# **Resource Summary Report**

Generated by FDI Lab - SciCrunch.org on May 24, 2025

# **Transcription Regulatory Regions Database**

RRID:SCR\_005723 Type: Tool

# **Proper Citation**

Transcription Regulatory Regions Database (RRID:SCR\_005723)

### **Resource Information**

#### URL: http://www.bionet.nsc.ru/trrd/

**Proper Citation:** Transcription Regulatory Regions Database (RRID:SCR\_005723)

**Description:** TRRD is a unique information resource, accumulating information on structural and functional organization of transcription regulatory regions of eukaryotic genes. Only experimentally confirmed information is included into TRRD. Transcription Regulatory Regions Database (TRRD) is developed for accumulation of experimental information on the structure-function features of regulatory regions of eukaryotic genes. Each entry of TRRD corresponds to a particular gene. The annotated part of an entry includes the structure-function description of gene regulatory regions composed by regulatory units (promoters, silencers, enhancers, etc.), individual transcription factor binding sites that constitute these regulatory units, and transcription factors that bind to these sites. In addition, the entry contains the gene expression patterns and references to original publications.

Synonyms: TRRD

Resource Type: data or information resource, database

Defining Citation: PMID:11752324

Funding:

Resource Name: Transcription Regulatory Regions Database

Resource ID: SCR\_005723

Alternate IDs: nif-0000-03594

Record Creation Time: 20220129T080232+0000

Record Last Update: 20250523T054500+0000

# **Ratings and Alerts**

No rating or validation information has been found for Transcription Regulatory Regions Database.

No alerts have been found for Transcription Regulatory Regions Database.

## Data and Source Information

Source: <u>SciCrunch Registry</u>

## **Usage and Citation Metrics**

We found 3 mentions in open access literature.

Listed below are recent publications. The full list is available at FDI Lab - SciCrunch.org.

He YH, et al. (2016) Improved lipids, diastolic pressure and kidney function are potential contributors to familial longevity: a study on 60 Chinese centenarian families. Scientific reports, 6, 21962.

Coulibaly I, et al. (2008) Bioinformatic tools for inferring functional information from plant microarray data II: Analysis beyond single gene. International journal of plant genomics, 2008, 893941.

Galperin MY, et al. (2005) The Molecular Biology Database Collection: 2005 update. Nucleic acids research, 33(Database issue), D5.