

Holger Alexander Ebhardt

List of Publications by Year in descending order

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Version: 2024-02-01

15
papers

854
citations

933447

10
h-index

1058476

14
g-index

15
all docs

15
docs citations

15
times ranked

2047
citing authors

#	ARTICLE	IF	CITATIONS
1	Breast Cancer Classification Based on Proteotypes Obtained by SWATH Mass Spectrometry. <i>Cell Reports</i> , 2019, 28, 832-843.e7.	6.4	72
2	Platelet-Derived Microparticles From Obese Individuals: Characterization of Number, Size, Proteomics, and Crosstalk With Cancer and Endothelial Cells. <i>Frontiers in Pharmacology</i> , 2019, 10, 7.	3.5	44
3	Differences in the bovine milk whey proteome between early pregnancy and the estrous cycle. <i>Theriogenology</i> , 2018, 114, 301-307.	2.1	13
4	Structure-based assessment and network analysis of targeting 14-3-3 proteins in prostate cancer. <i>Molecular Cancer</i> , 2018, 17, 156.	19.2	6
5	A two-drug combination simulation study for metastatic castrate resistant prostate cancer. <i>Prostate</i> , 2018, 78, 1196-1200.	2.3	2
6	Systems pharmacology using mass spectrometry identifies critical response nodes in prostate cancer. <i>Npj Systems Biology and Applications</i> , 2018, 4, 26.	3.0	13
7	Comprehensive proteome analysis of human skeletal muscle in cachexia and sarcopenia: a pilot study. <i>Journal of Cachexia, Sarcopenia and Muscle</i> , 2017, 8, 567-582.	7.3	47
8	Proteome-wide association studies identify biochemical modules associated with a wing-size phenotype in <i>Drosophila melanogaster</i> . <i>Nature Communications</i> , 2016, 7, 12649.	12.8	41
9	Applications of targeted proteomics in systems biology and translational medicine. <i>Proteomics</i> , 2015, 15, 3193-3208.	2.2	182
10	Enzymatic generation of peptides flanked by basic amino acids to obtain MS/MS spectra with 2Å– sequence coverage. <i>Rapid Communications in Mass Spectrometry</i> , 2014, 28, 2735-2743.	1.5	8
11	CIG-P: Circular Interaction Graph for Proteomics. <i>BMC Bioinformatics</i> , 2014, 15, 344.	2.6	0
12	A repository of assays to quantify 10,000 human proteins by SWATH-MS. <i>Scientific Data</i> , 2014, 1, 140031.	5.3	370
13	Selected Reaction Monitoring Mass Spectrometry: A Methodology Overview. <i>Methods in Molecular Biology</i> , 2014, 1072, 209-222.	0.9	18
14	Isolation and Biochemical Analysis of Plant Small RNAs. <i>Methods in Molecular Biology</i> , 2012, 894, 223-239.	0.9	1
15	Range of protein detection by selected/multiple reaction monitoring mass spectrometry in an unfractionated human cell culture lysate. <i>Proteomics</i> , 2012, 12, 1185-1193.	2.2	37