

Resource Summary Report

Generated by T1D on Aug 2, 2026

TDASP

RRID:SCR_004943

Type: Tool

Proper Citation

TDASP (RRID:SCR_004943)

Resource Information

URL: <http://biostatistics.mdanderson.org/SoftwareDownload/>

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Description: Software application for power and sample-size calculations for the TDT and ASP tests under a wide variety of ascertainment schemes. Uses the flexible genetic model of McGinnis. Most calculations are exact rather than asymptotic. (entry from Genetic Analysis Software)

Abbreviations: TDASP

Synonyms: Power and Sample-Size Calculations for the TDT and ASP Tests

Resource Type: software resource, software application

Keywords: gene, genetic, genomic, fortran95, unix, ms-windows

Resource Name: TDASP

Resource ID: SCR_004943

Alternate IDs: nlx_154675

Record Creation Time: 20220129T080227+0000

Record Last Update: 20260801T191050+0000

Ratings and Alerts

No rating or validation information has been found for TDASP.

No alerts have been found for TDASP.

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 [T1D](#) .

Zhu D, et al. (2019) CACNA1C (rs1006737) may be a susceptibility gene for schizophrenia: An updated meta-analysis. Brain and behavior, 9(6), e01292.

Wittenburg D, et al. (2018) An approximate Bayesian significance test for genomic evaluations. Biometrical journal. Biometrische Zeitschrift, 60(6), 1096.

Jacob L, et al. (2016) Evaluation of a multi-arm multi-stage Bayesian design for phase II drug selection trials - an example in hemato-oncology. BMC medical research methodology, 16, 67.