## Pankaj Kumar Singh

List of Publications by Year in descending order

Source: https://exaly.com/author-pdf/2215649/publications.pdf

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933447 1281871 11 447 10 11 g-index citations h-index papers 12 12 12 2808 docs citations times ranked citing authors all docs

#	Article	IF	CITATIONS
1	Glycogen synthase protects neurons from cytotoxicity of mutant huntingtin by enhancing the autophagy flux. Cell Death and Disease, 2018, 9, 201.	6.3	29
2	Cerebellar Ataxia and Coenzyme Q Deficiency through Loss of Unorthodox Kinase Activity. Molecular Cell, 2016, 63, 608-620.	9.7	101
3	Coumarin–pyrene conjugate: Synthesis, structure and Cu-selective fluorescent sensing in mammalian kidney cells. Journal of Luminescence, 2016, 171, 159-165.	3.1	29
4	Autophagy Defects and Lafora Disease. , 2016, , 187-195.		0
5	Interdependence of laforin and malin proteins for their stability and functions could underlie the molecular basis of locus heterogeneity in Lafora disease. Journal of Biosciences, 2015, 40, 863-871.	1.1	11
6	Changing Shapes of Glycogenââ,¬â€œAutophagy Nexus in Neurons: Perspective from a Rare Epilepsy. Frontiers in Neurology, 2015, 6, 14.	2.4	16
7	Decreased O-Linked GlcNAcylation Protects from Cytotoxicity Mediated by Huntingtin Exon1 Protein Fragment. Journal of Biological Chemistry, 2014, 289, 13543-13553.	3.4	54
8	Activation of serum/glucocorticoid-induced kinase 1 (SGK1) underlies increased glycogen levels, mTOR activation, and autophagy defects in Lafora disease. Molecular Biology of the Cell, 2013, 24, 3776-3786.	2.1	39
9	The Laforin-Malin Complex Negatively Regulates Glycogen Synthesis by Modulating Cellular Glucose Uptake via Glucose Transporters. Molecular and Cellular Biology, 2012, 32, 652-663.	2.3	41
10	Lafora disease E3 ubiquitin ligase malin is recruited to the processing bodies and regulates the microRNA-mediated gene silencing process via the decapping enzyme Dcp1a. RNA Biology, 2012, 9, 1440-1449.	3.1	20
11	The malin–laforin complex suppresses the cellular toxicity of misfolded proteins by promoting their degradation through the ubiquitin–proteasome system. Human Molecular Genetics, 2009, 18, 688-700.	2.9	106