Resource Summary Report

Generated by RRID on Apr 17, 2025

VarySysDB

RRID:SCR_005880
Type: Tool

Proper Citation

VarySysDB (RRID:SCR_005880)

Resource Information

URL: http://h-invitational.jp/varygene/

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Description: It consists of a Genome Browser, an LD Search System, and the VaryGene 2 system. The Generic Genome Browser is a combination of database and interactive Web page for manipulating and displaying annotations on genomes, while LDSearchSystem is a search system for linkage disequilibrium (LD) bins. VaryGene 2 is a system to search, display, and download our research results on human polymorphism based on publicly available data and annotations of transcripts presented by H-InvDB. VaryGene 2 provides information about single nucleotide polymorphisms (SNPs), deletion-insertion polymorphisms (DIPs), short tandem repeats (STRs), single amino acid repeats (SARs), structural variation (or copy number variations: CNVs), and their relations to the genome, transcripts, and functional domains. Users can search by polymorphisms, transcripts, STRs/SARs, and CNVs.

Synonyms: VarySysDB

Resource Type: database, data or information resource

Keywords: genome, human polymorphism

Funding:

Resource Name: VarySysDB

Resource ID: SCR_005880

Alternate IDs: nif-0000-03621

Record Creation Time: 20220129T080233+0000

Record Last Update: 20250412T055010+0000

Ratings and Alerts

No rating or validation information has been found for VarySysDB.

No alerts have been found for VarySysDB.

Data and Source Information

Source: SciCrunch Registry

Usage and Citation Metrics

We have not found any literature mentions for this resource.