

Jiaqi

List of Publications by Year in descending order

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Version: 2024-02-01

23
papers

672
citations

686830

13
h-index

713013

21
g-index

25
all docs

25
docs citations

25
times ranked

650
citing authors

#	ARTICLE	IF	CITATIONS
1	<i>Legionella pneumophila</i> modulates host energy metabolism by ADP-ribosylation of ADP/ATP translocases. <i>ELife</i> , 2022, 11, .	2.8	27
2	<i>Legionella pneumophila</i> temporally regulates the activity of ADP/ATP translocases by reversible ADP-ribosylation. , 2022, 1, 51-65.		7
3	<i>Coxiella burnetii</i> inhibits host immunity by a protein phosphatase adapted from glycolysis. <i>Proceedings of the National Academy of Sciences of the United States of America</i> , 2022, 119, .	3.3	158
4	<i>Bartonella</i> type IV secretion effector BepC induces stress fiber formation through activation of GEF-H1. <i>PLoS Pathogens</i> , 2021, 17, e1009065.	2.1	2
5	<i>Shigella</i> evades pyroptosis by arginine ADP-ribosylation of caspase-11. <i>Nature</i> , 2021, 599, 290-295.	13.7	93
6	<i>Legionella pneumophila</i> regulates the activity of UBE2N by deamidase-mediated deubiquitination. <i>EMBO Journal</i> , 2020, 39, e102806.	3.5	38
7	<i>Legionella</i> effector MavC targets the Ube2N-Ub conjugate for noncanonical ubiquitination. <i>Nature Communications</i> , 2020, 11, 2365.	5.8	21
8	Arginine GlcNAcylation of Rab small GTPases by the pathogen <i>Salmonella Typhimurium</i> . <i>Communications Biology</i> , 2020, 3, 287.	2.0	27
9	Structural insights into the mechanism and inhibition of transglutaminase-induced ubiquitination by the <i>Legionella</i> effector MavC. <i>Nature Communications</i> , 2020, 11, 1774.	5.8	15
10	<i>Salmonella</i> Proteomic Profiling during Infection Distinguishes the Intracellular Environment of Host Cells. <i>MSystems</i> , 2019, 4, .	1.7	20
11	Proteomic Analysis of FNR-Regulated Anaerobiosis in <i>Salmonella Typhimurium</i> . <i>Journal of the American Society for Mass Spectrometry</i> , 2019, 30, 1001-1012.	1.2	8
12	<i>Bartonella quintana</i> type IV secretion effector BepE induced selective autophagy by conjugation with K63 polyubiquitin chain. <i>Cellular Microbiology</i> , 2019, 21, e12984.	1.1	14
13	Proteomic approaches beyond expression profiling and PTM analysis. <i>Analytical and Bioanalytical Chemistry</i> , 2018, 410, 4051-4060.	1.9	9
14	Regulation of the small GTPase Rab1 function by a bacterial glucosyltransferase. <i>Cell Discovery</i> , 2018, 4, 53.	3.1	28
15	<i>Shigella flexneri</i> Regulator SlyA Controls Bacterial Acid Resistance by Directly Activating the Glutamate Decarboxylation System. <i>Frontiers in Microbiology</i> , 2018, 9, 2071.	1.5	4
16	Proteomic Delineation of the ArcA Regulon in <i>Salmonella Typhimurium</i> During Anaerobiosis. <i>Molecular and Cellular Proteomics</i> , 2018, 17, 1937-1947.	2.5	17
17	A Proteomic View of <i>Salmonella Typhimurium</i> in Response to Phosphate Limitation. <i>Proteomes</i> , 2018, 6, 19.	1.7	9
18	<i>Salmonella</i> proteomics under oxidative stress reveals coordinated regulation of antioxidant defense with iron metabolism and bacterial virulence. <i>Journal of Proteomics</i> , 2017, 157, 52-58.	1.2	36

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19	Quantitative proteomic analysis of host epithelial cells infected by <i>Salmonella enterica</i> serovar Typhimurium. <i>Proteomics</i> , 2017, 17, 1700092.	1.3	14
20	Identification of a Novel Salmonella Type III Effector by Quantitative Secretome Profiling. <i>Molecular and Cellular Proteomics</i> , 2017, 16, 2219-2228.	2.5	31
21	Temporal Regulation of a Salmonella Typhimurium Virulence Factor by the Transcriptional Regulator YdcR. <i>Molecular and Cellular Proteomics</i> , 2017, 16, 1683-1693.	2.5	9
22	Proteomic Analyses of Intracellular <i>Salmonella enterica</i> Serovar Typhimurium Reveal Extensive Bacterial Adaptations to Infected Host Epithelial Cells. <i>Infection and Immunity</i> , 2015, 83, 2897-2906.	1.0	66
23	Elucidating Ionic Liquid Environments That Affect the Morphology of TiO ₂ Nanocrystals: A DFT Study. <i>Journal of Physical Chemistry C</i> , 2014, 118, 23320-23327.	1.5	18