**PDBTM**

RRID:SCR_011962  
Type: Tool

**Proper Citation**

PDBTM (RRID:SCR_011962)

**Resource Information**

**URL:** http://pdbtm.enzim.hu/

**Proper Citation:** PDBTM (RRID:SCR_011962)

**Description:** Comprehensive and continuously updated transmembrane protein database of the Protein Data Bank (PDB) created by scanning all PDB entries with the TMDET algorithm. Resource for transmembrane proteins and their structures.

**Abbreviations:** PDBTM

**Synonyms:** PDBTM, Protein Data Bank of Transmembrane Proteins, Protein Data Bank of Transmembrane Proteins

**Resource Type:** data or information resource, database

**Defining Citation:** PMID:23203988

**Keywords:** FASEB list

**Funding Agency:** Hungarian Scientific Research Fund, Hungarian Academy of Sciences

**Availability:** Free, Available for download, Freely available, Acknowledgement requested

**Resource Name:** PDBTM

**Resource ID:** SCR_011962

**Alternate IDs:** OMICS_01613

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**Ratings and Alerts**
No rating or validation information has been found for PDBTM.

No alerts have been found for PDBTM.

**Data and Source Information**

**Source:** SciCrunch Registry

**Usage and Citation Metrics**

We found 45 mentions in open access literature.

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


Ding W, et al. (2020) Predicting the Real-Valued Inter-Residue Distances for Proteins. Advanced science (Weinheim, Baden-Wurttemberg, Germany), 7(19), 2001314.

Mesdaghi S, et al. (2020) In silico prediction of structure and function for a large family of transmembrane proteins that includes human Tmem41b. F1000Research, 9, 1395.


