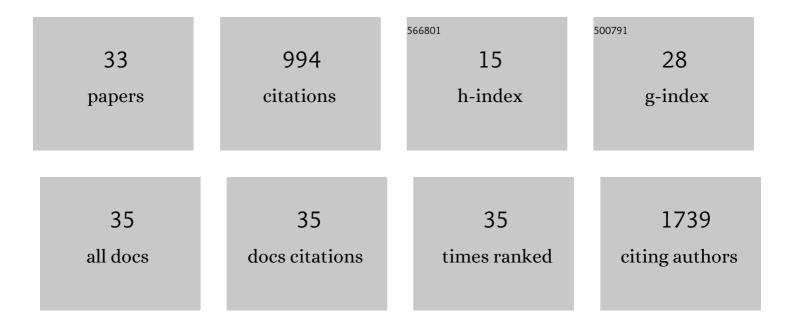
Po-Lin Chiu

List of Publications by Year in descending order

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PO-LIN CHIL

#	Article	IF	CITATIONS
1	Macromolecular Assemblies of the Mammalian Circadian Clock. Molecular Cell, 2017, 67, 770-782.e6.	4.5	198
2	ChAdOx1 interacts with CAR and PF4 with implications for thrombosis with thrombocytopenia syndrome. Science Advances, 2021, 7, eabl8213.	4.7	112
3	Visualizing macromolecular complexes with in situ liquid scanning transmission electron microscopy. Micron, 2012, 43, 1085-1090.	1.1	89
4	Nose-to-brain transport of aerosolised quantum dots following acute exposure. Nanotoxicology, 2014, 8, 885-893.	1.6	75
5	Ligand-induced structural changes in the cyclic nucleotide-modulated potassium channel MloK1. Nature Communications, 2014, 5, 3106.	5.8	59
6	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
7	Mapping the interaction between factor VIII and von Willebrand factor by electron microscopy and mass spectrometry. Blood, 2015, 126, 935-938.	0.6	53
8	The Structure of the Prokaryotic Cyclic Nucleotide-Modulated Potassium Channel MloK1 at 16 Ã Resolution. Structure, 2007, 15, 1053-1064.	1.6	51
9	The structure of the stress-induced photosystem l–IsiA antenna supercomplex. Nature Structural and Molecular Biology, 2019, 26, 443-449.	3.6	50
10	Dynamic self-assembly of detonation nanodiamond in water. Nanoscale, 2020, 12, 5363-5367.	2.8	34
11	Milestones in electron crystallography. Journal of Computer-Aided Molecular Design, 2006, 20, 519-527.	1.3	32
12	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
13	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
14	Structural basis of redox modulation on chloroplast ATP synthase. Communications Biology, 2020, 3, 482.	2.0	25
15	The use of trehalose in the preparation of specimens for molecular electron microscopy. Micron, 2011, 42, 762-772.	1.1	20
16	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
17	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
18	Structural and Functional Analysis of Disease-Linked p97 ATPase Mutant Complexes. International Journal of Molecular Sciences, 2021, 22, 8079.	1.8	11

Ро-Lім Сній

#	Article	lF	CITATIONS
19	Effect of Lipid Head Groups on Double-Layered Two-Dimensional Crystals Formed by Aquaporin-0. PLoS ONE, 2015, 10, e0117371.	1.1	9
20	Protein secondary structure signatures from energy loss spectra recorded in the electron microscope. Journal of Microscopy, 2021, 282, 215-223.	0.8	6
21	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
22	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
23	Cryo-EM reveals conformational flexibility in apo DNA polymerase ζ. Journal of Biological Chemistry, 2021, 297, 100912.	1.6	5
24	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
25	Probing Structural Perturbation of Biomolecules by Extracting Cryo-EM Data Heterogeneity. Biomolecules, 2022, 12, 628.	1.8	4
26	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
27	Conserved L464 in p97 D1–D2 linker is critical for p97 cofactor regulated ATPase activity. Biochemical Journal, 2021, 478, 3185-3204.	1.7	1
28	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
29	Isolation of Functional Temperature Activated Transmembrane Domain of Human TRPM8. Biophysical Journal, 2020, 118, 413a.	0.2	0
30	Structural Studies using Cryo-EM to Unravel Mechanistic Details of p47 Binding To p97. Biophysical Journal, 2020, 118, 501a.	0.2	0
31	Evidence For Flexible Tethering Of Fc To FVIII In Recombinant FVIII-Fc Fusion Protein rFVIIIFc. Blood, 2013, 122, 1102-1102.	0.6	0
32	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
33	Sample Preparation using a Lipid Monolayer Method for Electron Crystallographic Studies. Journal of Visualized Experiments, 2021, , .	0.2	0