

# Flora Meilleur

## List of Publications by Year in descending order

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69  
papers

1,764  
citations

257357

24  
h-index

302012

39  
g-index

76  
all docs

76  
docs citations

76  
times ranked

1920  
citing authors

#	ARTICLE	IF	CITATIONS
1	Refinement of macromolecular structures against neutron data with <i>SHELXL2013</i> . <i>Journal of Applied Crystallography</i> , 2014, 47, 462-466.	1.9	152
2	The P132H mutation in the main protease of Omicron SARS-CoV-2 decreases thermal stability without compromising catalysis or small-molecule drug inhibition. <i>Cell Research</i> , 2022, 32, 498-500.	5.7	85
3	Neutron protein crystallography: A complementary tool for locating hydrogens in proteins. <i>Archives of Biochemistry and Biophysics</i> , 2016, 602, 48-60.	1.4	83
4	Oxygen Activation at the Active Site of a Fungal Lytic Polysaccharide Monooxygenase. <i>Angewandte Chemie - International Edition</i> , 2017, 56, 767-770.	7.2	78
5	Quantum model of catalysis based on a mobile proton revealed by subatomic x-ray and neutron diffraction studies of h-aldose reductase. <i>Proceedings of the National Academy of Sciences of the United States of America</i> , 2008, 105, 1844-1848.	3.3	74
6	The IMAGINE instrument: first neutron protein structure and new capabilities for neutron macromolecular crystallography. <i>Acta Crystallographica Section D: Biological Crystallography</i> , 2013, 69, 2157-2160.	2.5	73
7	Expression, purification, assay, and crystal structure of perdeuterated human arginase I. <i>Archives of Biochemistry and Biophysics</i> , 2007, 465, 82-89.	1.4	65
8	High-resolution neutron protein crystallography with radically small crystal volumes: application of perdeuteration to human aldose reductase. <i>Acta Crystallographica Section D: Biological Crystallography</i> , 2005, 61, 1413-1417.	2.5	61
9	Deuterium Labeling for Neutron Structure-Function-Dynamics Analysis. <i>Methods in Molecular Biology</i> , 2009, 544, 281-292.	0.4	55
10	New sources and instrumentation for neutrons in biology. <i>Chemical Physics</i> , 2008, 345, 133-151.	0.9	53
11	Neutron scattering in the biological sciences: progress and prospects. <i>Acta Crystallographica Section D: Structural Biology</i> , 2018, 74, 1129-1168.	1.1	47
12	Structural Stability and Dynamics of Hydrogenated and Perdeuterated Cytochrome P450cam (CYP101). <i>Biochemistry</i> , 2004, 43, 8744-8753.	1.2	46
13	Neutron Crystal Structure of RAS GTPase Puts in Question the Protonation State of the GTP $\hat{\gamma}$ -Phosphate. <i>Journal of Biological Chemistry</i> , 2015, 290, 31025-31036.	1.6	44
14	Open Conformation of Ezrin Bound to Phosphatidylinositol 4,5-Bisphosphate and to F-actin Revealed by Neutron Scattering. <i>Journal of Biological Chemistry</i> , 2012, 287, 37119-37133.	1.6	43
15	A suite-level review of the neutron single-crystal diffraction instruments at Oak Ridge National Laboratory. <i>Review of Scientific Instruments</i> , 2018, 89, 092802.	0.6	43
16	Unambiguous determination of H-atom positions: comparing results from neutron and high-resolution X-ray crystallography. <i>Acta Crystallographica Section D: Biological Crystallography</i> , 2010, 66, 558-567.	2.5	40
17	Production and X-ray crystallographic analysis of fully deuterated cytochrome P450cam. <i>Acta Crystallographica Section D: Biological Crystallography</i> , 2005, 61, 539-544.	2.5	29
18	A quasi-Laue neutron crystallographic study of d-xylose isomerase. <i>European Biophysics Journal</i> , 2006, 35, 601-609.	1.2	28

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19	Comparison of hydrogen determination with X-ray and neutron crystallography in a human aldose reductase inhibitor complex. <i>European Biophysics Journal</i> , 2006, 35, 577-583.	1.2	27
20	Scattering functions of Platonic solids. <i>Journal of Applied Crystallography</i> , 2011, 44, 545-557.	1.9	26
21	Rapid visualization of hydrogen positions in protein neutron crystallographic structures. <i>Acta Crystallographica Section D: Biological Crystallography</i> , 2012, 68, 35-41.	2.5	26
22	Structural investigation of cellobiose dehydrogenase IIA: Insights from small angle scattering into intra- and intermolecular electron transfer mechanisms. <i>Biochimica Et Biophysica Acta - General Subjects</i> , 2018, 1862, 1031-1039.	1.1	26
23	The Neutron Macromolecular Crystallography Instruments at Oak Ridge National Laboratory: Advances, Challenges, and Opportunities. <i>Crystals</i> , 2018, 8, 388.	1.0	26
24	Modulating Enzyme Activity by Altering Protein Dynamics with Solvent. <i>Biochemistry</i> , 2018, 57, 4263-4275.	1.2	26
25	Neutron Laue macromolecular crystallography. <i>European Biophysics Journal</i> , 2006, 35, 611-620.	1.2	25
26	The Structure of Sindbis Virus Produced from Vertebrate and Invertebrate Hosts as Determined by Small-Angle Neutron Scattering. <i>Journal of Virology</i> , 2010, 84, 5270-5276.	1.5	25
27	IMAGINE: neutrons reveal enzyme chemistry. <i>Acta Crystallographica Section D: Structural Biology</i> , 2018, 74, 778-786.	1.1	25
28	The mechanisms of catalysis and ligand binding for the SARS-CoV-2 NSP3 macrodomain from neutron and x-ray diffraction at room temperature. <i>Science Advances</i> , 2022, 8, .	4.7	24
29	Neutron protein crystallography at ultra-low (<math>15\text{K}</math>) temperatures. <i>Journal of Applied Crystallography</i> , 2012, 45, 686-692.	1.9	23
30	Crystal structures of wild-type <i>Trichoderma reesei</i> Cel7A catalytic domain in open and closed states. <i>FEBS Letters</i> , 2016, 590, 4429-4438.	1.3	23
31	Interplay of Intrinsic and Environmental Effects on the Magnetic Properties of Free Radicals Issuing from H-Atom Addition to Cytosine. <i>Journal of the American Chemical Society</i> , 2001, 123, 7113-7117.	6.6	22
32	IMAGINE: The neutron protein crystallography beamline at the high flux isotope reactor. <i>Methods in Enzymology</i> , 2020, 634, 69-85.	0.4	21
33	Next-generation diamond cell and applications to single-crystal neutron diffraction. <i>Review of Scientific Instruments</i> , 2018, 89, 092902.	0.6	20
34	Characterization of image plates for neutron diffraction. <i>Journal of Applied Crystallography</i> , 2009, 42, 749-757.	1.9	19
35	Structural studies of <i>Neurospora crassa</i> LPMO9D and redox partner CDH1A using neutron crystallography and small-angle scattering. <i>Carbohydrate Research</i> , 2017, 448, 200-204.	1.1	19
36	Neutron crystallographic studies of T4 lysozyme at cryogenic temperature. <i>Protein Science</i> , 2017, 26, 2098-2104.	3.1	19

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37	Neutron Scattering Techniques and Applications in Structural Biology. <i>Current Protocols in Protein Science</i> , 2013, 72, Unit17.16.	2.8	18
38	Neutron structure of the cyclic glucose-bound xylose isomerase E186Q mutant. <i>Acta Crystallographica Section D: Biological Crystallography</i> , 2014, 70, 414-420.	2.5	17
39	A preliminary neutron diffraction study of rasburicase, a recombinant urate oxidase enzyme, complexed with 8-azaxanthin. <i>Acta Crystallographica Section F: Structural Biology Communications</i> , 2006, 62, 306-309.	0.7	16
40	Redox-Promoting Protein Motions in Rubredoxin. <i>Journal of Physical Chemistry B</i> , 2011, 115, 8925-8936.	1.2	14
41	Crystallization of a fungal lytic polysaccharide monooxygenase expressed from glycoengineered <i>Pichia pastoris</i> for X-ray and neutron diffraction. <i>Acta Crystallographica Section F, Structural Biology Communications</i> , 2017, 73, 70-78.	0.4	14
42	An extended N-H bond, driven by a conserved second-order interaction, orients the flavin N5 orbital in cholesterol oxidase. <i>Scientific Reports</i> , 2017, 7, 40517.	1.6	14
43	Preliminary neutron crystallographic analysis of selectively CH <sub>3</sub> -protonated deuterated rubredoxin from <i>Pyrococcus furiosus</i> . <i>Acta Crystallographica Section F: Structural Biology Communications</i> , 2008, 64, 537-540.	0.7	13
44	Conformational Changes in Sindbis Virus Induced by Decreased pH Are Revealed by Small-Angle Neutron Scattering. <i>Journal of Virology</i> , 2012, 86, 1982-1987.	1.5	13
45	Neutron structure of the T26H mutant of T4 phage lysozyme provides insight into the catalytic activity of the mutant enzyme and how it differs from that of wild type. <i>Protein Science</i> , 2017, 26, 1953-1963.	3.1	13
46	Optimizing crystal volume for neutron diffraction: D-xylose isomerase. <i>European Biophysics Journal</i> , 2006, 35, 621-632.	1.2	12
47	New insight into the structure of RNA in red clover necrotic mosaic virus and the role of divalent cations revealed by small-angle neutron scattering. <i>Archives of Virology</i> , 2013, 158, 1661-1669.	0.9	10
48	Metalloprotein catalysis: structural and mechanistic insights into oxidoreductases from neutron protein crystallography. <i>Acta Crystallographica Section D: Structural Biology</i> , 2021, 77, 1251-1269.	1.1	10
49	Production and characterization of recombinant perdeuterated cholesterol oxidase. <i>Analytical Biochemistry</i> , 2015, 485, 102-108.	1.1	9
50	Oxygen Activation at the Active Site of a Fungal Lytic Polysaccharide Monooxygenase. <i>Angewandte Chemie</i> , 2017, 129, 785-788.	1.6	9
51	Preliminary neutron diffraction studies of <i>Escherichia coli</i> dihydrofolate reductase bound to the anticancer drug methotrexate. <i>Acta Crystallographica Section D: Biological Crystallography</i> , 2005, 61, 574-579.	2.5	8
52	Dynamic nuclear polarization enhanced neutron crystallography: Amplifying hydrogen in biological crystals. <i>Methods in Enzymology</i> , 2020, 634, 153-175.	0.4	8
53	Crystal structure and thermal expansion of a CsCe <sub>2</sub> Cl <sub>7</sub> scintillator. <i>Journal of Solid State Chemistry</i> , 2015, 227, 142-149.	1.4	6
54	Neutron Crystallography Data Collection and Processing for Modelling Hydrogen Atoms in Protein Structures. <i>Journal of Visualized Experiments</i> , 2020, , .	0.2	6

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55	A beginner's guide to neutron macromolecular crystallography. <i>Biochemist</i> , 2020, 42, 16-20.	0.2	6
56	A prototype system for dynamically polarized neutron protein crystallography. <i>Nuclear Instruments and Methods in Physics Research, Section A: Accelerators, Spectrometers, Detectors and Associated Equipment</i> , 2019, 940, 430-434.	0.7	5
57	Preliminary results of neutron and X-ray diffraction data collection on a lytic polysaccharide monooxygenase under reduced and acidic conditions. <i>Acta Crystallographica Section F, Structural Biology Communications</i> , 2021, 77, 128-133.	0.4	4
58	Crystallization and preliminary X-ray diffraction analysis of red clover necrotic mosaic virus. <i>Acta Crystallographica Section F: Structural Biology Communications</i> , 2010, 66, 1458-1462.	0.7	3
59	Titration of ionizable groups in proteins using multiple neutron data sets from a single crystal: application to the small GTPase Ras. <i>Acta Crystallographica Section F, Structural Biology Communications</i> , 2019, 75, 111-115.	0.4	3
60	Neutron and X-ray analysis of the Fenna-Matthews-Olson photosynthetic antenna complex from <i>Prosthecochloris aestuarii</i> . <i>Acta Crystallographica Section F, Structural Biology Communications</i> , 2019, 75, 171-175.	0.4	3
61	Crystallization and preliminary X-ray diffraction analysis of <i>Hypocrea jecorina</i> Cel7A in two new crystal forms. <i>Acta Crystallographica Section F, Structural Biology Communications</i> , 2014, 70, 773-776.	0.4	2
62	Dynamically polarized samples for neutron protein crystallography at the Spallation Neutron Source. <i>Journal of Physics: Conference Series</i> , 2016, 746, 012008.	0.3	2
63	Third School on the Applications of Neutron Scattering Techniques in Structural Biology, Oak Ridge, TN. <i>Neutron News</i> , 2013, 24, 4-4.	0.1	1
64	ORNL hosts first virtual HANDS workshop. <i>Neutron News</i> , 2020, 31, 11-12.	0.1	1
65	ORNL hosts the participants of the 4th Neutrons in Structural Biology Workshop and the IMAGINE single crystal neutron diffractometer first external users. <i>Neutron News</i> , 2014, 25, 12-12.	0.1	0
66	Innenteilbild: Oxygen Activation at the Active Site of a Fungal Lytic Polysaccharide Monooxygenase ( <i>Angew. Chem.</i> 3/2017). <i>Angewandte Chemie</i> , 2017, 129, 674-674.	1.6	0
67	Characterization of biomass-degrading enzymes using neutron diffraction and scattering. <i>Neutron News</i> , 2021, 32, 13-14.	0.1	0
68	Neutron Crystal Structure of Ras GTPase sets New Paradigm for GTP Hydrolysis. <i>FASEB Journal</i> , 2015, 29, 893.7.	0.2	0
69	Teaching and Education highlighted. <i>Journal of Applied Crystallography</i> , 2022, 55, 215-217.	1.9	0