation Time: 20220129T080251+0000

Record Last Update: 20260905T133243+0000

Ratings and Alerts

No rating or validation information has been found for FESTA.

No alerts have 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 [Kravitz](#) .

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.

Gualdr??n 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.

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