

Resource Summary Report

Generated by [SPARC SAWG](#) on Aug 9, 2026

TFmodeller

RRID:SCR_015715

Type: Tool

Proper Citation

TFmodeller (RRID:SCR_015715)

Resource Information

URL: <http://www.ccg.unam.mx/tfmodeller>

Proper Citation: TFmodeller (RRID:SCR_015715)

Description: Web application that scans a library of protein-DNA complexes and builds comparative models of proteins bound to DNA. Its results include complex coordinates, schematic interface diagrams, interface alignments and DNA motifs.

Resource Type: web application, sequence analysis software, software resource, software application, data analysis software, data visualization software, data processing software

Defining Citation: [PMID:17459960](#)

Keywords: protein-dna complex, comparative model, modeling software, schematic interface diagram, interface alignment, dna motif, complex coordinate

Funding: NIGMS RO1-GM071962

Availability: Free for academic use, Freely available, Tutorial available

Resource Name: TFmodeller

Resource ID: SCR_015715

Alternate URLs: <http://maya.ccg.unam.mx/~tfmodell/>

License URLs: <http://maya.ccg.unam.mx/~tfmodell/credits.html>

Record Creation Time: 20220129T080327+0000

Record Last Update: 20260808T190025+0000

Ratings and Alerts

No rating or validation information has been found for TFmodeller.

No alerts have been found for TFmodeller.

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 [SPARC SAWG](#) .

Angarica VE, et al. (2008) Prediction of TF target sites based on atomistic models of protein-DNA complexes. BMC bioinformatics, 9, 436.