Resource Summary Report

Generated by dkNET on Apr 17, 2025

SIMLINK

RRID:SCR_009387
Type: Tool

Proper Citation

SIMLINK (RRID:SCR_009387)

Resource Information

URL: http://csg.sph.umich.edu/boehnke/simlink.php

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Description: Software program to estimate the probability (power) of detecting linkage given family history information on a set of identified pedigrees. (entry from Genetic Analysis Software)

Resource Type: software resource, software application

Keywords: gene, genetic, genomic, fortran, ms-dos, unix, sunos, vms

Funding:

Resource Name: SIMLINK

Resource ID: SCR_009387

Alternate IDs: nlx_154625

Record Creation Time: 20220129T080252+0000

Record Last Update: 20250416T063546+0000

Ratings and Alerts

No rating or validation information has been found for SIMLINK.

No alerts have been found for SIMLINK.

Data and Source Information

Source: SciCrunch Registry

Usage and Citation Metrics

We found 3 mentions in open access literature.

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

Tomberg K, et al. (2018) Whole exome sequencing of ENU-induced thrombosis modifier mutations in the mouse. PLoS genetics, 14(9), e1007658.

Dong Y, et al. (2013) Genome-wide scan for hypertension linkage to chromosome 12q23.1 - q23.3 in a Chinese family. The Indian journal of medical research, 137(5), 935.

Kuchtey J, et al. (2011) Mapping of the disease locus and identification of ADAMTS10 as a candidate gene in a canine model of primary open angle glaucoma. PLoS genetics, 7(2), e1001306.