Resource Summary Report

Generated by NIF on Apr 20, 2025

EPDT

RRID:SCR_009172

Type: Tool

Proper Citation

EPDT (RRID:SCR_009172)

Resource Information

URL: http://www.ibms.sinica.edu.tw/~csjfann/first%20flow/programlist.htm

Proper Citation: EPDT (RRID:SCR_009172)

Description: THIS RESOURCE IS NO LONGER IN SERVICE. Documented on May 16,2023. Software application that for detecting linkage/disequilibrium signals # between genetic markers and disease loci, particularly if only one # or a few large pedigrees are available. The strategy differs from # conventional approaches that require at least a moderate number of # families to attain adequate statistical power. The proposed testing # procedure is advantageous in that it provides high statistical power # coupled with reduced sample collection. Furthermore, the proposed # method avoids problems such as potential population stratification # and genetic heterogeneity, and is robust with respect to misspecification # of phenotype. (entry from Genetic Analysis Software)

Abbreviations: EPDT

Synonyms: Extended Pedigree Disequilibrium Test

Resource Type: software application, software resource

Keywords: gene, genetic, genomic, unix

Funding:

Availability: THIS RESOURCE IS NO LONGER IN SERVICE

Resource Name: EPDT

Resource ID: SCR 009172

Alternate IDs: nlx_154301

Record Creation Time: 20220129T080251+0000

Record Last Update: 20250420T015747+0000

Ratings and Alerts

No rating or validation information has been found for EPDT.

No alerts have been found for EPDT.

Data and Source Information

Source: SciCrunch Registry

Usage and Citation Metrics

We found 1 mentions in open access literature.

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

Martin KC, et al. (2024) Functional partitioning of sentence processing and emotional prosody in the right perisylvian cortex after perinatal stroke. Scientific reports, 14(1), 28602.