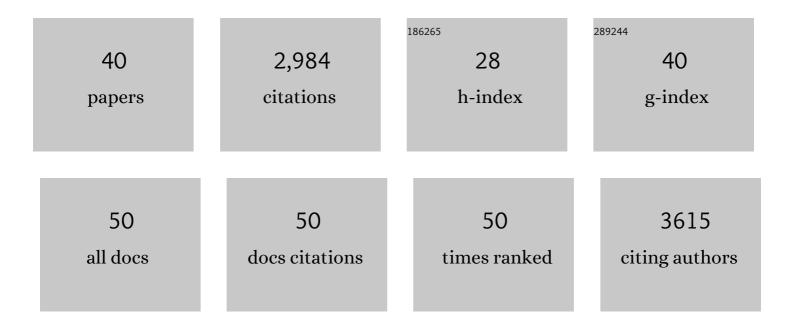
Seok-Yong Lee

List of Publications by Year in descending order

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#	Article	IF	CITATIONS
1	Structure and Mechanism of the Lipid Flippase MurJ. Annual Review of Biochemistry, 2022, 91, 705-729.	11.1	8
2	Recent advances on the inhibition of human solute carriers: Therapeutic implications and mechanistic insights. Current Opinion in Structural Biology, 2022, 74, 102378.	5.7	5
3	Vanilloid-dependent TRPV1 opening trajectory from cryoEM ensemble analysis. Nature Communications, 2022, 13, .	12.8	24
4	Structural basis for inhibition and regulation of a chitin synthase from Candida albicans. Nature Structural and Molecular Biology, 2022, 29, 653-664.	8.2	34
5	Toward a Molecular Basis of Cellular Nucleoside Transport in Humans. Chemical Reviews, 2021, 121, 5336-5358.	47.7	19
6	Synthesis and evaluation of cyclopentane-based muraymycin analogs targeting MraY. European Journal of Medicinal Chemistry, 2021, 215, 113272.	5.5	3
7	Heat-dependent opening of TRPV1 in the presence of capsaicin. Nature Structural and Molecular Biology, 2021, 28, 554-563.	8.2	92
8	Sample preparation of the human TRPA1 ion channel for cryo-EM studies. Methods in Enzymology, 2021, 653, 75-87.	1.0	0
9	Molecular Sensors of Temperature, Pressure, and Pain with Special Focus on TRPV1, TRPM8, and PIEZO2 Ion Channels. Neuroscience Bulletin, 2021, 37, 1745-1749.	2.9	12
10	Structural Insights into Electrophile Irritant Sensing by the Human TRPA1 Channel. Neuron, 2020, 105, 882-894.e5.	8.1	81
11	Current View of Ligand and Lipid Recognition by the Menthol Receptor TRPM8. Trends in Biochemical Sciences, 2020, 45, 806-819.	7.5	14
12	Structures of Bacterial MraY and Human GPT Provide Insights into Rational Antibiotic Design. Journal of Molecular Biology, 2020, 432, 4946-4963.	4.2	14
13	The role of π-helices in TRP channel gating. Current Opinion in Structural Biology, 2019, 58, 314-323.	5.7	47
14	Chemical logic of MraY inhibition by antibacterial nucleoside natural products. Nature Communications, 2019, 10, 2917.	12.8	49
15	Visualizing structural transitions of ligand-dependent gating of the TRPM2 channel. Nature Communications, 2019, 10, 3740.	12.8	34
16	Structures of human ENT1 in complex with adenosine reuptake inhibitors. Nature Structural and Molecular Biology, 2019, 26, 599-606.	8.2	65
17	Visualizing conformation transitions of the Lipid II flippase MurJ. Nature Communications, 2019, 10, 1736.	12.8	51
18	Structural basis of cooling agent and lipid sensing by the cold-activated TRPM8 channel. Science, 2019, 363	12.6	171

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19	Ca2+/CaM interaction with voltage-gated Na+channels. Proceedings of the National Academy of Sciences of the United States of America, 2019, 116, 26150-26151.	7.1	1
20	Symmetry transitions during gating of the TRPV2 ion channel in lipid membranes. ELife, 2019, 8, .	6.0	39
21	Regulatory switch at the cytoplasmic interface controls TRPV channel gating. ELife, 2019, 8, .	6.0	53
22	GlcNAc-1-P-transferase–tunicamycin complex structure reveals basis for inhibition of N-glycosylation. Nature Structural and Molecular Biology, 2018, 25, 217-224.	8.2	104
23	Differential Inhibition of Nav1.7 and Neuropathic Pain by Hybridoma-Produced and Recombinant Monoclonal Antibodies that Target Nav1.7. Neuroscience Bulletin, 2018, 34, 22-41.	2.9	22
24	Conformational plasticity in the selectivity filter of the TRPV2 ion channel. Nature Structural and Molecular Biology, 2018, 25, 405-415.	8.2	79
25	Expression and Role of Voltage-Gated Sodium Channels in Human Dorsal Root Ganglion Neurons with Special Focus on Nav1.7, Species Differences, and Regulation by Paclitaxel. Neuroscience Bulletin, 2018, 34, 4-12.	2.9	97
26	Structure of the cold- and menthol-sensing ion channel TRPM8. Science, 2018, 359, 237-241.	12.6	234
27	Conformational ensemble of the human TRPV3 ion channel. Nature Communications, 2018, 9, 4773.	12.8	100
28	Cryo-EM structure of a mitochondrial calcium uniporter. Science, 2018, 361, 506-511.	12.6	116
29	Visualizing multistep elevator-like transitions of a nucleoside transporter. Nature, 2017, 545, 66-70.	27.8	49
30	Crystal structure of the MOP flippase MurJ in an inward-facing conformation. Nature Structural and Molecular Biology, 2017, 24, 171-176.	8.2	71
31	Cryo-electron microscopy structure of the lysosomal calcium-permeable channel TRPML3. Nature, 2017, 550, 411-414.	27.8	104
32	Current view on regulation of voltageâ€gated sodium channels by calcium and auxiliary proteins. Protein Science, 2016, 25, 1573-1584.	7.6	40
33	Structural insights into inhibition of lipid I production in bacterial cell wall synthesis. Nature, 2016, 533, 557-560.	27.8	96
34	Cryo-electron microscopy structure of the TRPV2 ion channel. Nature Structural and Molecular Biology, 2016, 23, 180-186.	8.2	251
35	Liposome Reconstitution and Transport Assay for Recombinant Transporters. Methods in Enzymology, 2015, 556, 373-383.	1.0	16
36	Structural basis of nucleoside and nucleoside drug selectivity by concentrative nucleoside transporters. ELife, 2014, 3, e03604.	6.0	50

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37	Structural analyses of Ca2+/CaM interaction with NaV channel C-termini reveal mechanisms of calcium-dependent regulation. Nature Communications, 2014, 5, 4896.	12.8	86
38	Crystal Structure of MraY, an Essential Membrane Enzyme for Bacterial Cell Wall Synthesis. Science, 2013, 341, 1012-1016.	12.6	194
39	Crystal structure of a concentrative nucleoside transporter from Vibrio cholerae at 2.4 à Nature, 2012, 483, 489-493.	27.8	117
40	Crystal Structure of the Ternary Complex of a NaV C-Terminal Domain, a Fibroblast Growth Factor Homologous Factor, and Calmodulin. Structure, 2012, 20, 1167-1176.	3.3	138