# **Resource Summary Report**

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### **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/

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**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 resource, software application

Keywords: gene, genetic, genomic, bio.tools

**Funding:** 

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:** 20250416T063640+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.