Xenia P Kostoulias

List of Publications by Year in descending order

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XENIA P KOSTOULIAS

#	Article	IF	CITATIONS
1	Genomic and phenotypic analyses of diverse non-clinical Acinetobacter baumannii strains reveals strain-specific virulence and resistance capacity. Microbial Genomics, 2022, 8, .	2.0	7
2	Mpeg1 is not essential for antibacterial or antiviral immunity, but is implicated in antigen presentation. Immunology and Cell Biology, 2022, 100, 529-546.	2.3	4
3	Phage-antibiotic combination is a superior treatment against Acinetobacter baumannii in a preclinical study. EBioMedicine, 2022, 80, 104045.	6.1	40
4	Bacteriophage-resistant Acinetobacter baumannii are resensitized to antimicrobials. Nature Microbiology, 2021, 6, 157-161.	13.3	159
5	The Resistance to Host Antimicrobial Peptides in Infections Caused by Daptomycin-Resistant Staphylococcus aureus. Antibiotics, 2021, 10, 96.	3.7	6
6	Antibiotic-chemoattractants enhance neutrophil clearance of Staphylococcus aureus. Nature Communications, 2021, 12, 6157.	12.8	18
7	Targeting NLRP3 and Staphylococcal pore-forming toxin receptors in human-induced pluripotent stem cell-derived macrophages. Journal of Leukocyte Biology, 2020, 108, 967-981.	3.3	19
8	The Mechanisms of Disease Caused by Acinetobacter baumannii. Frontiers in Microbiology, 2019, 10, 1601.	3.5	220
9	Antibiotic resistance and host immune evasion in <i>Staphylococcus aureus</i> mediated by a metabolic adaptation. Proceedings of the National Academy of Sciences of the United States of America, 2019, 116, 3722-3727.	7.1	69
10	Unstable chromosome rearrangements in <i>Staphylococcus aureus</i> cause phenotype switching associated with persistent infections. Proceedings of the National Academy of Sciences of the United States of America, 2019, 116, 20135-20140.	7.1	69
11	Evolution of Daptomycin Resistance in Coagulase-Negative Staphylococci Involves Mutations of the Essential Two-Component Regulator WalKR. Antimicrobial Agents and Chemotherapy, 2019, 63, .	3.2	22
12	Evolution of carbapenem resistance in Acinetobacter baumannii during a prolonged infection. Microbial Genomics, 2018, 4, .	2.0	49
13	Global Gene Expression Profile of Acinetobacter baumannii During Bacteremia. Journal of Infectious Diseases, 2017, 215, S52-S57.	4.0	38
14	Vancomycin-intermediate Staphylococcus aureus isolates are attenuated for virulence when compared with susceptible progenitors. Clinical Microbiology and Infection, 2017, 23, 767-773.	6.0	30
15	Synthesis of novel 1,2,5-oxadiazoles and evaluation of action against Acinetobacter baumannii. Bioorganic and Medicinal Chemistry, 2017, 25, 6267-6272.	3.0	16
16	<i>Acinetobacter baumannii</i> phenylacetic acid metabolism influences infection outcome through a direct effect on neutrophil chemotaxis. Proceedings of the National Academy of Sciences of the United States of America, 2016, 113, 9599-9604.	7.1	109
17	Vancomycin susceptibility in methicillin-resistant Staphylococcus aureus is mediated by YycHI activation of the WalRK essential two-component regulatory system. Scientific Reports, 2016, 6, 30823.	3.3	48
18	Impact of a Cross-Kingdom Signaling Molecule of Candida albicans on Acinetobacter baumannii Physiology. Antimicrobial Agents and Chemotherapy, 2016, 60, 161-167.	3.2	40

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19	A Clobal Virulence Regulator in Acinetobacter baumannii and Its Control of the Phenylacetic Acid Catabolic Pathway. Journal of Infectious Diseases, 2014, 210, 46-55.	4.0	139
20	The RpoB H481Y Rifampicin Resistance Mutation and an Active Stringent Response Reduce Virulence and Increase Resistance to Innate Immune Responses in Staphylococcus aureus. Journal of Infectious Diseases, 2013, 207, 929-939.	4.0	94
21	Serine/Threonine Phosphatase Stp1 Contributes to Reduced Susceptibility to Vancomycin and Virulence in Staphylococcus aureus. Journal of Infectious Diseases, 2012, 205, 1677-1687.	4.0	98
22	Expression Patterns and Roles of Periostin During Kidney and Ureter Development. Journal of Urology, 2011, 186, 1537-1544.	0.4	22
23	Spatial gene expression in the T-stage mouse metanephros. Gene Expression Patterns, 2006, 6, 807-825.	0.8	37
24	Temporal and spatial transcriptional programs in murine kidney development. Physiological Genomics, 2005, 23, 159-171.	2.3	64
25	Variant esp gene as a marker of a distinct genetic lineage of vancomycinresistant Enterococcus faecium spreading in hospitals. Lancet, The, 2001, 357, 853-855.	13.7	291