

# Prerak T Desai

## List of Publications by Year in descending order

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Version: 2024-02-01

23  
papers

711  
citations

759233

12  
h-index

642732

23  
g-index

23  
all docs

23  
docs citations

23  
times ranked

1279  
citing authors

#	ARTICLE	IF	CITATIONS
1	Genome Sequence of Pigmented Siderophore-Producing Strain <i>Serratia marcescens</i> SM6. <i>Microbiology Resource Announcements</i> , 2019, 8, .	0.6	13
2	Whole Cell Cross-Linking to Discover Host-Microbe Protein Cognate Receptor/Ligand Pairs. <i>Frontiers in Microbiology</i> , 2018, 9, 1585.	3.5	9
3	Draft Genome Sequence of <i>Salmonella enterica</i> subsp. <i>enterica</i> Serovar Orion Strain CRJJGF_00093 (Phylum <i>Gammaproteobacteria</i> ). <i>Genome Announcements</i> , 2016, 4, .	0.8	6
4	Draft Genome Sequence of <i>Salmonella enterica</i> subsp. <i>diarizonae</i> Serovar 61:k:1,5,(7) Strain CRJJGF_00165 (Phylum <i>Gammaproteobacteria</i> ). <i>Genome Announcements</i> , 2016, 4, .	0.8	4
5	Reply to Yue. <i>Clinical Infectious Diseases</i> , 2016, 62, 1326-1327.	5.8	1
6	Draft Genome Sequence of <i>Salmonella enterica</i> subsp. <i>enterica</i> Serovar Bardo Strain CRJJGF_00099 (Phylum <i>Gammaproteobacteria</i> ). <i>Genome Announcements</i> , 2016, 4, .	0.8	7
7	Draft Genome Sequence of <i>Salmonella enterica</i> subsp. <i>enterica</i> Serovar Putten Strain CRJJGF_00159 (Phylum <i>Gammaproteobacteria</i> ). <i>Genome Announcements</i> , 2016, 4, .	0.8	4
8	Draft Genome Sequence of <i>Salmonella enterica</i> subsp. <i>enterica</i> Serovar Blockley Strain CRJJGF_00147 (Phylum <i>Gammaproteobacteria</i> ). <i>Genome Announcements</i> , 2016, 4, .	0.8	4
9	Draft Genome Sequence of <i>Salmonella enterica</i> subsp. <i>enterica</i> Serovar Kiambu Strain CRJJGF_00061 (Phylum <i>Gammaproteobacteria</i> ). <i>Genome Announcements</i> , 2016, 4, .	0.8	4
10	Draft Genome Sequence of <i>Salmonella enterica</i> subsp. <i>enterica</i> Serovar Lille Strain CRJJGF_000101 (Phylum <i>Gammaproteobacteria</i> ). <i>Genome Announcements</i> , 2016, 4, .	0.8	4
11	Draft Genome Sequence of <i>Salmonella enterica</i> subsp. <i>enterica</i> Serovar Widemarsh Strain CRJJGF_00058 (Phylum <i>Gammaproteobacteria</i> ). <i>Genome Announcements</i> , 2016, 4, .	0.8	4
12	Persistent Infections by Nontyphoidal <i>Salmonella</i> in Humans: Epidemiology and Genetics. <i>Clinical Infectious Diseases</i> , 2016, 62, 879-886.	5.8	98
13	Development of the Intestinal RNA Virus Community of Healthy Broiler Chickens. <i>PLoS ONE</i> , 2016, 11, e0150094.	2.5	9
14	Solid tumors provide niche-specific conditions that lead to preferential growth of <i>Salmonella</i> . <i>Oncotarget</i> , 2016, 7, 35169-35180.	1.8	35
15	Genetic Mechanisms Underlying the Pathogenicity of Cold-Stressed <i>Salmonella enterica</i> Serovar Typhimurium in Cultured Intestinal Epithelial Cells. <i>Applied and Environmental Microbiology</i> , 2014, 80, 6943-6953.	3.1	27
16	Integrative Analysis of Salmonellosis in Israel Reveals Association of <i>Salmonella enterica</i> Serovar 9,12:l,v with Extraintestinal Infections, Dissemination of Endemic <i>S. enterica</i> Serovar Typhimurium DT104 Biotypes, and Severe Underreporting of Outbreaks. <i>Journal of Clinical Microbiology</i> , 2014, 52, 2078-2088.	3.9	14
17	Evolutionary Genomics of <i>Salmonella enterica</i> Subspecies. <i>MBio</i> , 2013, 4, .	4.1	106
18	Preadaptation to Cold Stress in <i>Salmonella enterica</i> Serovar Typhimurium Increases Survival during Subsequent Acid Stress Exposure. <i>Applied and Environmental Microbiology</i> , 2013, 79, 7281-7289.	3.1	59

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19	Evolutionary Genomics of Salmonella enterica Subspecies. MBio, 2013, 4, .	4.1	38
20	Virulence Gene Profiling and Pathogenicity Characterization of Non-Typhoidal Salmonella Accounted for Invasive Disease in Humans. PLoS ONE, 2013, 8, e58449.	2.5	143
21	Consumption of Lysozyme-Rich Milk Can Alter Microbial Fecal Populations. Applied and Environmental Microbiology, 2012, 78, 6153-6160.	3.1	87
22	Virulence of 32 Salmonella Strains in Mice. PLoS ONE, 2012, 7, e36043.	2.5	19
23	Solid-Phase Capture of Pathogenic Bacteria by Using Gangliosides and Detection with Real-Time PCR. Applied and Environmental Microbiology, 2008, 74, 2254-2258.	3.1	16