## Hamid Mirzaei

List of Publications by Year in descending order

Source: https://exaly.com/author-pdf/6444/publications.pdf

Version: 2024-02-01

22 papers 1,401 citations

394421 19 h-index 677142 22 g-index

24 all docs

24 docs citations

times ranked

24

2792 citing authors

#	Article	IF	CITATIONS
1	A Gain-of-Function Mutation in DHT Synthesis in Castration-Resistant Prostate Cancer. Cell, 2013, 154, 1074-1084.	28.9	257
2	Inert and seed-competent tau monomers suggest structural origins of aggregation. ELife, 2018, 7, .	6.0	183
3	Comparative Evaluation of Current Peptide Production Platforms Used in Absolute Quantification in Proteomics. Molecular and Cellular Proteomics, 2008, 7, 813-823.	3.8	91
4	Confetti: A Multiprotease Map of the HeLa Proteome for Comprehensive Proteomics. Molecular and Cellular Proteomics, 2014, 13, 1573-1584.	3.8	89
5	Fucosylation and protein glycosylation create functional receptors for cholera toxin. ELife, 2015, 4, e09545.	6.0	81
6	Yeast Ataxin-2 Forms an Intracellular Condensate Required for the Inhibition of TORC1 Signaling during Respiratory Growth. Cell, 2019, 177, 697-710.e17.	28.9	73
7	Type VI Secretion System Toxins Horizontally Shared between Marine Bacteria. PLoS Pathogens, 2015, 11, e1005128.	4.7	71
8	Discovery of tumor-specific irreversible inhibitors of stearoyl CoA desaturase. Nature Chemical Biology, 2016, 12, 218-225.	8.0	64
9	Hydralazine induces stress resistance and extends C. elegans lifespan by activating the NRF2/SKN-1 signalling pathway. Nature Communications, 2017, 8, 2223.	12.8	62
10	Stable association of RNAi machinery is conserved between the cytoplasm and nucleus of human cells. Rna, 2016, 22, 1085-1098.	3.5	56
11	Cloud CPFP: A Shotgun Proteomics Data Analysis Pipeline Using Cloud and High Performance Computing. Journal of Proteome Research, 2012, 11, 6282-6290.	3.7	52
12	Detection of FGF15 in Plasma by Stable Isotope Standards and Capture by Anti-peptide Antibodies and Targeted Mass Spectrometry. Cell Metabolism, 2015, 21, 898-904.	16.2	51
13	Abnormal mechanosensing and cofilin activation promote the progression of ascending aortic aneurysms in mice. Science Signaling, 2015, 8, ra105.	3.6	43
14	Characterizing the connectivity of poly-ubiquitin chains by selected reaction monitoring mass spectrometry. Molecular BioSystems, 2010, 6, 2004.	2.9	35
15	Hydralazine targets cAMP-dependent protein kinase leading to sirtuin 1/5 activation and lifespan extension in C. elegans. Nature Communications, 2019, 10, 4905.	12.8	33
16	Age-Onset Phosphorylation of a Minor Actin Variant Promotes Intestinal Barrier Dysfunction. Developmental Cell, 2019, 51, 587-601.e7.	7.0	33
17	Systematic measurement of transcription factor-DNA interactions by targeted mass spectrometry identifies candidate gene regulatory proteins. Proceedings of the National Academy of Sciences of the United States of America, 2013, 110, 3645-3650.	7.1	31
18	A GTPase-activating protein–binding protein (G3BP1)/antiviral protein relay conveys arteriosclerotic Wnt signals in aortic smooth muscle cells. Journal of Biological Chemistry, 2018, 293, 7942-7968.	3.4	24

#	Article	IF	CITATION
19	Halogenated Peptides as Internal Standards (H-PINS). Molecular and Cellular Proteomics, 2009, 8, 1934-1946.	3.8	23
20	<scp>GOAT</scp> – A simple <scp>LC</scp> â€ <scp>MS</scp> / <scp>MS</scp> gradient optimization tool. Proteomics, 2014, 14, 1467-1471.	2.2	17
21	OSR1 regulates a subset of inward rectifier potassium channels via a binding motif variant. Proceedings of the National Academy of Sciences of the United States of America, 2018, 115, 3840-3845.	7.1	17
22	Exocyst protein subnetworks integrate Hippo and mTOR signaling to promote virus detection and cancer. Cell Reports, 2021, 36, 109491.	6.4	11