Resource Summary Report

Generated by RRID on Apr 16, 2025

SNIPPEEP

RRID:SCR_013309
Type: Tool

Proper Citation

SNIPPEEP (RRID:SCR_013309)

Resource Information

URL: http://snippeep.sourceforge.net/

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Description: Software application that is an interactive graphic interface to visualise results from whole genome genotyping. It allows one to visualise single subjects and groups of subjects, and provides a direct connection with the UCSC Genome Browser. (entry from Genetic Analysis Software)

Resource Type: software resource, software application

Keywords: gene, genetic, genomic, c

Funding:

Resource Name: SNIPPEEP

Resource ID: SCR_013309

Alternate IDs: nlx_154019

Record Creation Time: 20220129T080315+0000

Record Last Update: 20250416T063639+0000

Ratings and Alerts

No rating or validation information has been found for SNIPPEEP.

No alerts have been found for SNIPPEEP.

Data and Source Information

Source: SciCrunch Registry

Usage and Citation Metrics

We found 3 mentions in open access literature.

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

Leblond CS, et al. (2019) Both rare and common genetic variants contribute to autism in the Faroe Islands. NPJ genomic medicine, 4, 1.

Leblond CS, et al. (2014) Meta-analysis of SHANK Mutations in Autism Spectrum Disorders: a gradient of severity in cognitive impairments. PLoS genetics, 10(9), e1004580.

Leblond CS, et al. (2012) Genetic and functional analyses of SHANK2 mutations suggest a multiple hit model of autism spectrum disorders. PLoS genetics, 8(2), e1002521.