

Aafaque Ahmad Khan

List of Publications by Year in descending order

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Version: 2024-02-01

21
papers

2,899
citations

566801

15
h-index

713013

21
g-index

21
all docs

21
docs citations

21
times ranked

6865
citing authors

#	ARTICLE	IF	CITATIONS
1	Integrative analysis of non-small cell lung cancer patient-derived xenografts identifies distinct proteotypes associated with patient outcomes. <i>Nature Communications</i> , 2022, 13, 1811.	5.8	21
2	Multi-Omics Analysis to Characterize Cigarette Smoke Induced Molecular Alterations in Esophageal Cells. <i>Frontiers in Oncology</i> , 2020, 10, 1666.	1.3	1
3	Mutational Landscape of Esophageal Squamous Cell Carcinoma in an Indian Cohort. <i>Frontiers in Oncology</i> , 2020, 10, 1457.	1.3	21
4	Intracranial Aneurysm Biomarker Candidates Identified by a Proteome-Wide Study. <i>OMICS A Journal of Integrative Biology</i> , 2020, 24, 483-492.	1.0	14
5	Comprehensive network map of interferon gamma signaling. <i>Journal of Cell Communication and Signaling</i> , 2018, 12, 745-751.	1.8	67
6	Data from quantitative proteomic analysis of lung adenocarcinoma and squamous cell carcinoma primary tissues using high resolution mass spectrometry. <i>Data in Brief</i> , 2018, 19, 1631-1637.	0.5	1
7	A network map of thrombopoietin signaling. <i>Journal of Cell Communication and Signaling</i> , 2018, 12, 737-743.	1.8	12
8	Chronic Cigarette Smoke Mediated Global Changes in Lung Mucoepidermoid Cells: A Phosphoproteomic Analysis. <i>OMICS A Journal of Integrative Biology</i> , 2017, 21, 474-487.	1.0	38
9	Altered signaling associated with chronic arsenic exposure in human skin keratinocytes. <i>Proteomics - Clinical Applications</i> , 2017, 11, 1700004.	0.8	2
10	Long-Term Cigarette Smoke Exposure and Changes in MiRNA Expression and Proteome in Non-Small-Cell Lung Cancer. <i>OMICS A Journal of Integrative Biology</i> , 2017, 21, 390-403.	1.0	24
11	Phosphotyrosine profiling of curcumin-induced signaling. <i>Clinical Proteomics</i> , 2016, 13, 13.	1.1	19
12	A pathway map of glutamate metabolism. <i>Journal of Cell Communication and Signaling</i> , 2016, 10, 69-75.	1.8	115
13	A network map of Interleukin-10 signaling pathway. <i>Journal of Cell Communication and Signaling</i> , 2016, 10, 61-67.	1.8	85
14	LC-MS-based serum metabolomic analysis reveals dysregulation of phosphatidylcholines in esophageal squamous cell carcinoma. <i>Journal of Proteomics</i> , 2015, 127, 96-102.	1.2	38
15	Phosphotyrosine profiling identifies ephrin receptor A2 as a potential therapeutic target in esophageal squamous cell carcinoma. <i>Proteomics</i> , 2015, 15, 374-382.	1.3	38
16	A knowledgebase resource for interleukin-17 family mediated signaling. <i>Journal of Cell Communication and Signaling</i> , 2015, 9, 291-296.	1.8	25
17	Plasma Proteome Database as a resource for proteomics research: 2014 update. <i>Nucleic Acids Research</i> , 2014, 42, D959-D965.	6.5	273
18	Pancreatic Cancer Database. <i>Cancer Biology and Therapy</i> , 2014, 15, 963-967.	1.5	57

#	ARTICLE	IF	CITATIONS
19	Signaling Network Map of Endothelial TEK Tyrosine Kinase. Journal of Signal Transduction, 2014, 2014, 1-6.	2.0	46
20	A draft map of the human proteome. Nature, 2014, 509, 575-581.	13.7	1,948
21	A multicellular signal transduction network of AGE/RAGE signaling. Journal of Cell Communication and Signaling, 2013, 7, 19-23.	1.8	54