

Mitra Mastali

List of Publications by Year in descending order

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Version: 2024-02-01

11
papers

258
citations

1163117

8
h-index

1372567

10
g-index

11
all docs

11
docs citations

11
times ranked

622
citing authors

#	ARTICLE	IF	CITATIONS
1	Sex-based differences in remote monitoring of biometric, psychometric and biomarker indices in stable ischemic heart disease. <i>Biology of Sex Differences</i> , 2022, 13, 15.	4.1	1
2	Retina mirrors brain pathology and response to GA immunotherapy in advanced stage AD-model mice.. <i>Alzheimer's and Dementia</i> , 2021, 17 Suppl 3, e055329.	0.8	0
3	Feasibility of Patient-Centric Remote Dried Blood Sampling: The Prediction, Risk, and Evaluation of Major Adverse Cardiac Events (PRE-MACE) Study. <i>Biodemography and Social Biology</i> , 2020, 65, 313-322.	1.0	7
4	Parallels between retinal and brain pathology and response to immunotherapy in old, late-stage Alzheimer's disease mouse models. <i>Aging Cell</i> , 2020, 19, e13246.	6.7	32
5	Identification of Putative Early Atherosclerosis Biomarkers by Unsupervised Deconvolution of Heterogeneous Vascular Proteomes. <i>Journal of Proteome Research</i> , 2020, 19, 2794-2806.	3.7	16
6	Quality Control and Outlier Detection of Targeted Mass Spectrometry Data from Multiplex Protein Panels. <i>Journal of Proteome Research</i> , 2020, 19, 2278-2293.	3.7	12
7	A protocol integrating remote patient monitoring patient reported outcomes and cardiovascular biomarkers. <i>Npj Digital Medicine</i> , 2019, 2, 84.	10.9	12
8	Serum NfL (Neurofilament Light Chain) Levels and Incident Stroke in Adults With Diabetes Mellitus. <i>Stroke</i> , 2019, 50, 1669-1675.	2.0	60
9	Acute neuropathological consequences of short-term mechanical ventilation in wild-type and Alzheimer's disease mice. <i>Critical Care</i> , 2019, 23, 63.	5.8	21
10	Highly Reproducible Automated Proteomics Sample Preparation Workflow for Quantitative Mass Spectrometry. <i>Journal of Proteome Research</i> , 2018, 17, 420-428.	3.7	68
11	The continuing evolution of cardiac troponin I biomarker analysis: from protein to proteoform. <i>Expert Review of Proteomics</i> , 2017, 14, 973-986.	3.0	29