

# State of the Art

## AlphaFold Protein Structure Database

Developed by Google DeepMind and EMBL-EBI

Search for protein, gene, UniProt accession or organism or sequence search

BETA

Search

Examples: MENFQKVEKIGEGTYGV... Free fatty acid receptor 2 At1g58602 Q5VSL9 E. coli

See search help →

Go to online course →

See our updates – September 2024

Congratulations to Demis Hassabis, John Jumper and David Baker on winning the 2024 Nobel Prize in Chemistry!



AlphaFold DB provides open access to over 200 million protein structure predictions to accelerate scientific research.

# Text to Text - ChatGPT

What can I help with?



Message ChatGPT



Create image



Summarize text



Brainstorm



Code



Make a plan

More