

# Resource Summary Report

Generated by [PRECISE-TBI](#) on Sep 8, 2026

## Orientations of Proteins in Membranes database

RRID:SCR\_011961

Type: Tool

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### Proper Citation

Orientations of Proteins in Membranes database (RRID:SCR\_011961)

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### Resource Information

**URL:** <http://opm.phar.umich.edu/>

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**Description:** Database that provides a collection of transmembrane, monotopic and peripheral proteins from the Protein Data Bank whose spatial arrangements in the lipid bilayer have been calculated theoretically and compared with experimental data. The database allows analysis, sorting and searching of membrane proteins based on their structural classification, species, destination membrane, numbers of transmembrane segments and subunits, numbers of secondary structures and the calculated hydrophobic thickness or tilt angle with respect to the bilayer normal.

**Abbreviations:** OPM

**Synonyms:** Orientations of Proteins in Membranes (OPM) database, OPM Database

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

**Defining Citation:** [PMID:16397007](#)

**Keywords:** protein, membrane, bio.tools, FASEB list

**Funding:** NSF

**Availability:** Acknowledgement requested

**Resource Name:** Orientations of Proteins in Membranes database

**Resource ID:** SCR\_011961

**Alternate IDs:** OMICS\_01612, biotools:opm

**Alternate URLs:** <https://bio.tools/opm>

**Record Creation Time:** 20220129T080307+0000

**Record Last Update:** 20260905T132717+0000

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## Ratings and Alerts

No rating or validation information has been found for Orientations of Proteins in Membranes database.

No alerts have been found for Orientations of Proteins in Membranes database.

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## Data and Source Information

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

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## Usage and Citation Metrics

We found 135 mentions in open access literature.

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

Walker AR, et al. (2026) Differential conservation analysis identifies residues defining constitutive internalization in beta-adrenergic receptors. *iScience*, 29(3), 115033.

Stawarski M, et al. (2026) Functional signatures of de novo GABBR1 and GABBR2 variants associated with neurodevelopmental disorders. *NPJ genomic medicine*, 11(1).

Zhang X, et al. (2026) Mechanistic insights into ligand selectivity in  $\mu$ - and  $\kappa$ -opioid receptors beyond the "message-address" conceptual framework. *Acta pharmaceutica Sinica. B*, 16(3), 1530.

Moreno-Giménez E, et al. (2026) Bioconversion of carotenoids into high-value crocins using a marine sponge carotenoid cleavage dioxygenase. *The New phytologist*, 250(5), 3230.

Zhao Z, et al. (2026) Structure of the human P2X3 receptor reveals the basis for subtype-selective inhibition by sivopixant. *PLoS biology*, 24(4), e3003777.

Morris GP, et al. (2026) A sorghum pangenome reference improves global crop trait discovery. *Nature*, 652(8112), 1245.

Merz M, et al. (2025) Downstream Signaling of Muscarinic M4 Receptors Is Regulated by Receptor Density and Cellular Environment. *Pharmacology research & perspectives*, 13(3), e70123.

van der Velden WJC, et al. (2025) Allosteric communication mechanism in the glucagon receptor. *The Journal of biological chemistry*, 301(6), 108530.

Markwardt F, et al. (2025) Activation of the Rat P2X7 Receptor by Functionally Different ATP Activation Sites. *Cells*, 14(12).

Yamamuro T, et al. (2025) Mitochondria-derived PEP licenses glycerolipid synthesis via SLC25A35. *bioRxiv : the preprint server for biology*.

Farfán-García AE, et al. (2025) Structural and functional analysis of *Escherichia coli* membrane disruption by Ib-M peptides. *PloS one*, 20(10), e0334029.

Vladkova R, et al. (2025) X-Ray Crystal and Cryo-Electron Microscopy Structure Analysis Unravels How the Unique Thylakoid Lipid Composition Is Utilized by Cytochrome b6f for Driving Reversible Proteins' Reorganization During State Transitions. *Membranes*, 15(5).

Romp E, et al. (2025) Modifications in FLAP's second cytosolic loop influence 5-LOX interaction, inhibitor binding, and leukotriene formation. *FEBS letters*, 599(15), 2179.

Clemente-Moragón A, et al. (2025) Pharmacogenomics and chronotherapy of drug-induced cardioprotection in acute myocardial infarction. *Nature communications*, 16(1), 10450.

Guo CR, et al. (2025) Understanding interspecies drug response variations between human and rodent P2X7 receptors. *Nature communications*, 16(1), 10827.

Lin YY, et al. (2024) Finely ordered intracellular domain harbors an allosteric site to modulate physiopathological function of P2X3 receptors. *Nature communications*, 15(1), 7652.

Rullo-Tubau J, et al. (2024) Structure and mechanisms of transport of human Asc1/CD98hc amino acid transporter. *Nature communications*, 15(1), 2986.

Floch A, et al. (2024) Molecular dynamics of the human RhD and RhAG blood group proteins. *Frontiers in chemistry*, 12, 1360392.

Lazzara F, et al. (2024) Anti-angiogenic and antioxidant effects of axitinib in human retinal endothelial cells: implications in diabetic retinopathy. *Frontiers in pharmacology*, 15, 1415846.

de Araújo NS, et al. (2024) Aptamer-Based Recognition of Breast Tumor Cells: A New Era for Breast Cancer Diagnosis. *International journal of molecular sciences*, 25(2).