

Resource Summary Report

Generated by [nidm-terms](#) on Sep 21, 2026

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 application, software resource

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

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: 20260919T195806+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 [nidm-terms](#) .

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.