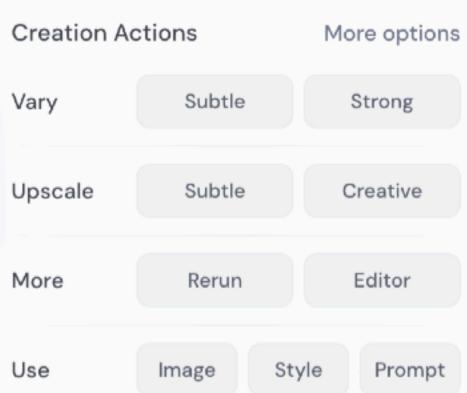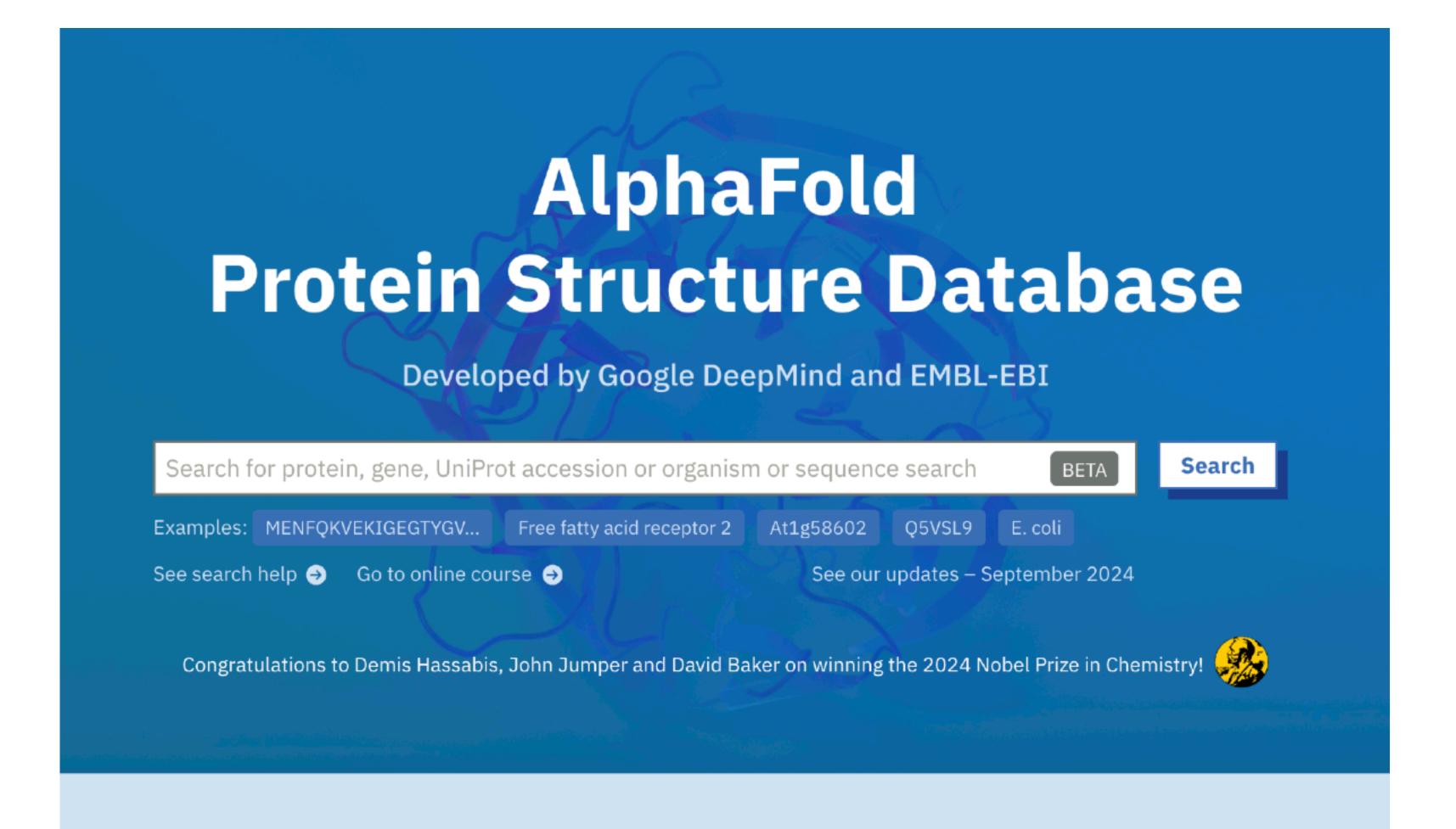Show how diffusion models take a random noisy image and iteratively convert them into a clear image guided by a text prompt

v 6.1

Creation Actions   More options

Vary   Subtle   Strong

Upscale   Subtle   Creative

More   Rerun   Editor

Use   Image   Style   Prompt

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.