

# Resource Summary Report

Generated by [PRECISE-TBI](#) on Sep 8, 2026

## RARTF

RRID:SCR\_013457

Type: Tool

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### Proper Citation

RARTF (RRID:SCR\_013457)

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### Resource Information

**URL:** <http://rarge.psc.riken.jp/rartf/>

**Proper Citation:** RARTF (RRID:SCR\_013457)

**Description:** Database of complete sets of Arabidopsis transcription factors with a variety of information on Arabidopsis thaliana transcription factor families including: full-length cDNA sequences, Ds-tagged mutants, multiple sequences alignments of family members, phylogenetic trees, functional motifs, and so on. In addition, expression profiles of all transcription factor genes are available.

**Abbreviations:** RARTF

**Synonyms:** RARTF: RIKEN Arabidopsis Transcription Factor database, RIKEN Arabidopsis Transcription Factor database

**Resource Type:** data or information resource, database

**Defining Citation:** [PMID:16769687](#)

**Keywords:** transcription factor, protein, gene, microarray, expression profile

**Availability:** Free

**Resource Name:** RARTF

**Resource ID:** SCR\_013457

**Alternate IDs:** OMICS\_00562

**Record Creation Time:** 20220129T080316+0000

**Record Last Update:** 20260905T133207+0000

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### Ratings and Alerts

No rating or validation information has been found for RARTF.

No alerts have been found for RARTF.

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## Data and Source Information

**Source:** [SciCrunch Registry](#)

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## Usage and Citation Metrics

We found 2 mentions in open access literature.

**Listed below are recent publications.** The full list is available at [PRECISE-TBI](#) .

Wirojsirasak W, et al. (2019) Pan- and core- gene association networks: Integrative approaches to understanding biological regulation. PloS one, 14(1), e0210481.

Graf A, et al. (2017) Parallel analysis of Arabidopsis circadian clock mutants reveals different scales of transcriptome and proteome regulation. Open biology, 7(3).