

Resource Summary Report

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Wigwams

RRID:SCR_006400

Type: Tool

Proper Citation

Wigwams (RRID:SCR_006400)

Resource Information

URL:

http://www2.warwick.ac.uk/fac/sci/systemsbiology/staff/ott/tools_and_software/wigwams

Proper Citation: Wigwams (RRID:SCR_006400)

Description: A computational tool for analyzing multiple gene expression time series data sets for the same organism. The goal is to determine if there is evidence for gene regulatory mechanisms that are shared by multiple different expression responses.

Abbreviations: Wigwams

Resource Type: software resource

Defining Citation: [PMID:24351708](#)

Keywords: matlab, gene, gene module, co-regulation, time series, gene expression, differential expression

Resource Name: Wigwams

Resource ID: SCR_006400

Alternate IDs: OMICS_02214

Record Creation Time: 20220129T080236+0000

Record Last Update: 20260725T190619+0000

Ratings and Alerts

No rating or validation information has been found for Wigwams.

No alerts have been found for Wigwams.

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 [nidm-terms](#) .

Zhang N, et al. (2020) Revealing Shared and Distinct Genes Responding to JA and SA Signaling in Arabidopsis by Meta-Analysis. Frontiers in plant science, 11, 908.