Venkatesh Mallikarjun

List of Publications by Year in descending order

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932766 1125271 13 542 10 13 citations g-index h-index papers 22 22 22 1059 docs citations times ranked citing authors all docs

#	Article	IF	CITATIONS
1	Circadian control of the secretory pathway maintains collagen homeostasis. Nature Cell Biology, 2020, 22, 74-86.	4.6	130
2	Monitoring Intracellular Redox Potential Changes Using SERS Nanosensors. ACS Nano, 2012, 6, 888-896.	7.3	90
3	Regulation of Lifespan by the Mitochondrial Electron Transport Chain: Reactive Oxygen Species-Dependent and Reactive Oxygen Species-Independent Mechanisms. Antioxidants and Redox Signaling, 2013, 19, 1953-1969.	2.5	59
4	Nuclear decoupling is part of a rapid protein-level cellular response to high-intensity mechanical loading. Nature Communications, 2019, 10, 4149.	5.8	58
5	MicroRNA-dependent regulation of biomechanical genes establishes tissue stiffness homeostasis. Nature Cell Biology, 2019, 21, 348-358.	4.6	44
6	Cellular redox potential and the biomolecular electrochemical series: A systems hypothesis. Free Radical Biology and Medicine, 2012, 53, 280-288.	1.3	38
7	Laser capture microdissection coupled mass spectrometry (LCM-MS) for spatially resolved analysis of formalin-fixed and stained human lung tissues. Clinical Proteomics, 2020, 17, 24.	1.1	37
8	The interplay between mitochondrial protein and iron homeostasis and its possible role in ageing. Experimental Gerontology, 2014, 56, 123-134.	1.2	17
9	Circadian time series proteomics reveals daily dynamics in cartilage physiology. Osteoarthritis and Cartilage, 2021, 29, 739-749.	0.6	17
10	Therapeutic Manipulation of Ageing: Repurposing Old Dogs and Discovering New Tricks. EBioMedicine, 2016, 14, 24-31.	2.7	15
11	BayesENproteomics: Bayesian Elastic Nets for Quantification of Peptidoforms in Complex Samples. Journal of Proteome Research, 2020, 19, 2167-2184.	1.8	9
12	Peptide location fingerprinting reveals modificationâ€associated biomarker candidates of ageing in human tissue proteomes. Aging Cell, 2021, 20, e13355.	3.0	9
13	BiolD-based proteomic analysis of the Bid interactome identifies novel proteins involved in cell-cycle-dependent apoptotic priming. Cell Death and Disease, 2020, 11, 872.	2.7	6