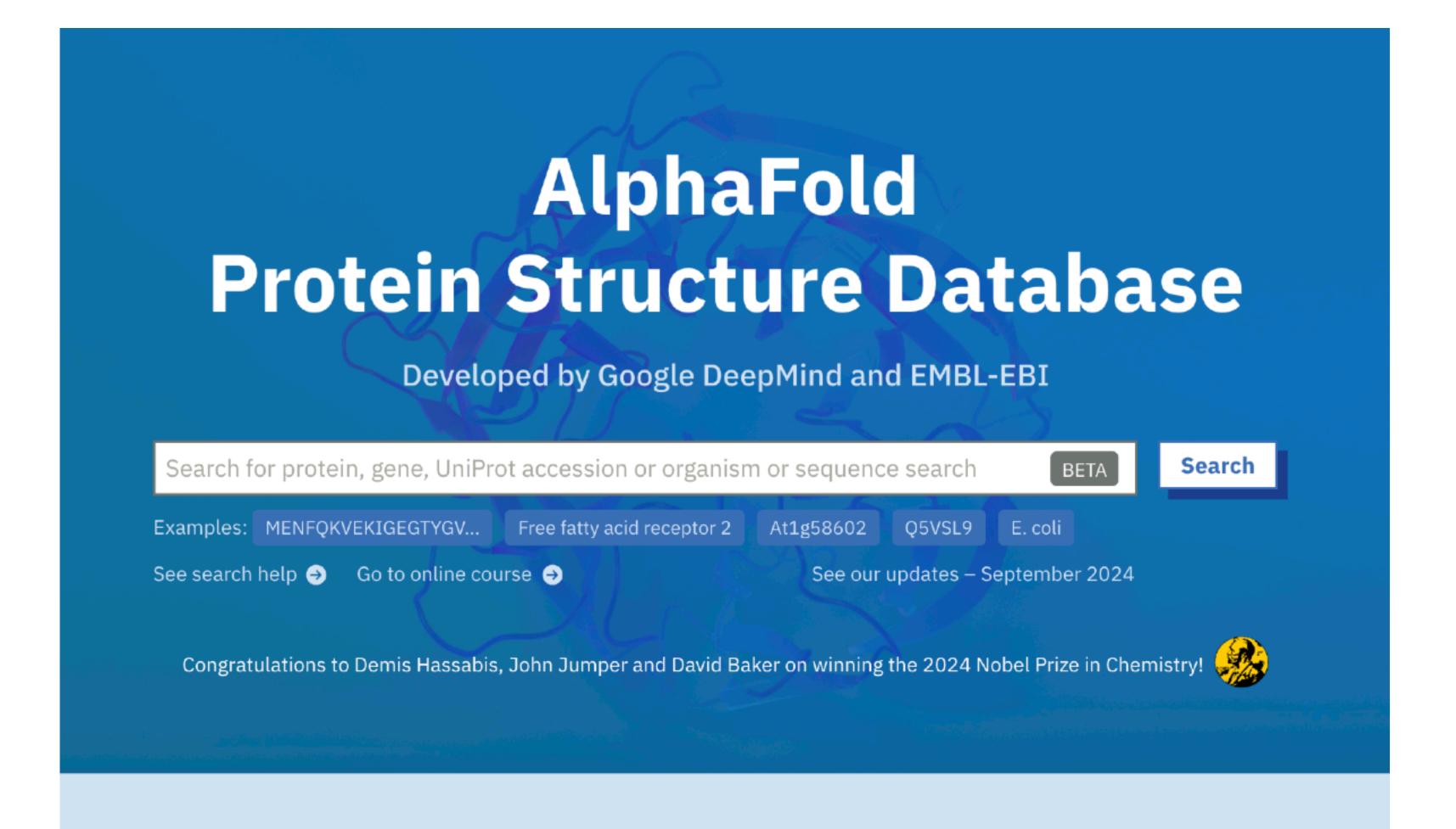
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AlphaFold DB provides open access to over 200 million protein structure predictions to accelerate scientific research.

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What can I help with?

