

# Resource Summary Report

Generated by [Kravitz](#) on Sep 18, 2026

## TFFM

RRID:SCR\_010888

Type: Tool

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### Proper Citation

TFFM (RRID:SCR\_010888)

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### Resource Information

**URL:** <http://cisreg.cmmmt.ubc.ca/TFFM/doc/>

**Proper Citation:** TFFM (RRID:SCR\_010888)

**Description:** Software for Transcription Factor Flexible Models (TFFMs) that represent Transcription Factor Binding Sites (TFBSs) and are based on hidden Markov models (HMM). They are flexible and are able to model both position interdependence within TFBSs and variable length motifs within a single dedicated framework.

**Abbreviations:** TFFM

**Synonyms:** Transcription Factor Flexible Models

**Resource Type:** software resource

**Defining Citation:** [PMID:24039567](#)

**Keywords:** python

**Availability:** GNU Lesser General Public Licence

**Resource Name:** TFFM

**Resource ID:** SCR\_010888

**Alternate IDs:** OMICS\_00495

**Record Creation Time:** 20220129T080301+0000

**Record Last Update:** 20260912T195725+0000

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### Ratings and Alerts

No rating or validation information has been found for TFFM.

No alerts have been found for TFFM.

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## Data and Source Information

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

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## Usage and Citation Metrics

We found 1 mentions in open access literature.

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

Mathelier A, et al. (2016) JASPAR 2016: a major expansion and update of the open-access database of transcription factor binding profiles. Nucleic acids research, 44(D1), D110.