Aafaque Ahmad Khan

List of Publications by Year
in descending order

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	A draft map of the human proteome. Nature, 2014, 509, 575-581.	13.7	1,948
2	Plasma Proteome Database as a resource for proteomics research: 2014 update. Nucleic Acids Research, 2014, 42, D959-D965.	6.5	273
3	A pathway map of glutamate metabolism. Journal of Cell Communication and Signaling, 2016, 10, 69-75.	1.8	115
4	A network map of Interleukin-10 signaling pathway. Journal of Cell Communication and Signaling, 2016, 10, 61-67.	1.8	85
5	Comprehensive network map of interferon gamma signaling. Journal of Cell Communication and Signaling, 2018, 12, 745-751.	1.8	67
6	Pancreatic Cancer Database. Cancer Biology and Therapy, 2014, 15, 963-967.	1.5	57
7	A multicellular signal transduction network of AGE/RAGE signaling. Journal of Cell Communication and Signaling, 2013, 7, 19-23.	1.8	54
8	Signaling Network Map of Endothelial TEK Tyrosine Kinase. Journal of Signal Transduction, 2014, 2014, 1-6.	2.0	46
9	LC–MS-based serum metabolomic analysis reveals dysregulation of phosphatidylcholines in esophageal squamous cell carcinoma. Journal of Proteomics, 2015, 127, 96-102.	1.2	38
10	Phosphotyrosine profiling identifies ephrin receptor A2 as a potential therapeutic target in esophageal squamous cell carcinoma. Proteomics, 2015, 15, 374-382.	1.3	38
11	Chronic Cigarette Smoke Mediated Global Changes in Lung Mucoepidermoid Cells: A Phosphoproteomic Analysis. OMICS A Journal of Integrative Biology, 2017, 21, 474-487.	1.0	38
12	A knowledgebase resource for interleukin-17 family mediated signaling. Journal of Cell Communication and Signaling, 2015, 9, 291-296.	1.8	25
13	Long-Term Cigarette Smoke Exposure and Changes in MiRNA Expression and Proteome in Non-Small-Cell Lung Cancer. OMICS A Journal of Integrative Biology, 2017, 21, 390-403.	1.0	24
14	Mutational Landscape of Esophageal Squamous Cell Carcinoma in an Indian Cohort. Frontiers in Oncology, 2020, 10, 1457.	1.3	21
15	Integrative analysis of non-small cell lung cancer patient-derived xenografts identifies distinct proteotypes associated with patient outcomes. Nature Communications, 2022, 13, 1811.	5.8	21
16	Phosphotyrosine profiling of curcumin-induced signaling. Clinical Proteomics, 2016, 13, 13.	1.1	19
17	Intracranial Aneurysm Biomarker Candidates Identified by a Proteome-Wide Study. OMICS A Journal of Integrative Biology, 2020, 24, 483-492.	1.0	14
18	A network map of thrombopoietin signaling. Journal of Cell Communication and Signaling, 2018, 12, 737-743.	1.8	12

#	Article	IF	Citations
19	Altered signaling associated with chronic arsenic exposure in human skin keratinocytes. Proteomics - Clinical Applications, 2017, 11, 1700004.	0.8	2
20	Data from quantitative proteomic analysis of lung adenocarcinoma and squamous cell carcinoma primary tissues using high resolution mass spectrometry. Data in Brief, 2018, 19, 1631-1637.	0.5	1
21	Multi-Omics Analysis to Characterize Cigarette Smoke Induced Molecular Alterations in Esophageal Cells. Frontiers in Oncology, 2020, 10, 1666.	1.3	1