Zhihua Liu

List of Publications by Year
in descending order

11	1,018	1040056 9	1281871 11
papers	citations	h-index	g-index

11	11	11	1711
all docs	docs citations	times ranked	citing authors

#	Article	IF	Citations
1	CG14906 (mettl4) mediates m6A methylation of U2 snRNA in Drosophila. Cell Discovery, 2020, 6, 44.	6.7	35
2	Acsl, the Drosophila ortholog of intellectual disability-related ACSL4, inhibits synaptic growth by altered lipids. Journal of Cell Science, 2016, 129, 4034-4045.	2.0	14
3	dAcsl, theDrosophilaOrtholog of Acyl-CoA Synthetase Long-Chain Family Member 3 and 4, Inhibits Synapse Growth by Attenuating Bone Morphogenetic Protein Signaling via Endocytic Recycling. Journal of Neuroscience, 2014, 34, 2785-2796.	3.6	29
4	RIM Promotes Calcium Channel Accumulation at Active Zones of theDrosophilaNeuromuscular Junction. Journal of Neuroscience, 2012, 32, 16586-16596.	3.6	88
5	Drosophila FMRP regulates microtubule network formation and axonal transport of mitochondria. Human Molecular Genetics, 2011, 20, 51-63.	2.9	44
6	DrosophilaAcyl-CoA Synthetase Long-Chain Family Member 4 Regulates Axonal Transport of Synaptic Vesicles and Is Required for Synaptic Development and Transmission. Journal of Neuroscience, 2011, 31, 2052-2063.	3.6	37
7	Membrane-associated farnesylated UCH-L1 promotes Î±-synuclein neurotoxicity and is a therapeutic target for Parkinson's disease. Proceedings of the National Academy of Sciences of the United States of America, 2009, 106, 4635-4640.	7.1	121
8	DrosophilaTubulin-specific chaperone E functions at neuromuscular synapses and is required for microtubule network formation. Development (Cambridge), 2009, 136, 1571-1581.	2.5	48
9	Mutational Analysis Establishes a Critical Role for the N Terminus of Fragile X Mental Retardation Protein FMRP. Journal of Neuroscience, 2008, 28, 3221-3226.	3.6	25
10	Reply to "Does mapping reveal correlation between gene expression and proteinâ€“protein interaction?". Nature Genetics, 2003, 33, 16-17.	21.4	7
11	Correlation between transcriptome and interactome mapping data from Saccharomyces cerevisiae. Nature Genetics, 2001, 29, 482-486.	21.4	570