

Resource Summary Report

Generated by [dkNET](#) on Sep 8, 2026

[tfcheckpoint](#)

RRID:SCR_023880

Type: Tool

Proper Citation

tfcheckpoint (RRID:SCR_023880)

Resource Information

URL: <https://www.tfcheckpoint.org/>

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Description: Collection of transcription factors annotated according to experimental and other evidence on their function as true DbTFs. Provides reference for both small scale experiments and genome scale studies. Curated compendium of specific DNA-binding RNA polymerase II transcription factors.

Resource Type: data or information resource, database

Defining Citation: [PMID:23933972](#)

Keywords: transcription factor, DNA-binding RNA polymerase II transcription factors, DNA-binding, RNA polymerase II transcription factors,

Funding: Liaison Committee between the Central Norway Regional Health Authority ; Norwegian Cancer Society ; Norwegian University of Science and Technology

Availability: Free, Freely available

Resource Name: tfcheckpoint

Resource ID: SCR_023880

Record Creation Time: 20230727T050222+0000

Record Last Update: 20260905T133219+0000

Ratings and Alerts

No rating or validation information has been found for tfcheckpoint.

No alerts have been found for tfcheckpoint.

Data and Source Information

Source: [SciCrunch Registry](#)

Usage and Citation Metrics

We found 4 mentions in open access literature.

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

Corvino D, et al. (2025) Type I Interferon Drives a Cellular State Inert to TCR-Stimulation and Could Impede Effective T-Cell Differentiation in Cancer. European journal of immunology, 55(1), e202451371.

Newsham I, et al. (2024) Early detection and diagnosis of cancer with interpretable machine learning to uncover cancer-specific DNA methylation patterns. Biology methods & protocols, 9(1), bpae028.

Sun Y, et al. (2024) Comprehensive analysis of differential long non-coding RNA and messenger RNA expression in cholelithiasis using high-throughput sequencing and bioinformatics. Frontiers in genetics, 15, 1375019.

Bergmann T, et al. (2022) Production of human entorhinal stellate cell-like cells by forward programming shows an important role of Foxp1 in reprogramming. Frontiers in cell and developmental biology, 10, 976549.