Resource Summary Report

Generated by dkNET on Apr 17, 2025

SNPHARVESTER

RRID:SCR_008536
Type: Tool

Proper Citation

SNPHARVESTER (RRID:SCR_008536)

Resource Information

URL: http://bioinformatics.ust.hk/SNPHarvester.html

Proper Citation: SNPHARVESTER (RRID:SCR_008536)

Description: Software tool for detecting epistatic interactions in genome-wide association studies (entry from Genetic Analysis Software)

Abbreviations: SNPHARVESTER

Resource Type: software resource, software application

Keywords: gene, genetic, genomic, java

Funding:

Resource Name: SNPHARVESTER

Resource ID: SCR_008536

Alternate IDs: nlx_154643

Record Creation Time: 20220129T080248+0000

Record Last Update: 20250416T063523+0000

Ratings and Alerts

No rating or validation information has been found for SNPHARVESTER.

No alerts have been found for SNPHARVESTER.

Data and Source Information

Source: SciCrunch Registry

Usage and Citation Metrics

We found 6 mentions in open access literature.

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

Tang DY, et al. (2024) SEEI: spherical evolution with feedback mechanism for identifying epistatic interactions. BMC genomics, 25(1), 462.

Tuo S, et al. (2018) FDHE-IW: A Fast Approach for Detecting High-Order Epistasis in Genome-Wide Case-Control Studies. Genes, 9(9).

Tuo S, et al. (2017) Niche harmony search algorithm for detecting complex disease associated high-order SNP combinations. Scientific reports, 7(1), 11529.

Jiang X, et al. (2015) Learning Predictive Interactions Using Information Gain and Bayesian Network Scoring. PloS one, 10(12), e0143247.

Lee JY, et al. (2013) A SNP Harvester Analysis to Better Detect SNPs of CCDC158 Gene That Are Associated with Carcass Quality Traits in Hanwoo. Asian-Australasian journal of animal sciences, 26(6), 766.

Anunciação O, et al. (2013) Using information interaction to discover epistatic effects in complex diseases. PloS one, 8(10), e76300.