

Abidali Mohamedali

List of Publications by Year in descending order

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Version: 2024-02-01

25
papers

639
citations

840776

11
h-index

677142

22
g-index

26
all docs

26
docs citations

26
times ranked

1309
citing authors

#	ARTICLE	IF	CITATIONS
1	A Bioinformatics Approach to Mine the Microbial Proteomic Profile of COVID-19 Mass Spectrometry Data. <i>Applied Microbiology</i> , 2022, 2, 150-164.	1.6	3
2	Mass spectrometryâ€‘based protein identification in proteomicsâ€‘a review. <i>Briefings in Bioinformatics</i> , 2021, 22, 1620-1638.	6.5	55
3	Use of a Recombinant Biomarker Protein DDA Library Increases DIA Coverage of Low Abundance Plasma Proteins. <i>Journal of Proteome Research</i> , 2021, 20, 2374-2389.	3.7	6
4	Proteomic investigations into resistance in colorectal cancer. <i>Expert Review of Proteomics</i> , 2020, 17, 49-65.	3.0	8
5	Identification of Proteins From Proteomic Analysis. , 2019, , 855-870.		1
6	Quantification of Proteins From Proteomic Analysis. , 2019, , 871-890.		1
7	Proteomics Reveals Cellâ€‘Surface Urokinase Plasminogen Activator Receptor Expression Impacts Most Hallmarks of Cancer. <i>Proteomics</i> , 2019, 19, e1900026.	2.2	9
8	Potential early clinical stage colorectal cancer diagnosis using a proteomics blood test panel. <i>Clinical Proteomics</i> , 2019, 16, 34.	2.1	44
9	Oncoproteomics: Current status and future opportunities. <i>Clinica Chimica Acta</i> , 2019, 495, 611-624.	1.1	20
10	Looking for Missing Proteins. , 2019, , .		2
11	Accelerating the search for the missing proteins in the human proteome. <i>Nature Communications</i> , 2017, 8, 14271.	12.8	86
12	De Novo Peptide Sequencing: Deep Mining of High-Resolution Mass Spectrometry Data. <i>Methods in Molecular Biology</i> , 2017, 1549, 119-134.	0.9	10
13	A Systematic Bioinformatics Approach to Identify High Quality Mass Spectrometry Data and Functionally Annotate Proteins and Proteomes. <i>Methods in Molecular Biology</i> , 2017, 1549, 163-176.	0.9	3
14	Human Prestin: A Candidate PE1 Protein Lacking Stringent Mass Spectrometric Evidence?. <i>Journal of Proteome Research</i> , 2017, 16, 4531-4535.	3.7	6
15	Effects of Acute and Chronic Biomechanical Strain on Human Cerebral Endothelial Cells in Altering their Proteome Profile. <i>Current Proteomics</i> , 2017, 14, .	0.3	2
16	Mechanical stretch: physiological and pathological implications for human vascular endothelial cells. <i>Vascular Cell</i> , 2015, 7, 8.	0.2	185
17	A novel multiplexed immunoassay identifies CEA, IL-8 and prolactin as prospective markers for Dukesâ€‘™ stages A-D colorectal cancers. <i>Clinical Proteomics</i> , 2015, 12, 10.	2.1	33
18	Transforming growth factor-Î², MAPK and Wnt signaling interactions in colorectal cancer. <i>EuPA Open Proteomics</i> , 2015, 8, 104-115.	2.5	31

#	ARTICLE	IF	CITATIONS
19	Epithelial and Stromal Cell Urokinase Plasminogen Activator Receptor Expression Differentially Correlates with Survival in Rectal Cancer Stages B and C Patients. PLoS ONE, 2015, 10, e0117786.	2.5	12
20	Correlations between Integrin $\alpha 5 \beta 1$ Expression and Clinico-Pathological Features in Stage B and Stage C Rectal Cancer. PLoS ONE, 2014, 9, e97248.	2.5	10
21	Characterization of the Interaction between Heterodimeric $\alpha 5 \beta 1$ Integrin and Urokinase Plasminogen Activator Receptor (uPAR) Using Functional Proteomics. Journal of Proteome Research, 2014, 13, 5956-5964.	3.7	18
22	An improved method for the detection and enrichment of low-abundant membrane and lipid raft-residing proteins. Journal of Proteomics, 2013, 79, 299-304.	2.4	10
23	Ultradepletion of Human Plasma using Chicken Antibodies: A Proof of Concept Study. Journal of Proteome Research, 2013, 12, 2399-2413.	3.7	25
24	Unlocking the Puzzling Biology of the Black Périgord Truffle <i>Tuber melanosporum</i> . Journal of Proteome Research, 2013, 12, 5349-5356.	3.7	24
25	A novel, cost-effective and efficient chicken egg IgY purification procedure. Journal of Immunological Methods, 2012, 380, 73-76.	1.4	35