

Resource Summary Report

Generated by [NIF](#) on Sep 19, 2026

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

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: 20260912T200241+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.