

Resource Summary Report

Generated by [NIF](#) on Aug 12, 2026

[TFBSGroup](#)

RRID:SCR_010887

Type: Tool

Proper Citation

TFBSGroup (RRID:SCR_010887)

Resource Information

URL: <http://bioinformatics.bioengr.uic.edu/TFBSGroup/>

Proper Citation: TFBSGroup (RRID:SCR_010887)

Description: THIS RESOURCE IS NO LONGER IN SERVICE. Documented on January 11, 2023. A Software program for Predicting Transcription Factor Binding Sites.

Abbreviations: TFBSGroup

Synonyms: TFBSGroup: A Program for Predicting Transcription Factor Binding Sites

Resource Type: software resource

Keywords: transcription factor binding site

Availability: THIS RESOURCE IS NO LONGER IN SERVICE

Resource Name: TFBSGroup

Resource ID: SCR_010887

Alternate IDs: OMICS_00494

Record Creation Time: 20220129T080301+0000

Record Last Update: 20260808T185939+0000

Ratings and Alerts

No rating or validation information has been found for TFBSGroup.

No alerts have been found for TFBSGroup.

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 [NIF](#) .

Jia C, et al. (2013) A fast weak motif-finding algorithm based on community detection in graphs. BMC bioinformatics, 14, 227.