Richard M Walsh Jr

List of Publications by Year in descending order

Source: https://exaly.com/author-pdf/6673428/publications.pdf

Version: 2024-02-01

18 papers 2,829 citations

840585 11 h-index 996849 15 g-index

21 all docs

21 docs citations

times ranked

21

4892 citing authors

#	Article	IF	CITATIONS
1	Distinct conformational states of SARS-CoV-2 spike protein. Science, 2020, 369, 1586-1592.	6.0	995
2	Structure of a human synaptic GABAA receptor. Nature, 2018, 559, 67-72.	13.7	389
3	Structural impact on SARS-CoV-2 spike protein by D614G substitution. Science, 2021, 372, 525-530.	6.0	344
4	Membrane fusion and immune evasion by the spike protein of SARS-CoV-2 Delta variant. Science, 2021, 374, 1353-1360.	6.0	246
5	Structural basis for enhanced infectivity and immune evasion of SARS-CoV-2 variants. Science, 2021, 373, 642-648.	6.0	211
6	Shared structural mechanisms of general anaesthetics and benzodiazepines. Nature, 2020, 585, 303-308.	13.7	195
7	Structural principles of distinct assemblies of the human $\hat{l}\pm4\hat{l}^22$ nicotinic receptor. Nature, 2018, 557, 261-265.	13.7	177
8	Agonist Selectivity and Ion Permeation in the $\hat{l}\pm3\hat{l}^24$ Ganglionic Nicotinic Receptor. Neuron, 2019, 104, 501-511.e6.	3.8	131
9	Structures of chaperone-associated assembly intermediates reveal coordinated mechanisms of proteasome biogenesis. Nature Structural and Molecular Biology, 2021, 28, 418-425.	3.6	29
10	Human UDP-α-d-xylose Synthase andEscherichia coliArnA Conserve a Conformational Shunt That Controls Whether Xylose or 4-Keto-Xylose Is Produced. Biochemistry, 2012, 51, 8844-8855.	1.2	21
11	Evidence of Kinetic Cooperativity in Dimeric Ketopantoate Reductase from <i>Staphylococcus aureus</i> . Biochemistry, 2015, 54, 3360-3369.	1.2	14
12	Hysteresis in Human UDP-Glucose Dehydrogenase Is Due to a Restrained Hexameric Structure That Favors Feedback Inhibition. Biochemistry, 2014, 53, 8043-8051.	1.2	9
13	Chaperone-mediated assembly of the proteasome core particle – recent developments and structural insights. Journal of Cell Science, 2022, 135, .	1.2	7
14	Human UDP-α- <scp>d</scp> -xylose Synthase Forms a Catalytically Important Tetramer That Has Not Been Observed in Crystal Structures. Biochemistry, 2013, 52, 3888-3898.	1.2	6
15	<i>Man o' War</i> Mutation in UDP-α- <scp>d</scp> -Xylose Synthase Favors the Abortive Catalytic Cycle and Uncovers a Latent Potential for Hexamer Formation. Biochemistry, 2015, 54, 807-819.	1.2	3
16	Mechanism of proteasome gate modulation by assembly chaperones Pba1 and Pba2. Journal of Biological Chemistry, 2022, 298, 101906.	1.6	2
17	Resolution of Heterogeneity in Nicotinic Receptor Assembly by Cryo-EM. Biophysical Journal, 2018, 114, 10a-11a.	0.2	O
18	Cryo-EM Structure of the $\hat{l}\pm1\hat{l}^22\hat{l}^32$ GabaAReceptor in a Lipidic Environment. Biophysical Journal, 2020, 118, 582a-583a.	0.2	0