

Hiro Furukawa

List of Publications by Year in descending order

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Version: 2024-02-01

35
papers

3,115
citations

218677

26
h-index

345221

36
g-index

38
all docs

38
docs citations

38
times ranked

2742
citing authors

#	ARTICLE	IF	CITATIONS
1	Crystal structure of a heterotetrameric NMDA receptor ion channel. <i>Science</i> , 2014, 344, 992-997.	12.6	500
2	Structure, function, and allosteric modulation of NMDA receptors. <i>Journal of General Physiology</i> , 2018, 150, 1081-1105.	1.9	363
3	Subunit arrangement and phenylethanolamine binding in GluN1/GluN2B NMDA receptors. <i>Nature</i> , 2011, 475, 249-253.	27.8	302
4	Structure, Function, and Pharmacology of Glutamate Receptor Ion Channels. <i>Pharmacological Reviews</i> , 2021, 73, 1469-1658.	16.0	237
5	Activation of NMDA receptors and the mechanism of inhibition by ifenprodil. <i>Nature</i> , 2016, 534, 63-68.	27.8	173
6	Structure of the zinc-bound amino-terminal domain of the NMDA receptor NR2B subunit. <i>EMBO Journal</i> , 2009, 28, 3910-3920.	7.8	171
7	The Cryo-EM structure of pannexin 1 reveals unique motifs for ion selection and inhibition. <i>ELife</i> , 2020, 9, .	6.0	103
8	Control of Assembly and Function of Glutamate Receptors by the Amino-Terminal Domain. <i>Molecular Pharmacology</i> , 2010, 78, 535-549.	2.3	95
9	Ligand-specific deactivation time course of GluN1/GluN2D NMDA receptors. <i>Nature Communications</i> , 2011, 2, 294.	12.8	78
10	Structural Determinants of Agonist Efficacy at the Glutamate Binding Site of N-Methyl-D-Aspartate Receptors. <i>Molecular Pharmacology</i> , 2013, 84, 114-127.	2.3	76
11	Structural Insights into Competitive Antagonism in NMDA Receptors. <i>Neuron</i> , 2014, 81, 366-378.	8.1	75
12	A structural biology perspective on NMDA receptor pharmacology and function. <i>Current Opinion in Structural Biology</i> , 2015, 33, 68-75.	5.7	70
13	Molecular Basis for Subtype Specificity and High-Affinity Zinc Inhibition in the GluN1-GluN2A NMDA Receptor Amino-Terminal Domain. <i>Neuron</i> , 2016, 92, 1324-1336.	8.1	70
14	Structural Determinants and Mechanism of Action of a GluN2C-selective NMDA Receptor Positive Allosteric Modulator. <i>Molecular Pharmacology</i> , 2014, 86, 548-560.	2.3	69
15	Structural Basis of Functional Transitions in Mammalian NMDA Receptors. <i>Cell</i> , 2020, 182, 357-371.e13.	28.9	66
16	Emerging structural insights into the function of ionotropic glutamate receptors. <i>Trends in Biochemical Sciences</i> , 2015, 40, 328-337.	7.5	64
17	Structural Mechanism of Functional Modulation by Gene Splicing in NMDA Receptors. <i>Neuron</i> , 2018, 98, 521-529.e3.	8.1	57
18	Structural Mechanism for Modulation of Synaptic Neuroligin-Neurexin Signaling by MDGA Proteins. <i>Neuron</i> , 2017, 95, 896-913.e10.	8.1	55

#	ARTICLE	IF	CITATIONS
19	Role of heterotrimeric G $\beta\gamma$ proteins in maize development and enhancement of agronomic traits. <i>PLoS Genetics</i> , 2018, 14, e1007374.	3.5	55
20	Structure and assembly of calcium homeostasis modulator proteins. <i>Nature Structural and Molecular Biology</i> , 2020, 27, 150-159.	8.2	55
21	Mapping the Binding of GluN2B-Selective N-Methyl-D-aspartate Receptor Negative Allosteric Modulators. <i>Molecular Pharmacology</i> , 2012, 82, 344-359.	2.3	44
22	On the molecular nature of large-pore channels. <i>Journal of Molecular Biology</i> , 2021, 433, 166994.	4.2	44
23	Divergent roles of a peripheral transmembrane segment in AMPA and NMDA receptors. <i>Journal of General Physiology</i> , 2017, 149, 661-680.	1.9	41
24	Dissecting diverse functions of NMDA receptors by structural biology. <i>Current Opinion in Structural Biology</i> , 2019, 54, 34-42.	5.7	37
25	Structure and function of glutamate receptor amino terminal domains. <i>Journal of Physiology</i> , 2012, 590, 63-72.	2.9	35
26	Structural elements of a pH-sensitive inhibitor binding site in NMDA receptors. <i>Nature Communications</i> , 2019, 10, 321.	12.8	32
27	A Eukaryotic Specific Transmembrane Segment is Required for Tetramerization in AMPA Receptors. <i>Journal of Neuroscience</i> , 2013, 33, 9840-9845.	3.6	31
28	Novel Mode of Antagonist Binding in NMDA Receptors Revealed by the Crystal Structure of the GluN1-GluN2A Ligand-Binding Domain Complexed to NVP-AAM077. <i>Molecular Pharmacology</i> , 2017, 92, 22-29.	2.3	27
29	Hodgkin-Huxley-Katz Prize Lecture: Genetic and pharmacological control of glutamate receptor channel through a highly conserved gating motif. <i>Journal of Physiology</i> , 2020, 598, 3071-3083.	2.9	23
30	Structural insights into binding of therapeutic channel blockers in NMDA receptors. <i>Nature Structural and Molecular Biology</i> , 2022, 29, 507-518.	8.2	21
31	Structural basis of subtype-selective competitive antagonism for GluN2C/2D-containing NMDA receptors. <i>Nature Communications</i> , 2020, 11, 423.	12.8	19
32	Development and characterization of functional antibodies targeting NMDA receptors. <i>Nature Communications</i> , 2022, 13, 923.	12.8	11
33	Effective production of oligomeric membrane proteins by EarlyBac-insect cell system. <i>Methods in Enzymology</i> , 2021, 653, 3-19.	1.0	7
34	Deeper Insights into the Allosteric Modulation of Ionotropic Glutamate Receptors. <i>Neuron</i> , 2016, 91, 1187-1189.	8.1	2
35	Production of Heteromeric Transmembrane Receptors with Defined Subunit Stoichiometry. <i>Structure</i> , 2016, 24, 653-655.	3.3	1