Uros Kuzmanov

List of Publications by Year in descending order

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623734 642732 23 704 14 23 h-index citations g-index papers 24 24 24 1177 docs citations times ranked citing authors all docs

#	Article	IF	CITATIONS
1	Membrane proteomic profiling of the heart: past, present, and future. American Journal of Physiology - Heart and Circulatory Physiology, 2021, 320, H417-H423.	3.2	5
2	An organ-on-a-chip model for pre-clinical drug evaluation in progressive non-genetic cardiomyopathy. Journal of Molecular and Cellular Cardiology, 2021, 160, 97-110.	1.9	23
3	Mapping signalling perturbations in myocardial fibrosis via the integrative phosphoproteomic profiling of tissue from diverse sources. Nature Biomedical Engineering, 2020, 4, 889-900.	22.5	17
4	Bioinformatic analysis of membrane and associated proteins in murine cardiomyocytes and human myocardium. Scientific Data, 2020, 7, 425.	5.3	8
5	EPIC: software toolkit for elution profile-based inference of protein complexes. Nature Methods, 2019, 16, 737-742.	19.0	67
6	AKAP6 and phospholamban colocalize and interact in HEKâ€293T cells and primary murine cardiomyocytes. Physiological Reports, 2019, 7, e14144.	1.7	4
7	Lipidomic profiling of subchronic As4S4 exposure identifies inflammatory mediators as sensitive biomarkers in rats. Metallomics, 2019, 11, 576-585.	2.4	10
8	Nanoscale reorganization of sarcoplasmic reticulum in pressure-overload cardiac hypertrophy visualized by dSTORM. Scientific Reports, 2019, 9, 7867.	3.3	15
9	A strategy for the metabolomics-based screening of active constituents and quality consistency control for natural medicinal substance toad venom. Analytica Chimica Acta, 2018, 1031, 108-118.	5.4	13
10	Avoiding false discovery in biomarker research. BMC Biochemistry, 2016, 17, 17.	4.4	6
11	Global phosphoproteomic profiling reveals perturbed signaling in a mouse model of dilated cardiomyopathy. Proceedings of the National Academy of Sciences of the United States of America, 2016, 113, 12592-12597.	7.1	35
12	Phosphoproteomic network analysis in the sea urchin <i>Strongylocentrotus purpuratus</i> reveals new candidates in egg activation. Proteomics, 2015, 15, 4080-4095.	2.2	24
13	Large-scale label-free phosphoproteomics: from technology to data interpretation. Bioanalysis, 2014, 6, 2403-2420.	1.5	8
14	Semiquantitative proteomic analysis of human hippocampal tissues from Alzheimer's disease and age-matched control brains. Clinical Proteomics, 2013, 10, 5.	2.1	67
15	The sweet and sour of serological glycoprotein tumor biomarker quantification. BMC Medicine, 2013, 11, 31.	5.5	67
16	Copy Number and Expression Alterations of miRNAs in the Ovarian Cancer Cell Line OVCAR-3: Impact on Kallikrein 6 Protein Expression. Clinical Chemistry, 2013, 59, 296-305.	3.2	15
17	Fuzzy decision support system for ship lock control. Expert Systems With Applications, 2013, 40, 3953-3960.	7.6	41
18	Protein-protein interaction networks: probing disease mechanisms using model systems. Genome Medicine, 2013, 5, 37.	8.2	124

#	Article	IF	CITATION
19	Using phosphoproteomics to monitor disregulated signaling networks in cardiac disease preceding heart failure. Bioanalysis, 2013, 5, 2863-2866.	1.5	3
20	Glycoproteomic identification of potential glycoprotein biomarkers in ovarian cancer proximal fluids. Clinical Chemistry and Laboratory Medicine, 2013, 51, 1467-76.	2.3	26
21	Separation of kallikrein 6 glycoprotein subpopulations in biological fluids by anionâ€exchange chromatography coupled to ELISA and identification by mass spectrometry. Proteomics, 2012, 12, 799-809.	2.2	19
22	Differential N-glycosylation of Kallikrein 6 Derived from Ovarian Cancer Cells or the Central Nervous System. Molecular and Cellular Proteomics, 2009, 8, 791-798.	3.8	60
23	Disruption of the Mthfd1 Gene Reveals a Monofunctional 10-Formyltetrahydrofolate Synthetase in Mammalian Mitochondria. Journal of Biological Chemistry, 2005, 280, 7597-7602.	3.4	47