

Resource Summary Report

Generated by [kravitz2](#) on Jul 31, 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 resource, software application

Keywords: gene, genetic, genomic

Resource Name: FESTA

Resource ID: SCR_009081

Alternate IDs: nlx_154092

Record Creation Time: 20220129T080251+0000

Record Last Update: 20260731T042806+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 [kravitz2](#) .

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.