

Resource Summary Report

Generated by [dkNET](#) on Aug 25, 2026

SNP HITLINK

RRID:SCR_013340

Type: Tool

Proper Citation

SNP HITLINK (RRID:SCR_013340)

Resource Information

URL: <http://www.dynacom.co.jp/u-tokyo.ac.jp/snphitlink/>

Proper Citation: SNP HITLINK (RRID:SCR_013340)

Description: Software program providing a useful pipeline to directly connect SNP data and linkage analysis program. SNP HiTLink currently supports the data from SNP chips provided by Affymetrix (Mapping 100k/500k array set, Genome-Wide Human SNP array 5.0/6.0) and Illumina (recently supported), carrying out typical linkage analysis programs of MLINK (FASTLINK/ LINKAGE package), Superlink, Merlin and Allegro. (entry from Genetic Analysis Software)

Abbreviations: SNP HITLINK

Synonyms: SNP High-Throughput LINKage analysis system

Resource Type: software application, software resource

Keywords: gene, genetic, genomic, bio.tools

Resource Name: SNP HITLINK

Resource ID: SCR_013340

Alternate IDs: nlx_154644, biotools:snp_hitlink

Alternate URLs: https://bio.tools/snp_hitlink

Record Creation Time: 20220129T080315+0000

Record Last Update: 20260821T194007+0000

Ratings and Alerts

No rating or validation information has been found for SNP HITLINK.

No alerts have been found for SNP HITLINK.

Data and Source Information

Source: [SciCrunch Registry](#)

Usage and Citation Metrics

We have not found any literature mentions for this resource.