Resource Summary Report

Generated by dkNET on Apr 16, 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: 20250416T063536+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 dkNET.

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.