

Search form



Database of Comparative Protein Structure Models

Welcome to ModBase, a database of three-dimensional protein models calculated by comparative modeling.

General Information

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Note:

MODBASE contains theoretically calculated models, not experimentally determined structures. The models may contain **significant** errors.

ModBase search form

Search

Search type Model(Default)

Display type Model Detail (graphical)

All available datasets are selected

[Select specific dataset\(s\)](#)

Search by properties

Property Database Accession Number

Organism ALL

or

[Advanced search](#)

Users of ModBase are requested to cite this article in their publications:

[MODBASE, a database of annotated comparative protein structure models, and associated resources](#), Ursula Pleper, Narayanan Eswar, Fred Davis, M.S. Madhusudhan, Andrea Rossi, Marc A. Marti-Renom, Rachel Karchin, Ben Webb, David Eramian, Min-Yi Shen, Libusha Kelly, Francisco Melo and Andrej Sali *Nucleic Acids Research* **34**, D291-D295, 2006.

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