

Resource Summary Report

Generated by [RRID](#) on Aug 10, 2026

Transcription Factor encyclopedia

RRID:SCR_002130

Type: Tool

Proper Citation

Transcription Factor encyclopedia (RRID:SCR_002130)

Resource Information

URL: <http://www.cisreg.ca/tfe>

Proper Citation: Transcription Factor encyclopedia (RRID:SCR_002130)

Description: THIS RESOURCE IS NO LONGER IN SERVICE. Documented on September 23,2022.The TFe provides details of transcription factor binding sites in close collaboration with Pazar, a public database of transcription factor and regulatory sequence information. The long-term goal of TFe is to create an online encyclopedic collection of well-studied transcription factor proteins in the human, mouse, and rat genomes, combining a mixture of both expert-curated and automatically-populated content to provide users with a wide set of information relevant to a transcription factor protein of their interest.

Synonyms: Tfe

Resource Type: database, data or information resource

Defining Citation: [PMID:22458515](#)

Keywords: binding site, regulatory sequence, transcription factor

Availability: THIS RESOURCE IS NO LONGER IN SERVICE

Resource Name: Transcription Factor encyclopedia

Resource ID: SCR_002130

Alternate IDs: nif-0000-20924

Record Creation Time: 20220129T080211+0000

Record Last Update: 20260808T190351+0000

Ratings and Alerts

No rating or validation information has been found for Transcription Factor encyclopedia.

No alerts have been found for Transcription Factor encyclopedia.

Data and Source Information

Source: [SciCrunch Registry](#)

Usage and Citation Metrics

We found 2 mentions in open access literature.

Listed below are recent publications. The full list is available at [RRID](#) .

Gao L, et al. (2019) The expression, significance and function of cancer susceptibility candidate 9 in lung squamous cell carcinoma: A bioinformatics and in vitro investigation. International journal of oncology, 54(5), 1651.

Yusuf D, et al. (2012) The transcription factor encyclopedia. Genome biology, 13(3), R24.