

Pallavi Singh

List of Publications by Year in descending order

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Version: 2024-02-01

21
papers

701
citations

759233

12
h-index

713466

21
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22
all docs

22
docs citations

22
times ranked

1266
citing authors

#	ARTICLE	IF	CITATIONS
1	Intestinal microbial communities associated with acute enteric infections and disease recovery. <i>Microbiome</i> , 2015, 3, 45.	11.1	151
2	Influence of penicillin on microbial diversity of the cecal microbiota in broiler chickens. <i>Poultry Science</i> , 2013, 92, 272-276.	3.4	102
3	Pilus distribution among lineages of group b streptococcus: an evolutionary and clinical perspective. <i>BMC Microbiology</i> , 2014, 14, 159.	3.3	58
4	Factors Associated with Shiga Toxin-Producing <i>Escherichia coli</i> Shedding by Dairy and Beef Cattle. <i>Applied and Environmental Microbiology</i> , 2016, 82, 5049-5056.	3.1	55
5	Antimicrobial Susceptibility Profiles of Human <i>Campylobacter jejuni</i> Isolates and Association with Phylogenetic Lineages. <i>Frontiers in Microbiology</i> , 2016, 7, 589.	3.5	48
6	Impact of age and sex on the composition and abundance of the intestinal microbiota in individuals with and without enteric infections. <i>Annals of Epidemiology</i> , 2016, 26, 380-385.	1.9	47
7	Characterization of enteropathogenic and Shiga toxin-producing <i>Escherichia coli</i> in cattle and deer in a shared agroecosystem. <i>Frontiers in Cellular and Infection Microbiology</i> , 2015, 5, 29.	3.9	43
8	Clade 8 and Clade 6 Strains of <i>Escherichia coli</i> O157:H7 from Cattle in Argentina have Hypervirulent-Like Phenotypes. <i>PLoS ONE</i> , 2015, 10, e0127710.	2.5	39
9	Intestinal Microbial Community Dynamics of White-Tailed Deer (<i>Odocoileus virginianus</i>) in an Agroecosystem. <i>Microbial Ecology</i> , 2017, 74, 496-506.	2.8	37
10	Microbial Ecology: Where are we now?. <i>Postdoc Journal</i> , 2016, 4, 3-17.	0.4	23
11	A Nonhemolytic Group B <i>Streptococcus</i> Strain Exhibits Hypervirulence. <i>Journal of Infectious Diseases</i> , 2018, 217, 983-987.	4.0	18
12	Multilocus sequence typing of <i>Salmonella</i> strains by high-throughput sequencing of selectively amplified target genes. <i>Journal of Microbiological Methods</i> , 2012, 88, 127-133.	1.6	17
13	Massively parallel sequencing of enriched target amplicons for high-resolution genotyping of <i>Salmonella</i> serovars. <i>Molecular and Cellular Probes</i> , 2013, 27, 80-85.	2.1	11
14	Comparative analysis of <i>Campylobacter</i> populations within individual market-age broilers using fla gene typing method. <i>Poultry Science</i> , 2013, 92, 2135-2144.	3.4	9
15	Draft Genome Sequence of an Invasive <i>Streptococcus agalactiae</i> Isolate Lacking Pigmentation. <i>Genome Announcements</i> , 2016, 4, .	0.8	9
16	Whole-Genome Shotgun Sequencing of a Colonizing Multilocus Sequence Type 17 <i>Streptococcus agalactiae</i> Strain. <i>Journal of Bacteriology</i> , 2012, 194, 6005-6005.	2.2	8
17	Characterizing the Cattle Gut Microbiome in Farms with a High and Low Prevalence of Shiga Toxin Producing <i>Escherichia coli</i> . <i>Microorganisms</i> , 2021, 9, 1737.	3.6	8
18	Variability in the Occupancy of <i>Escherichia coli</i> O157 Integration Sites by Shiga Toxin-Encoding Prophages. <i>Toxins</i> , 2021, 13, 433.	3.4	7

#	ARTICLE	IF	CITATIONS
19	High prevalence of clade 8 Escherichia coli O157:H7 isolated from retail meat and butcher shop environment. <i>Infection, Genetics and Evolution</i> , 2016, 45, 1-5.	2.3	5
20	Draft Genome Sequence of a Diarrheagenic <i>Morganella morganii</i> Isolate. <i>Genome Announcements</i> , 2015, 3, .	0.8	3
21	Target-Enrichment Through Amplification of Hairpin-Ligated Universal Targets for Next-Generation Sequencing Analysis. <i>Methods in Molecular Biology</i> , 2011, 733, 267-278.	0.9	2