

Resource Summary Report

Generated by T1D on Aug 9, 2026

SET

RRID:SCR_003605

Type: Tool

Proper Citation

SET (RRID:SCR_003605)

Resource Information

URL: <http://microarray.ym.edu.tw:8080/tools/module/set/index.jsp?mode=home>

Proper Citation: SET (RRID:SCR_003605)

Description: A Java tool to evaluate and visualize the sample discrimination abilities of gene expression signatures. This tool provides a filtration function for signature identification and lies between clinical analyses and class prediction (or feature selection) tools.

Abbreviations: SET

Synonyms: Signature Evaluation Tool, SET - a Java tool to evaluate and visualize the sample discrimination abilities of gene expression signatures, Signature Evaluation Tool (SET), SET (Signature Evaluation Tool), SET (Signature Evaluation Tool) - a Java tool to evaluate and visualize the sample discrimination abilities of gene expression signatures

Resource Type: software resource, software application

Defining Citation: [PMID:18221568](#)

Keywords: java, gene expression, gene, discrimination, candidate gene, microarray

Resource Name: SET

Resource ID: SCR_003605

Alternate IDs: OMICS_02294

Record Creation Time: 20220129T080219+0000

Record Last Update: 20260808T190442+0000

Ratings and Alerts

No rating or validation information has been found for SET.

No alerts have been found for SET.

Data and Source Information

Source: [SciCrunch Registry](#)

Usage and Citation Metrics

We have not found any literature mentions for this resource.