## Kathryn Tunyasuvunakool

List of Publications by Year in descending order

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#	Article	IF	CITATIONS
1	AlphaFold Protein Structure Database: massively expanding the structural coverage of protein-sequence space with high-accuracy models. Nucleic Acids Research, 2022, 50, D439-D444.	14.5	3,692
2	The prospects and opportunities of protein structure prediction with AI. Nature Reviews Molecular Cell Biology, 2022, 23, 445-446.	37.0	14
3	Discovery of archaeal fusexins homologous to eukaryotic HAP2/GCS1 gamete fusion proteins. Nature Communications, 2022, 13, .	12.8	17
4	Highly accurate protein structure prediction with AlphaFold. Nature, 2021, 596, 583-589.	27.8	17,754
5	Highly accurate protein structure prediction for the human proteome. Nature, 2021, 596, 590-596.	27.8	1,773
6	Applying and improving <scp>AlphaFold</scp> at <scp>CASP14</scp> . Proteins: Structure, Function and Bioinformatics, 2021, 89, 1711-1721.	2.6	231