

Bridget Calder

List of Publications by Year in descending order

Source: <https://exaly.com/author-pdf/540915/publications.pdf>

Version: 2024-02-01

16
papers

360
citations

759233

12
h-index

940533

16
g-index

17
all docs

17
docs citations

17
times ranked

632
citing authors

#	ARTICLE	IF	CITATIONS
1	The Role of Tumor Microenvironment in Chemoresistance: 3D Extracellular Matrices as Accomplices. <i>International Journal of Molecular Sciences</i> , 2018, 19, 2861.	4.1	114
2	Identification of Quantitative Proteomic Differences between <i>Mycobacterium tuberculosis</i> Lineages with Altered Virulence. <i>Frontiers in Microbiology</i> , 2016, 7, 813.	3.5	34
3	A Human Lung Challenge Model to Evaluate the Safety and Immunogenicity of PPD and Live <i>Bacillus Calmette-Guérin</i> . <i>American Journal of Respiratory and Critical Care Medicine</i> , 2020, 201, 1277-1291.	5.6	28
4	Mass Spectrometry Offers Insight into the Role of Ser/Thr/Tyr Phosphorylation in the Mycobacteria. <i>Frontiers in Microbiology</i> , 2016, 7, 141.	3.5	26
5	Microbial function and genital inflammation in young South African women at high risk of HIV infection. <i>Microbiome</i> , 2020, 8, 165.	11.1	23
6	Quantitative label-free proteomic analysis of human urine to identify novel candidate protein biomarkers for schistosomiasis. <i>PLoS Neglected Tropical Diseases</i> , 2017, 11, e0006045.	3.0	22
7	Identification of Novel Physiological Substrates of <i>Mycobacterium bovis</i> BCG Protein Kinase G (PknG) by Label-free Quantitative Phosphoproteomics. <i>Molecular and Cellular Proteomics</i> , 2018, 17, 1365-1377.	3.8	21
8	Global proteome and phosphoproteome dynamics indicate novel mechanisms of vitamin C induced dormancy in <i>Mycobacterium smegmatis</i> . <i>Journal of Proteomics</i> , 2018, 180, 1-10.	2.4	16
9	Proteomic comparison of three clinical diarrhoeagenic drug-resistant <i>Escherichia coli</i> isolates grown on CHROMagar [®] , cSTEC media. <i>Journal of Proteomics</i> , 2018, 180, 25-35.	2.4	16
10	In silico verification and parallel reaction monitoring prevalidation of potential prostate cancer biomarkers. <i>Future Oncology</i> , 2016, 12, 43-57.	2.4	14
11	Associating H ₂ O ₂ and NO-related changes in the proteome of <i>Mycobacterium smegmatis</i> with enhanced survival in macrophage. <i>Emerging Microbes and Infections</i> , 2018, 7, 1-17.	6.5	13
12	Mycobacterial proteomics: analysis of expressed proteomes and post-translational modifications to identify candidate virulence factors. <i>Expert Review of Proteomics</i> , 2015, 12, 21-35.	3.0	12
13	Challenges and Opportunities for Biological Mass Spectrometry Core Facilities in the Developing World. <i>Journal of Biomolecular Techniques</i> , 2018, 29, 4-15.	1.5	8
14	Analysis of the regenerative capacity of human serum exosomes after a simple multistep separation from lipoproteins. <i>Journal of Tissue Engineering and Regenerative Medicine</i> , 2021, 15, 63-77.	2.7	7
15	Identification of novel salivary candidate protein biomarkers for tuberculosis diagnosis: A preliminary biomarker discovery study. <i>Tuberculosis</i> , 2021, 130, 102118.	1.9	5
16	Protocols for Preparation and Mass Spectrometry Analysis of Clinical Urine Samples to Identify Candidate Biomarkers of Schistosoma-Associated Bladder Cancer. <i>Methods in Molecular Biology</i> , 2021, 2292, 143-150.	0.9	1