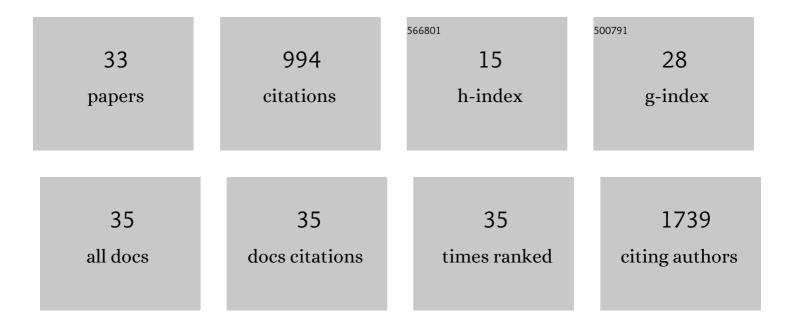
Po-Lin Chiu

List of Publications by Year in descending order

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#	Article	IF	CITATIONS
1	Probing Structural Perturbation of Biomolecules by Extracting Cryo-EM Data Heterogeneity. Biomolecules, 2022, 12, 628.	1.8	4
2	Protein secondary structure signatures from energy loss spectra recorded in the electron microscope. Journal of Microscopy, 2021, 282, 215-223.	0.8	6
3	Structural and Functional Analysis of Disease-Linked p97 ATPase Mutant Complexes. International Journal of Molecular Sciences, 2021, 22, 8079.	1.8	11
4	Cryo-EM reveals conformational flexibility in apo DNA polymerase ζ. Journal of Biological Chemistry, 2021, 297, 100912.	1.6	5
5	Conserved L464 in p97 D1–D2 linker is critical for p97 cofactor regulated ATPase activity. Biochemical Journal, 2021, 478, 3185-3204.	1.7	1
6	Sample Preparation and Data Collection for Electron Crystallographic Studies on Membrane Structures and Lipid–Protein Interaction. Methods in Molecular Biology, 2021, 2215, 247-265.	0.4	0
7	Sample Preparation using a Lipid Monolayer Method for Electron Crystallographic Studies. Journal of Visualized Experiments, 2021, , .	0.2	0
8	ChAdOx1 interacts with CAR and PF4 with implications for thrombosis with thrombocytopenia syndrome. Science Advances, 2021, 7, eabl8213.	4.7	112
9	Structural basis of redox modulation on chloroplast ATP synthase. Communications Biology, 2020, 3, 482.	2.0	25
10	A library of aminoglycoside-derived lipopolymer nanoparticles for delivery of small molecules and nucleic acids. Journal of Materials Chemistry B, 2020, 8, 8558-8572.	2.9	3
11	Dynamic self-assembly of detonation nanodiamond in water. Nanoscale, 2020, 12, 5363-5367.	2.8	34
12	Isolation of Functional Temperature Activated Transmembrane Domain of Human TRPM8. Biophysical Journal, 2020, 118, 413a.	0.2	0
13	Structural Studies using Cryo-EM to Unravel Mechanistic Details of p47 Binding To p97. Biophysical Journal, 2020, 118, 501a.	0.2	0
14	The structure of the stress-induced photosystem l–IsiA antenna supercomplex. Nature Structural and Molecular Biology, 2019, 26, 443-449.	3.6	50
15	The <i>C. elegans</i> 3′ UTRome v2 resource for studying mRNA cleavage and polyadenylation, 3′-UTR biology, and miRNA targeting. Genome Research, 2019, 29, 2104-2116.	2.4	14
16	Membrane directed expression in Escherichia coli of BBA57 and other virulence factors from the Lyme disease agent Borrelia burgdorferi. Scientific Reports, 2019, 9, 17606.	1.6	4
17	The structural basis for the functional comparability of factor VIII and the longâ€acting variant recombinant factor VIII Fc fusion protein. Journal of Thrombosis and Haemostasis, 2017, 15, 1167-1179.	1.9	14
18	Electron crystallography reveals that substrate release from the PTS IIC glucose transporter is coupled to a subtle conformational change. Journal of Structural Biology, 2017, 199, 39-45.	1.3	5

Ро-Lім Сній

#	Article	IF	CITATIONS
19	Macromolecular Assemblies of the Mammalian Circadian Clock. Molecular Cell, 2017, 67, 770-782.e6.	4.5	198
20	Magnetically Directed Two-Dimensional Crystallization of OmpF Membrane Proteins in Block Copolymers. Journal of the American Chemical Society, 2016, 138, 28-31.	6.6	27
21	Evaluation of super-resolution performance of the K2 electron-counting camera using 2D crystals of aquaporin-0. Journal of Structural Biology, 2015, 192, 163-173.	1.3	26
22	Mapping the interaction between factor VIII and von Willebrand factor by electron microscopy and mass spectrometry. Blood, 2015, 126, 935-938.	0.6	53
23	2D and 3D crystallization of the wild-type IIC domain of the glucose PTS transporter from Escherichia coli. Journal of Structural Biology, 2015, 191, 376-380.	1.3	5
24	Effect of Lipid Head Groups on Double-Layered Two-Dimensional Crystals Formed by Aquaporin-0. PLoS ONE, 2015, 10, e0117371.	1.1	9
25	Nose-to-brain transport of aerosolised quantum dots following acute exposure. Nanotoxicology, 2014, 8, 885-893.	1.6	75
26	Ligand-induced structural changes in the cyclic nucleotide-modulated potassium channel MloK1. Nature Communications, 2014, 5, 3106.	5.8	59
27	Assessing Structure-Activity Relationships in FVIII By Integration of Structural, Biophysical, and Biochemical Data Obtained with Anti-FVIII Antibodies. Blood, 2014, 124, 4223-4223.	0.6	1
28	Evidence For Flexible Tethering Of Fc To FVIII In Recombinant FVIII-Fc Fusion Protein rFVIIIFc. Blood, 2013, 122, 1102-1102.	0.6	0
29	Visualizing macromolecular complexes with in situ liquid scanning transmission electron microscopy. Micron, 2012, 43, 1085-1090.	1.1	89
30	The use of trehalose in the preparation of specimens for molecular electron microscopy. Micron, 2011, 42, 762-772.	1.1	20
31	An optical and microPET assessment of thermally-sensitive liposome biodistribution in the Met-1 tumor model: Importance of formulation. Journal of Controlled Release, 2010, 143, 13-22.	4.8	56
32	The Structure of the Prokaryotic Cyclic Nucleotide-Modulated Potassium Channel MloK1 at 16 Ã Resolution. Structure, 2007, 15, 1053-1064.	1.6	51
33	Milestones in electron crystallography. Journal of Computer-Aided Molecular Design, 2006, 20, 519-527.	1.3	32