Amanda J Lewis

List of Publications by Year in descending order

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840776 1058476 1,309 14 11 14 citations h-index g-index papers 16 16 16 2989 docs citations times ranked citing authors all docs

#	Article	IF	Citations
1	Lewy pathology in Parkinson's disease consists of crowded organelles and lipid membranes. Nature Neuroscience, 2019, 22, 1099-1109.	14.8	604
2	Prion-like domains in RNA binding proteins are essential for building subnuclear paraspeckles. Journal of Cell Biology, 2015, 210, 529-539.	5.2	269
3	TDP-43 extracted from frontotemporal lobar degeneration subject brains displays distinct aggregate assemblies and neurotoxic effects reflecting disease progression rates. Nature Neuroscience, 2019, 22, 65-77.	14.8	143
4	The ins and outs of IncRNA structure: How, why and what comes next?. Biochimica Et Biophysica Acta - Gene Regulatory Mechanisms, 2016, 1859, 46-58.	1.9	71
5	Determinants of affinity and specificity in RNA-binding proteins. Current Opinion in Structural Biology, 2016, 38, 83-91.	5.7	51
6	The design and structural characterization of a synthetic pentatricopeptide repeat protein. Acta Crystallographica Section D: Biological Crystallography, 2015, 71, 196-208.	2.5	41
7	Cerebral Corpora amylacea are dense membranous labyrinths containing structurally preserved cell organelles. Scientific Reports, 2018, 8, 18046.	3.3	21
8	Imaging of post-mortem human brain tissue using electron and X-ray microscopy. Current Opinion in Structural Biology, 2019, 58, 138-148.	5.7	20
9	Caveolinâ€1â€driven membrane remodelling regulates hnRNPKâ€mediated exosomal microRNA sorting in cancer. Clinical and Translational Medicine, 2021, 11, e381.	4.0	19
10	Intracellular speciation of gold nanorods alters the conformational dynamics of genomic DNA. Nature Nanotechnology, 2018, 13, 1148-1153.	31.5	16
11	FTLDâ€₹DP assemblies seed neoaggregates with subtypeâ€specific features via a prionâ€ike cascade. EMBO Reports, 2021, 22, e53877.	4.5	14
12	The yeast transcription elongation factor Spt4/5 is a sequenceâ€specific RNA binding protein. Protein Science, 2016, 25, 1710-1721.	7.6	11
13	The first transcriptomes from field-collected individual whiteflies (Bemisia tabaci, Hemiptera:) Tj ETQq $1\ 1\ 0.7843$	14 rgBT /(1.1	Overlock 10 T
14	Ubiquitin fusion constructs allow the expression and purification of multi-KOW domain complexes of the Saccharomyces cerevisiae transcription elongation factor Spt4/5. Protein Expression and Purification, 2014, 100, 54-60.	1.3	2