# **Resource Summary Report**

Generated by FDI Lab - SciCrunch.org on Apr 15, 2025

## **TFFM**

RRID:SCR\_010888

Type: Tool

### **Proper Citation**

TFFM (RRID:SCR\_010888)

#### **Resource Information**

URL: http://cisreg.cmmt.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

**Funding:** 

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:** 20250410T070028+0000

## **Ratings and Alerts**

No rating or validation information has been found for TFFM.

No alerts have been found for TFFM.

#### **Data and Source Information**

Source: SciCrunch Registry

### **Usage and Citation Metrics**

We found 1 mentions in open access literature.

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

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.