Resource Summary Report

Generated by RRID on Apr 17, 2025

PDT

RRID:SCR_009319
Type: Tool

Proper Citation

PDT (RRID:SCR_009319)

Resource Information

URL: http://www.chg.duke.edu/software/pdt.html

Proper Citation: PDT (RRID:SCR_009319)

Description: Software analysis program to evaluate evidence of linkage disequilibrium (LD) in general pedigree data. All family data may be used without nullifying the validity of the association test, even when there is more than one affected in a family. The PDT program performs both allele-specific and genotype-specific LD analysis of individual markers. Version 5.1 adds the ability to perform genotype-specific analysis over marker sets. (entry from Genetic Analysis Software)

Abbreviations: PDT

Synonyms: pedigree disequilibrium test

Resource Type: software resource, software application

Keywords: gene, genetic, genomic, unix, solaris

Funding:

Resource Name: PDT

Resource ID: SCR_009319

Alternate IDs: nlx_154513

Record Creation Time: 20220129T080252+0000

Record Last Update: 20250416T063543+0000

Ratings and Alerts

No rating or validation information has been found for PDT.

No alerts have been found for PDT.

Data and Source Information

Source: SciCrunch Registry

Usage and Citation Metrics

We found 2 mentions in open access literature.

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

Voyiaziakis E, et al. (2011) Association of SLC6A4 variants with obsessive-compulsive disorder in a large multicenter US family study. Molecular psychiatry, 16(1), 108.

Elston RC, et al. (2004) A review of the 'Statistical Analysis for Genetic Epidemiology' (S.A.G.E.) software package. Human genomics, 1(6), 456.