## Johanna Hakanpää

List of Publications by Year in descending order

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623734 839539 18 1,208 14 18 g-index citations h-index papers 18 18 18 2005 docs citations times ranked citing authors all docs

#	Article	IF	Citations
1	X-ray screening identifies active site and allosteric inhibitors of SARS-CoV-2 main protease. Science, 2021, 372, 642-646.	12.6	240
2	Atomic Resolution Structure of the HFBII Hydrophobin, a Self-assembling Amphiphile. Journal of Biological Chemistry, 2004, 279, 534-539.	3 <b>.</b> 4	205
3	Two crystal structures of Trichoderma reesei hydrophobin HFBI-The structure of a protein amphiphile with and without detergent interaction. Protein Science, 2006, 15, 2129-2140.	7.6	158
4	P13, the EMBL macromolecular crystallography beamline at the low-emittance PETRA III ring forÂhigh-and low-energy phasing with variable beam focusing. Journal of Synchrotron Radiation, 2017, 24, 323-332.	2.4	155
5	Molecular Interactions between a Recombinant IgE Antibody and the β-Lactoglobulin Allergen. Structure, 2007, 15, 1413-1421.	3.3	125
6	Hydrophobin HFBII in detail: ultrahigh-resolution structure at 0.75â€Ã Acta Crystallographica Section D: Biological Crystallography, 2006, 62, 356-367.	<b>2.</b> 5	71
7	Crystal Structures of Hydrophobin HFBII in the Presence of Detergent Implicate the Formation of Fibrils and Monolayer Films. Journal of Biological Chemistry, 2007, 282, 28733-28739.	3.4	50
8	Crystal structures of Trichoderma reesei $\hat{l}^2$ -galactosidase reveal conformational changes in the active site. Journal of Structural Biology, 2011, 174, 156-163.	2.8	47
9	Determination of thioxylo-oligosaccharide binding to family 11 xylanases using electrospray ionization Fourier transform ion cyclotron resonance mass spectrometry and X-ray crystallography. FEBS Journal, 2005, 272, 2317-2333.	4.7	28
10	Novel non- $\hat{l}^2$ -lactam inhibitor of $\hat{l}^2$ -lactamase TEM-171 based on acylated phenoxyaniline. Biochimie, 2017, 132, 45-53.	2.6	26
11	The Contribution of Polystyrene Nanospheres towards the Crystallization of Proteins. PLoS ONE, 2009, 4, e4198.	2.5	24
12	Structural and functional insights into the unique CBS–CP12 fusion protein family in cyanobacteria. Proceedings of the National Academy of Sciences of the United States of America, 2018, 115, 7141-7146.	7.1	20
13	Structure and function of the N-terminal domain of the yeast telomerase reverse transcriptase. Nucleic Acids Research, 2018, 46, 1525-1540.	14.5	19
14	Crystallization and preliminary X-ray characterization of Trichoderma reeseihydrophobin HFBII. Acta Crystallographica Section D: Biological Crystallography, 2004, 60, 163-165.	<b>2.</b> 5	16
15	Characterization and crystallization of a recombinant IgE Fab fragment in complex with the bovine $\hat{I}^2$ -lactoglobulin allergen. Acta Crystallographica Section F: Structural Biology Communications, 2008, 64, 25-28.	0.7	13
16	Amphiphilic nanotubes in the crystal structure of a biosurfactant protein hydrophobin HFBII. Chemical Communications, 2011, 47, 9843.	4.1	6
17	Crystallization and preliminary diffraction analysis of a $\hat{l}^2$ -galactosidase from Trichoderma reesei. Acta Crystallographica Section F: Structural Biology Communications, 2009, 65, 767-769.	0.7	3
18	Polysaccharides as Precipitants in Protein Crystallization for X-Ray Diffraction Studies. Crystal Growth and Design, 2011, 11, 1152-1158.	3.0	2