# Resource Summary Report

Generated by T1D on Jul 29, 2026

## DMET-Analyzer

RRID:SCR_002030
Type: Tool

### Proper Citation

DMET-Analyzer (RRID:SCR_002030)

### Resource Information

**URL:** http://sourceforge.net/projects/dmetanalyzer/

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**Description:** Software tool for the automatic association analysis among the variation of the patient genomes and the clinical conditions of patients, i.e. the different response to drugs. The system allows: (i) to automatize the workflow of analysis of DMET (drug metabolism enzymes and transporters)-SNP (Single Nucleotide Polymorphism) data avoiding the use of multiple tools; (ii) the automatic annotation of DMET-SNP data and the search in existing databases of SNPs (e.g. dbSNP), (iii) the association of SNP with pathway through the search in PharmaKGB, a major knowledge base for pharmacogenomic studies. It has a simple graphical user interface that allows users (doctors/biologists) to upload and analyze DMET files produced by Affymetrix DMET-Console in an interactive way.

**Abbreviations:** DMET-Analyzer

**Synonyms:** DMETANALYZER, DMETANALYZER - A tool for supporting pharmacogenomics data analysis

**Resource Type:** software resource

**Defining Citation:** PMID:23035929

**Keywords:** drug, metabolism, enzyme, transporter, affymetrix, variation, genome, clinical, affymetrix dmet, single nucleotide polymorphism, annotation, analysis, pharmacogenomic, pathway

**Availability:** Free, Available for download, Freely available

**Resource Name:** DMET-Analyzer

**Resource ID:** SCR_002030

**Alternate IDs:** OMICS_01920

**Record Creation Time:** 20220129T080211+0000

**Record Last Update:** 20260725T190518+0000

## Ratings and Alerts

No rating or validation information has been found for DMET-Analyzer.

No alerts have been found for DMET-Analyzer.

## 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 T1D .

Scionti F, et al. (2017) Genetic variants associated with Fabry disease progression despite enzyme replacement therapy. Oncotarget, 8(64), 107558.