

# Robert Stones

## List of Publications by Year in descending order

Source: <https://exaly.com/author-pdf/12029240/publications.pdf>

Version: 2024-02-01

16  
papers

814  
citations

759233

12  
h-index

996975

15  
g-index

16  
all docs

16  
docs citations

16  
times ranked

1020  
citing authors

#	ARTICLE	IF	CITATIONS
1	DNA sequences and predicted protein structures of <i>prot6E</i> and <i>sefA</i> genes for <i>Salmonella</i> ser. Enteritidis detection. <i>Food Control</i> , 2019, 96, 271-280.	5.5	3
2	Targeting Tropomyosin Receptor Kinase in Cutaneous CYLD Defective Tumors With Pegcantratinib. <i>JAMA Dermatology</i> , 2018, 154, 913.	4.1	14
3	Changing of the Genomic Pattern of <i>Salmonella</i> Enteritidis Strains Isolated in Brazil Over a 48 year-period revealed by Whole Genome SNP Analyses. <i>Scientific Reports</i> , 2018, 8, 10478.	3.3	18
4	Baseline Practices for the Application of Genomic Data Supporting Regulatory Food Safety. <i>Journal of AOAC INTERNATIONAL</i> , 2017, 100, 721-731.	1.5	25
5	Whole Genome and Core Genome Multilocus Sequence Typing and Single Nucleotide Polymorphism Analyses of <i>Listeria monocytogenes</i> Isolates Associated with an Outbreak Linked to Cheese, United States, 2013. <i>Applied and Environmental Microbiology</i> , 2017, 83, .	3.1	93
6	Complete Genome Sequences of Two <i>Salmonella enterica</i> subsp. <i>enterica</i> Serovar Enteritidis Strains Isolated from Egg Products in the United States. <i>Genome Announcements</i> , 2017, 5, .	0.8	6
7	The Rise of Genomics and the Promise of Whole Genome Sequencing for Understanding Microbial Foodborne Pathogens. , 2017, , 333-351.		5
8	<i>Listeria monocytogenes</i> in Stone Fruits Linked to a Multistate Outbreak: Enumeration of Cells and Whole-Genome Sequencing. <i>Applied and Environmental Microbiology</i> , 2016, 82, 7030-7040.	3.1	83
9	Comparison of whole genome sequences from human and non-human <i>Escherichia coli</i> O26 strains. <i>Frontiers in Cellular and Infection Microbiology</i> , 2015, 5, 21.	3.9	15
10	Genetic Diversity of <i>Salmonella</i> Pathogenicity Islands SPI-5 and SPI-6 in <i>Salmonella</i> Newport. <i>Foodborne Pathogens and Disease</i> , 2014, 11, 798-807.	1.8	13
11	On the Evolutionary History, Population Genetics and Diversity among Isolates of <i>Salmonella</i> Enteritidis PFGE Pattern JEGX01.0004. <i>PLoS ONE</i> , 2013, 8, e55254.	2.5	146
12	Phylogenetics and Differentiation of <i>Salmonella</i> Newport Lineages by Whole Genome Sequencing. <i>PLoS ONE</i> , 2013, 8, e55687.	2.5	63
13	Phylogenomic Analysis Identifies Gene Gains That Define <i>Salmonella enterica</i> Subspecies I. <i>PLoS ONE</i> , 2013, 8, e76821.	2.5	8
14	Draft Genome Sequences of 21 <i>Salmonella enterica</i> Serovar Enteritidis Strains. <i>Journal of Bacteriology</i> , 2012, 194, 5994-5995.	2.2	27
15	High resolution clustering of <i>Salmonella enterica</i> serovar Montevideo strains using a next-generation sequencing approach. <i>BMC Genomics</i> , 2012, 13, 32.	2.8	140
16	Identification of a Salmonellosis Outbreak by Means of Molecular Sequencing. <i>New England Journal of Medicine</i> , 2011, 364, 981-982.	27.0	155