Resource Summary Report

Generated by dkNET on Apr 17, 2025

SNAP 3

RRID:SCR_009400

Type: Tool

Proper Citation

SNAP 3 (RRID:SCR_009400)

Resource Information

URL: http://capella.uni-kiel.de/snap/snap.htm

Proper Citation: SNAP 3 (RRID:SCR_009400)

Description: THIS RESOURCE IS NO LONGER IN SERVICE, documented September 29, 2016. Software program can be used to generate SNP haplotype sequence data of unrelated individuals and nuclear families with a fixed or random number of children.

Synonyms: Simulation of sNp haplotype data And Phenotypic traits

Resource Type: software resource, software application

Keywords: gene, genetic, genomic, c, linux, unix, solaris, ms-windows

Funding:

Availability: THIS RESOURCE IS NO LONGER IN SERVICE

Resource Name: SNAP 3

Resource ID: SCR_009400

Alternate IDs: nlx_154639

Old URLs: http://capella.uni-kiel.de/snap/snap.htm,

Record Creation Time: 20220129T080252+0000

Record Last Update: 20250416T063546+0000

Ratings and Alerts

No rating or validation information has been found for SNAP 3.

No alerts have been found for SNAP 3.

Data and Source Information

Source: SciCrunch Registry

Usage and Citation Metrics

We found 1 mentions in open access literature.

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

Wang L, et al. (2022) The Capparis spinosa var. herbacea genome provides the first genomic instrument for a diversity and evolution study of the Capparaceae family. GigaScience, 11.