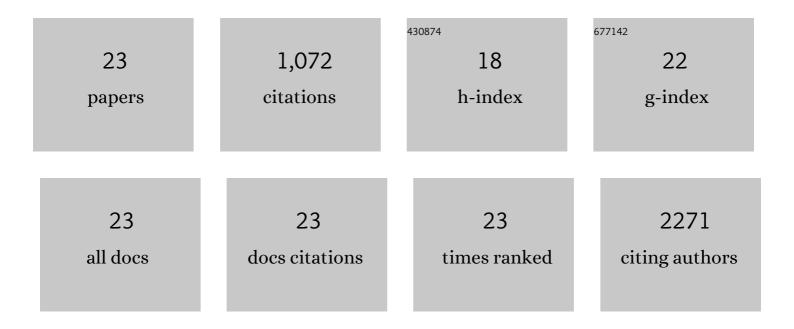
Mohit Raja Jain

List of Publications by Year in descending order

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Μομιτ Ρλιλ Ιλινι

#	Article	IF	CITATIONS
1	MicroRNA-7 activates Nrf2 pathway by targeting Keap1 expression. Free Radical Biology and Medicine, 2015, 89, 548-556.	2.9	116
2	Redox Regulatory Mechanism of Transnitrosylation by Thioredoxin. Molecular and Cellular Proteomics, 2010, 9, 2262-2275.	3.8	115
3	Mst1 Promotes Cardiac Myocyte Apoptosis through Phosphorylation and Inhibition of Bcl-xL. Molecular Cell, 2014, 54, 639-650.	9.7	110
4	MicroRNA-7 Protects against 1-Methyl-4-Phenylpyridinium-Induced Cell Death by Targeting RelA. Journal of Neuroscience, 2014, 34, 12725-12737.	3.6	85
5	Distinction of thioredoxin transnitrosylation and denitrosylation target proteins by the ICAT quantitative approach. Journal of Proteomics, 2011, 74, 2498-2509.	2.4	67
6	Proteomic identification of novel targets regulated by the mammalian target of rapamycin pathway during oligodendrocyte differentiation. Glia, 2011, 59, 1754-1769.	4.9	60
7	Profiling the <i>Aspergillus fumigatus</i> Proteome in Response to Caspofungin. Antimicrobial Agents and Chemotherapy, 2011, 55, 146-154.	3.2	57
8	Toxicogenomics of endoplasmic reticulum stress inducer tunicamycin in the small intestine and liver of Nrf2 knockout and C57BL/6J mice. Toxicology Letters, 2007, 168, 21-39.	0.8	56
9	Pharmacogenomics of Phenolic Antioxidant Butylated Hydroxyanisole (BHA) in the Small Intestine and Liver of Nrf2 Knockout and C57BL/6J Mice. Pharmaceutical Research, 2006, 23, 2621-2637.	3.5	55
10	Externalized Glycolytic Enzymes Are Novel, Conserved, and Early Biomarkers of Apoptosis. Journal of Biological Chemistry, 2012, 287, 10325-10343.	3.4	46
11	The Proteomic Signature of Aspergillus fumigatus During Early Development. Molecular and Cellular Proteomics, 2011, 10, M111.010108.	3.8	37
12	Myelin Lipid Abnormalities in the Aspartoacylase-Deficient Tremor Rat. Neurochemical Research, 2009, 34, 138-148.	3.3	36
13	Quantitative Proteomic Analysis of Formalin Fixed Paraffin Embedded Oral HPV Lesions from HIV Patients. The Open Proteomics Journal, 2008, 1, 40-45.	0.4	36
14	Pharmacokinetics, Pharmacodynamics and Drug Metabolism. Journal of Pharmaceutical Sciences, 2008, 97, 4528-4545.	3.3	35
15	Identification of novel S-nitrosation sites in soluble guanylyl cyclase, the nitric oxide receptor. Journal of Proteomics, 2016, 138, 40-47.	2.4	28
16	Altered proteolytic events in experimental autoimmune encephalomyelitis discovered by iTRAQ shotgun proteomics analysis of spinal cord. Proteome Science, 2009, 7, 25.	1.7	27
17	Proteomic Identification of Immunoproteasome Accumulation in Formalin-Fixed Rodent Spinal Cords with Experimental Autoimmune Encephalomyelitis. Journal of Proteome Research, 2012, 11, 1791-1803.	3.7	27
18	Ciliary neurotrophic factor (CNTF) plus soluble CNTF receptor α increases cyclooxygenase-2 expression, PGE2release and interferon-γ-induced CD40 in murine microglia. Journal of Neuroinflammation, 2009, 6, 7.	7.2	24

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19	Host Biomarkers of Invasive Pulmonary Aspergillosis To Monitor Therapeutic Response. Antimicrobial Agents and Chemotherapy, 2014, 58, 3373-3378.	3.2	18
20	Identification of Novel Nuclear Targets of Human Thioredoxin 1. Molecular and Cellular Proteomics, 2014, 13, 3507-3518.	3.8	17
21	Master redox regulator Trx1 upregulates SMYD1 & modulates lysine methylation. Biochimica Et Biophysica Acta - Proteins and Proteomics, 2015, 1854, 1816-1822.	2.3	11
22	A multidimensional approach to an in-depth proteomics analysis of transcriptional regulators in neuroblastoma cells. Journal of Neuroscience Methods, 2013, 216, 118-127.	2.5	8
23	iTRAQ Proteomics Profiling of Regulatory Proteins During Oligodendrocyte Differentiation. Neuromethods, 2012, , 119-138.	0.3	1