

## List of Publications by Year in descending order

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#	Article	IF	CITATIONS
1	Legionella pneumophila modulates host energy metabolism by ADP-ribosylation of ADP/ATP translocases. ELife, 2022, 11, .	6.0	27
2	<i>Legionella pneumophila</i> temporally regulates the activity of ADP/ATP translocases by reversible ADPâ€ribosylation. , 2022, 1, 51-65.		7
3	<i>Coxiella burnetii</i> inhibits host immunity by a protein phosphatase adapted from glycolysis. Proceedings of the National Academy of Sciences of the United States of America, 2022, 119, .	7.1	158
4	Bartonella type IV secretion effector BepC induces stress fiber formation through activation of GEF-H1. PLoS Pathogens, 2021, 17, e1009065.	4.7	2
5	Shigella evades pyroptosis by arginine ADP-riboxanation of caspase-11. Nature, 2021, 599, 290-295.	27.8	93
6	<i>Legionella pneumophila</i> regulates the activity of <scp>UBE</scp> 2N by deamidaseâ€mediated deubiquitination. EMBO Journal, 2020, 39, e102806.	7.8	38
7	Legionella effector MavC targets the Ube2N~Ub conjugate for noncanonical ubiquitination. Nature Communications, 2020, 11, 2365.	12.8	21
8	Arginine GlcNAcylation of Rab small GTPases by the pathogen Salmonella Typhimurium. Communications Biology, 2020, 3, 287.	4.4	27
9	Structural insights into the mechanism and inhibition of transglutaminase-induced ubiquitination by the Legionella effector MavC. Nature Communications, 2020, 11, 1774.	12.8	15
10	Salmonella Proteomic Profiling during Infection Distinguishes the Intracellular Environment of Host Cells. MSystems, 2019, 4, .	3.8	20
11	Proteomic Analysis of FNR-Regulated Anaerobiosis in <i>Salmonella</i> Typhimurium. Journal of the American Society for Mass Spectrometry, 2019, 30, 1001-1012.	2.8	8
12	Bartonella quintana type IV secretion effector BepE â€induced selective autophagy by conjugation with K63 polyubiquitin chain. Cellular Microbiology, 2019, 21, e12984.	2.1	14
13	Proteomic approaches beyond expression profiling and PTM analysis. Analytical and Bioanalytical Chemistry, 2018, 410, 4051-4060.	3.7	9
14	Regulation of the small GTPase Rab1 function by a bacterial glucosyltransferase. Cell Discovery, 2018, 4, 53.	6.7	28
15	Shigellaflexneri Regulator SlyA Controls Bacterial Acid Resistance by Directly Activating the Glutamate Decarboxylation System. Frontiers in Microbiology, 2018, 9, 2071.	3.5	4
16	Proteomic Delineation of the ArcA Regulon in Salmonella Typhimurium During Anaerobiosis. Molecular and Cellular Proteomics, 2018, 17, 1937-1947.	3.8	17
17	A Proteomic View of Salmonella Typhimurium in Response to Phosphate Limitation. Proteomes, 2018, 6, 19.	3.5	9
18	Salmonella proteomics under oxidative stress reveals coordinated regulation of antioxidant defense with iron metabolism and bacterial virulence. Journal of Proteomics, 2017, 157, 52-58	2.4	36

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19	Quantitative proteomic analysis of host epithelial cells infected by <i>Salmonella enterica</i> serovar Typhimurium. Proteomics, 2017, 17, 1700092.	2.2	14
20	Identification of a Novel Salmonella Type III Effector by Quantitative Secretome Profiling. Molecular and Cellular Proteomics, 2017, 16, 2219-2228.	3.8	31
21	Temporal Regulation of a Salmonella Typhimurium Virulence Factor by the Transcriptional Regulator YdcR. Molecular and Cellular Proteomics, 2017, 16, 1683-1693.	3.8	9
22	Proteomic Analyses of Intracellular Salmonella enterica Serovar Typhimurium Reveal Extensive Bacterial Adaptations to Infected Host Epithelial Cells. Infection and Immunity, 2015, 83, 2897-2906.	2.2	66
23	Elucidating Ionic Liquid Environments That Affect the Morphology of TiO <sub>2</sub> Nanocrystals: A DFT+ <i>D</i> Study. Journal of Physical Chemistry C, 2014, 118, 23320-23327.	3.1	18