Hyunjin Yoon

List of Publications by Year in descending order

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Ηγιινιν Υρον

#	Article	IF	CITATIONS
1	Retroreflection-based sandwich type affinity sensing of isothermal gene amplification products for foodborne pathogen detection. Analyst, The, 2022, 147, 450-460.	3.5	2
2	\ddot{I} fS-Mediated Stress Response Induced by Outer Membrane Perturbation Dampens Virulence in Salmonella enterica serovar Typhimurium. Frontiers in Microbiology, 2021, 12, 750940.	3.5	5
3	Understanding the multifaceted roles of the phosphoenolpyruvate: Phosphotransferase system in regulation of Salmonella virulence using a mutant defective in ptsI and crr expression. Microbiological Research, 2019, 223-225, 63-71.	5.3	13
4	New virulence factor CSK29544_02616 as LpxA binding partner in Cronobacter sakazakii. Scientific Reports, 2018, 8, 835.	3.3	5
5	Secretion of Salmonella Pathogenicity Island 1-Encoded Type III Secretion System Effectors by Outer Membrane Vesicles in Salmonella enterica Serovar Typhimurium. Frontiers in Microbiology, 2018, 9, 2810.	3.5	24
6	Enzyme IIANtr Regulates Salmonella Invasion Via 1,2-Propanediol And Propionate Catabolism. Scientific Reports, 2017, 7, 44827.	3.3	22
7	Endolysin LysSA97 is synergistic with carvacrol in controlling Staphylococcus aureus in foods. International Journal of Food Microbiology, 2017, 244, 19-26.	4.7	59
8	Fine-tuning of amino sugar homeostasis by EllANtr in Salmonella Typhimurium. Scientific Reports, 2016, 6, 33055.	3.3	26
9	Roles of Outer Membrane Vesicles (OMVs) in Bacterial Virulence. Journal of Bacteriology and Virology, 2015, 45, 1.	0.1	8
10	Temporal regulation of Salmonella pathogenicity Island 1 (SPI-1) hilA by Hfq in Salmonella enterica serovar typhimurium. Journal of the Korean Society for Applied Biological Chemistry, 2015, 58, 169-172.	0.9	1
11	Transcriptional response of selected genes of Salmonella enterica serovar Typhimurium biofilm cells during inactivation by superheated steam. International Journal of Food Microbiology, 2015, 192, 117-123.	4.7	11
12	<i>hfq</i> Plays Important Roles in Virulence and Stress Adaptation in Cronobacter sakazakii ATCC 29544. Infection and Immunity, 2015, 83, 2089-2098.	2.2	44
13	Identification and Characterization of Outer Membrane Vesicle-Associated Proteins in Salmonella enterica Serovar Typhimurium. Infection and Immunity, 2014, 82, 4001-4010.	2.2	70
14	A comparison of saturated steam and superheated steam for inactivation of Escherichia coli O157:H7, Salmonella Typhimurium, and Listeria monocytogenes biofilms on polyvinyl chloride and stainless steel. Food Control, 2014, 40, 344-350.	5.5	52
15	Characterization and genomic analysis of two Staphylococcus aureus bacteriophages isolated from poultry/livestock farms. Journal of General Virology, 2013, 94, 2569-2576.	2.9	13
16	Analysis of HilC/D-dependent invF promoter expression under different culture conditions. Microbial Pathogenesis, 2012, 52, 359-366.	2.9	10
17	Technologies and Approaches to Elucidate and Model the Virulence Program of Salmonella. Frontiers in Microbiology, 2011, 2, 121.	3.5	18
18	Experimental annotation of post-translational features and translated coding regions in the pathogen Salmonella Typhimurium. BMC Genomics, 2011, 12, 433.	2.8	29

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19	Systems analysis of multiple regulator perturbations allows discovery of virulence factors in Salmonella. BMC Systems Biology, 2011, 5, 100.	3.0	30
20	Discovery of Salmonella Virulence Factors Translocated via Outer Membrane Vesicles to Murine Macrophages. Infection and Immunity, 2011, 79, 2182-2192.	2.2	77
21	Quantitative PCR-Based Competitive Index for High-Throughput Screening of Salmonella Virulence Factors. Infection and Immunity, 2011, 79, 360-368.	2.2	10
22	ppGpp-mediated stationary phase induction of the genes encoded by horizontally acquired pathogenicity islands and cob/pdu locus in Salmonella enterica serovar Typhimurium. Journal of Microbiology, 2010, 48, 89-95.	2.8	11
23	<i>Salmonella</i> pathogenicity island 2 expression negatively controlled by EIIA ^{Ntr} –SsrB interaction is required for <i>Salmonella</i> virulence. Proceedings of the National Academy of Sciences of the United States of America, 2010, 107, 20506-20511.	7.1	48
24	Proteomic Investigation of the Time Course Responses of RAW 264.7 Macrophages to Infection with <i>Salmonella enterica</i> . Infection and Immunity, 2009, 77, 3227-3233.	2.2	54
25	Coordinated Regulation of Virulence during Systemic Infection of Salmonella enterica Serovar Typhimurium. PLoS Pathogens, 2009, 5, e1000306.	4.7	143
26	A Method for Investigating Proteinâ^'Protein Interactions Related to <i>Salmonella</i> Typhimurium Pathogenesis. Journal of Proteome Research, 2009, 8, 1504-1514.	3.7	22
27	Bottlenecks and Hubs in Inferred Networks Are Important for Virulence in <i>Salmonella typhimurium</i> . Journal of Computational Biology, 2009, 16, 169-180.	1.6	73
28	Global Systems-Level Analysis of Hfq and SmpB Deletion Mutants in Salmonella: Implications for Virulence and Global Protein Translation. PLoS ONE, 2009, 4, e4809.	2.5	109
29	Proteomics Analysis of the Causative Agent of Typhoid Fever. Journal of Proteome Research, 2008, 7, 546-557.	3.7	54
30	Use of high-throughput mass spectrometry to elucidate host–pathogen interactions inSalmonella. Future Microbiology, 2008, 3, 625-634.	2.0	8
31	Effects of chaperones on mRNA stability and gene expression in Escherichia coli. Journal of Microbiology and Biotechnology, 2008, 18, 228-33.	2.1	11
32	Mlc regulation of Salmonella pathogenicity island I gene expression via hilE repression. Nucleic Acids Research, 2007, 35, 1822-1832.	14.5	44
33	ppGpp-dependent Stationary Phase Induction of Genes on Salmonella Pathogenicity Island 1. Journal of Biological Chemistry, 2004, 279, 34183-34190.	3.4	129
34	Proteome analysis ofSalmonella entericaserovar Typhimuriumfismutant. FEMS Microbiology Letters, 2003, 226, 391-396.	1.8	21