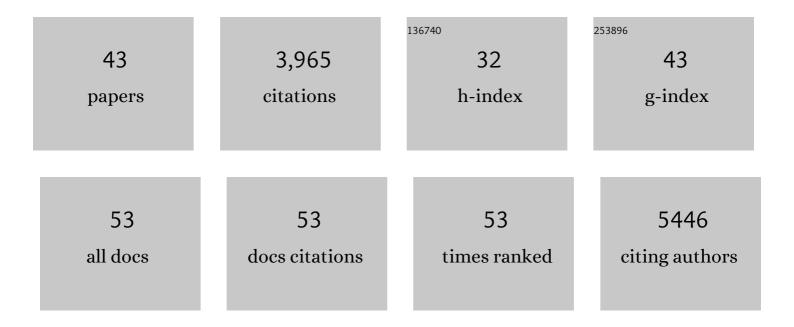
## Elisabeth Paula Carpenter

List of Publications by Year in descending order

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#	Article	IF	CITATIONS
1	Overcoming the challenges of membrane protein crystallography. Current Opinion in Structural Biology, 2008, 18, 581-586.	2.6	419
2	Structure and Molecular Mechanism of a Nucleobase–Cation–Symport-1 Family Transporter. Science, 2008, 322, 709-713.	6.0	347
3	Crystal structure of a prokaryotic homologue of the mammalian oligopeptide-proton symporters, PepT1 and PepT2. EMBO Journal, 2011, 30, 417-426.	3.5	269
4	MemProtMD: Automated Insertion of Membrane Protein Structures into Explicit Lipid Membranes. Structure, 2015, 23, 1350-1361.	1.6	257
5	K2P channel gating mechanisms revealed by structures of TREK-2 and a complex with Prozac. Science, 2015, 347, 1256-1259.	6.0	255
6	Structures of ABCB10, a human ATP-binding cassette transporter in apo- and nucleotide-bound states. Proceedings of the National Academy of Sciences of the United States of America, 2013, 110, 9710-9715.	3.3	219
7	Structural and functional diversity calls for a new classification of ABC transporters. FEBS Letters, 2020, 594, 3767-3775.	1.3	169
8	Structure of the polycystic kidney disease TRP channel Polycystin-2 (PC2). Nature Structural and Molecular Biology, 2017, 24, 114-122.	3.6	155
9	Structure of dehydroquinate synthase reveals an active site capable of multistep catalysis. Nature, 1998, 394, 299-302.	13.7	134
10	Protein farnesyl and N-myristoyl transferases: piggy-back medicinal chemistry targets for the development of antitrypanosomatid and antimalarial therapeutics. Molecular and Biochemical Parasitology, 2003, 126, 155-163.	0.5	126
11	Molecular insights into lipid-assisted Ca2+ regulation of the TRP channel Polycystin-2. Nature Structural and Molecular Biology, 2017, 24, 123-130.	3.6	105
12	The structural basis of lipid scrambling and inactivation in the endoplasmic reticulum scramblase TMEM16K. Nature Communications, 2019, 10, 3956.	5.8	101
13	Atomic resolution insight into host cell recognition by Toxoplasma gondii. EMBO Journal, 2007, 26, 2808-2820.	3.5	98
14	A pharmacological master key mechanism that unlocks the selectivity filter gate in K <sup>+</sup> channels. Science, 2019, 363, 875-880.	6.0	91
15	The Structural Basis of ZMPSTE24-Dependent Laminopathies. Science, 2013, 339, 1604-1607.	6.0	89
16	Structures of DPAGT1 Explain Glycosylation Disease Mechanisms and Advance TB Antibiotic Design. Cell, 2018, 175, 1045-1058.e16.	13.5	67
17	Polymodal activation of the TREK-2 K2P channel produces structurally distinct open states. Journal of General Physiology, 2016, 147, 497-505.	0.9	65
18	Bilayer-Mediated Structural Transitions Control Mechanosensitivity of the TREK-2 K2P Channel. Structure, 2017, 25, 708-718.e2.	1.6	64

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19	Mass spectrometry captures off-target drug binding and provides mechanistic insights into the human metalloprotease ZMPSTE24. Nature Chemistry, 2016, 8, 1152-1158.	6.6	61
20	Structure of a central stalk subunit F of prokaryotic V-type ATPase/synthase from Thermus thermophilus. EMBO Journal, 2005, 24, 3974-3983.	3.5	53
21	Mammalian Glucose Transporter Activity Is Dependent upon Anionic and Conical Phospholipids. Journal of Biological Chemistry, 2016, 291, 17271-17282.	1.6	53
22	A lower X-gate in TASK channels traps inhibitors within the vestibule. Nature, 2020, 582, 443-447.	13.7	53
23	A Comparative Study of Uracil-DNA Glycosylases from Human and Herpes Simplex Virus Type 1. Journal of Biological Chemistry, 2006, 281, 4983-4992.	1.6	52
24	The structural basis of fatty acid elongation by the ELOVL elongases. Nature Structural and Molecular Biology, 2021, 28, 512-520.	3.6	52
25	Structures and functions of mitochondrial ABC transporters. Biochemical Society Transactions, 2015, 43, 943-951.	1.6	50
26	Crystal Structure of the Acid-Induced Arginine Decarboxylase from <i>Escherichia coli</i> : Reversible Decamer Assembly Controls Enzyme Activity. Biochemistry, 2009, 48, 3915-3927.	1.2	48
27	AP endonuclease paralogues with distinct activities in DNA repair and bacterial pathogenesis. EMBO Journal, 2007, 26, 1363-1372.	3.5	47
28	Four Distinct Structural Domains in Clostridium difficile Toxin B Visualized Using SAXS. Journal of Molecular Biology, 2010, 396, 1260-1270.	2.0	46
29	Asymmetric mechanosensitivity in a eukaryotic ion channel. Proceedings of the National Academy of Sciences of the United States of America, 2017, 114, E8343-E8351.	3.3	45
30	Insights into How Nucleotide-Binding Domains Power ABC Transport. Structure, 2009, 17, 1213-1222.	1.6	40
31	An overview of heavy-atom derivatization of protein crystals. Acta Crystallographica Section D: Structural Biology, 2016, 72, 303-318.	1.1	40
32	Insights into outer membrane protein crystallization. Molecular Membrane Biology, 2008, 25, 631-638.	2.0	37
33	Lipid Interactions of a Ciliary Membrane TRP Channel: Simulation and Structural Studies of Polycystin-2. Structure, 2020, 28, 169-184.e5.	1.6	37
34	Purification and interaction analyses of two human lysosomal vitamin B <sub>12</sub> transporters: LMBD1 and ABCD4. Molecular Membrane Biology, 2014, 31, 250-261.	2.0	31
35	ABCB10 exports mitochondrial biliverdin, driving metabolic maladaptation in obesity. Science Translational Medicine, 2021, 13, .	5.8	27
36	High-performance liquid chromatography separation and intact mass analysis of detergent-solubilized integral membrane proteins. Analytical Biochemistry, 2011, 410, 272-280.	1.1	24

#	Article	IF	CITATIONS
37	<i>ZMPSTE24</i> missense mutations that cause progeroid diseases decrease prelamin A cleavage activity and/or protein stability. DMM Disease Models and Mechanisms, 2018, 11, .	1.2	24
38	Thioredoxin A Active-Site Mutants Form Mixed Disulfide Dimers That Resemble Enzyme–Substrate Reaction Intermediates. Journal of Molecular Biology, 2008, 379, 520-534.	2.0	20
39	Structural basis for the recognition and cleavage of abasic DNA in <i>Neisseria meningitidis</i> . Proceedings of the National Academy of Sciences of the United States of America, 2012, 109, 16852-16857.	3.3	19
40	Norfluoxetine inhibits TREK-2 K2P channels by multiple mechanisms including state-independent effects on the selectivity filter gate. Journal of General Physiology, 2021, 153, .	0.9	17
41	The SGC beyond structural genomics: redefining the role of 3D structures by coupling genomic stratification with fragment-based discovery. Essays in Biochemistry, 2017, 61, 495-503.	2.1	12
42	Site specificity determinants for prelamin A cleavage by the zinc metalloprotease ZMPSTE24. Journal of Biological Chemistry, 2021, 296, 100165.	1.6	12
43	Targeted next generation sequencing identifies a genetic spectrum of DNA variants in patients with hemiplegic migraine. Cephalalgia Reports, 2019, 2, 251581631988163.	0.2	8