## Siddhartha Thakur

List of Publications by Year in descending order

Source: https://exaly.com/author-pdf/8994529/publications.pdf

Version: 2024-02-01

101 papers

2,434 citations

147801 31 h-index 233421 45 g-index

108 all docs

108 docs citations

108 times ranked

2856 citing authors

#	Article	IF	CITATIONS
1	Global Perspective of the Vitamin D Status of African-Caribbean Populations: A Systematic Review and Meta-analysis. European Journal of Clinical Nutrition, 2022, 76, 516-526.	2.9	9
2	Characterization of Salmonella enterica Contamination in Pork and Poultry Meat from São Paulo/Brazil: Serotypes, Genotypes and Antimicrobial Resistance Profiles. Pathogens, 2022, 11, 358.	2.8	8
3	Genomics of human and chicken Salmonella isolates in Senegal: Broilers as a source of antimicrobial resistance and potentially invasive nontyphoidal salmonellosis infections. PLoS ONE, 2022, 17, e0266025.	2.5	7
4	Genomic Screening of Antimicrobial Resistance Markers in UK and US <i>Campylobacter</i> Isolates Highlights Stability of Resistance over an 18-Year Period. Antimicrobial Agents and Chemotherapy, 2022, 66, e0168721.	3.2	9
5	Multidrug resistance and virulence genes carried by mobile genomic elements in Salmonella enterica isolated from live food animals, processed, and retail meat in North Carolina, 2018–2019. International Journal of Food Microbiology, 2022, 378, 109821.	4.7	11
6	Antimicrobial resistance and interspecies gene transfer in Campylobacter coli and Campylobacter jejuni isolated from food animals, poultry processing, and retail meat in North Carolina, 2018–2019. PLoS ONE, 2021, 16, e0246571.	2.5	34
7	Genetic relatedness of multidrug resistant Escherichia coli isolated from humans, chickens and poultry environments. Antimicrobial Resistance and Infection Control, 2021, 10, 58.	4.1	61
8	Colonization with multidrug-resistant Enterobacteriaceae among infants: an observational study in southern Sri Lanka. Antimicrobial Resistance and Infection Control, 2021, 10, 72.	4.1	6
9	The effect of vegetation barriers at reducing the transmission of Salmonella and Escherichia coli from animal operations to fresh produce. International Journal of Food Microbiology, 2021, 347, 109196.	4.7	14
10	Ultra-accurate microbial amplicon sequencing with synthetic long reads. Microbiome, 2021, 9, 130.	11.1	53
11	Impact of the COVID-19 pandemic on the surveillance, prevention and control of antimicrobial resistance: a global survey. Journal of Antimicrobial Chemotherapy, 2021, 76, 3045-3058.	3.0	88
12	Genomic Characterization of a Nalidixic Acid-Resistant Salmonella Enteritidis Strain Causing Persistent Infections in Broiler Chickens. Frontiers in Veterinary Science, 2021, 8, 725737.	2.2	4
13	Does Irrigation with Treated and Untreated Wastewater Increase Antimicrobial Resistance in Soil and Water: A Systematic Review. International Journal of Environmental Research and Public Health, 2021, 18, 11046.	2.6	12
14	Identification of CTX-M Type ESBL E. coli from Sheep and Their Abattoir Environment Using Whole-Genome Sequencing. Pathogens, 2021, 10, 1480.	2.8	5
15	Transmission of antimicrobial resistant non-O157 Escherichia coli at the interface of animal-fresh produce in sustainable farming environments. International Journal of Food Microbiology, 2020, 319, 108472.	4.7	15
16	The role of <i>Enterococcus faecalis</i> during co-infection with avian pathogenic <i>Escherichia coli</i> in avian colibacillosis. Avian Pathology, 2020, 49, 589-599.	2.0	12
17	Molecular Epidemiology of Infectious Zoonotic and Livestock Diseases. Microbiology Spectrum, 2020, 8, .	3.0	12
18	Extended-spectrum ß-lactamase-producing Escherichia coli among humans, chickens and poultry environments in Abuja, Nigeria. One Health Outlook, 2020, 2, 8.	3.4	32

#	Article	IF	CITATIONS
19	Antimicrobial resistance and virulence factors profile of Salmonella spp. and Escherichia coli isolated from different environments exposed to anthropogenic activity. Journal of Global Antimicrobial Resistance, 2020, 22, 578-583.	2.2	17
20	International lineages of Salmonella enterica serovars isolated from chicken farms, Wakiso District, Uganda. PLoS ONE, 2020, 15, e0220484.	2.5	13
21	Class 1 integron-borne cassettes harboring blaCARB-2 gene in multidrug-resistant and virulent Salmonella Typhimurium ST19 strains recovered from clinical human stool samples, United States. PLoS ONE, 2020, 15, e0240978.	2.5	12
22	Title is missing!. , 2020, 15, e0220484.		0
23	Title is missing!. , 2020, 15, e0220484.		0
24	Title is missing!. , 2020, 15, e0220484.		0
25	Title is missing!. , 2020, 15, e0220484.		0
26	Title is missing!. , 2020, 15, e0220484.		0
27	Title is missing!. , 2020, 15, e0220484.		0
28	Title is missing!. , 2020, 15, e0240978.		0
29	Title is missing!. , 2020, 15, e0240978.		0
30	Title is missing!. , 2020, 15, e0240978.		0
31	Title is missing!. , 2020, 15, e0240978.		0
32	Genomic Features of High-Priority Salmonella enterica Serovars Circulating in the Food Production Chain, Brazil, 2000–2016. Scientific Reports, 2019, 9, 11058.	3.3	61
33	First Genome Sequence of Brucella abortus Biovar 3 Strain BAU21/S4023, Isolated from a Dairy Cow in Bangladesh. Microbiology Resource Announcements, 2019, 8, .	0.6	5
34	The Mandate for a Global "One Health―Approach to Antimicrobial Resistance Surveillance. American Journal of Tropical Medicine and Hygiene, 2019, 100, 227-228.	1.4	51
35	Antibiotic Usage in Poultry Production and Antimicrobial-Resistant Salmonella in Poultry. , 2019, , 47-66.		11
36	Prevalence and risk factors for multi-drug resistant Escherichia coli among poultry workers in the Federal Capital Territory, Abuja, Nigeria. PLoS ONE, 2019, 14, e0225379.	2.5	44

#	Article	IF	CITATIONS
37	Phenotypic and Genotypic Characterization of <i>Escherichia coli</i> and <i>Salmonella enterica</i> from Dairy Cattle Farms in the Wakiso District, Uganda: A Cross-Sectional Study. Foodborne Pathogens and Disease, 2019, 16, 54-59.	1.8	14
38	First report of mcr-1-harboring Salmonella enterica serovar Schwarzengrund isolated from poultry meat in Brazil. Diagnostic Microbiology and Infectious Disease, 2019, 93, 376-379.	1.8	41
39	Multidrug- and colistin-resistant Salmonella enterica 4,[5],12:i:- sequence type 34 carrying the mcr-3.1 gene on the IncHI2 plasmid recovered from a human. Journal of Medical Microbiology, 2019, 68, 986-990.	1.8	20
40	Endemic fluoroquinolone-resistant Salmonella enterica serovar Kentucky ST198 in northern India. Microbial Genomics, 2019, 5, .	2.0	21
41	Title is missing!. , 2019, 14, e0225379.		0
42	Title is missing!. , 2019, 14, e0225379.		0
43	Title is missing!. , 2019, 14, e0225379.		0
44	Title is missing!. , 2019, 14, e0225379.		0
45	Molecular Tools To Study Preharvest Food Safety Challenges. Microbiology Spectrum, 2018, 6, .	3.0	4
46	Understanding the Complexities of Food Safety Using a "One Health―Approach. Microbiology Spectrum, 2018, 6, .	3.0	11
47	Environmental Dissemination of Multidrug Methicillin-Resistant <i>Staphylococcus sciuri</i> Application of Manure from Commercial Swine Production Systems. Foodborne Pathogens and Disease, 2018, 15, 210-217.	1.8	20
48	Whole genome sequencing analysis of multiple Salmonella serovars provides insights into phylogenetic relatedness, antimicrobial resistance, and virulence markers across humans, food animals and agriculture environmental sources. BMC Genomics, 2018, 19, 801.	2.8	100
49	The role of citizen science in addressing grand challenges in food and agriculture research. Proceedings of the Royal Society B: Biological Sciences, 2018, 285, 20181977.	2.6	97
50	Multilaboratory Survey To Evaluate Salmonella Prevalence in Diarrheic and Nondiarrheic Dogs and Cats in the United States between 2012 and 2014. Journal of Clinical Microbiology, 2017, 55, 1350-1368.	3.9	58
51	Horizontal Dissemination of Antimicrobial Resistance Determinants in Multiple Salmonella Serotypes following Isolation from the Commercial Swine Operation Environment after Manure Application. Applied and Environmental Microbiology, 2017, 83, .	3.1	45
52	Assessing the Impact of Manure Application in Commercial Swine Farms on the Transmission of Antimicrobial Resistant Salmonella in the Environment. PLoS ONE, 2016, 11, e0164621.	2.5	46
53	Biofilm Formation by Environmental Isolates of $\langle i \rangle$ Salmonella $\langle i \rangle$ and Their Sensitivity to Natural Antimicrobials. Foodborne Pathogens and Disease, 2016, 13, 509-516.	1.8	21
54	Comparative Phenotypic and Genotypic Characterization of Temporally Related Nontyphoidal <i>Salmonella</i> Solated from Human Clinical Cases, Pigs, and the Environment in North Carolina. Foodborne Pathogens and Disease, 2014, 11, 156-164.	1.8	10

#	Article	IF	Citations
55	Dissemination of plasmid-encoded AmpC $\hat{l}^2$ -lactamases in antimicrobial resistant Salmonella serotypes originating from humans, pigs and the swine environment. Veterinary Microbiology, 2014, 173, 76-83.	1.9	17
56	Yersinia enterocoliticaof Porcine Origin: Carriage of Virulence Genes and Genotypic Diversity. Foodborne Pathogens and Disease, 2013, 10, 80-86.	1.8	14
57	Farm and environmental distribution of Campylobacter and Salmonella in broiler flocks. Research in Veterinary Science, 2013, 94, 33-42.	1.9	39
58	Longitudinal Study of Distributions of Similar Antimicrobial-Resistant Salmonella Serovars in Pigs and Their Environment in Two Distinct Swine Production Systems. Applied and Environmental Microbiology, 2013, 79, 5167-5178.	3.1	57
59	Methicillin-Resistant Staphylococcus aureus in Pigs and Farm Workers on Conventional and Antibiotic-Free Swine Farms in the USA. PLoS ONE, 2013, 8, e63704.	2.5	124
60	Longitudinal Study of the Persistence of Antimicrobial-Resistant Campylobacter Strains in Distinct Swine Production Systems on Farms, at Slaughter, and in the Environment. Applied and Environmental Microbiology, 2012, 78, 2698-2705.	3.1	42
61	Antimicrobial Resistance, Toxinotype, and Genotypic Profiling of Clostridium difficile Isolates of Swine Origin. Journal of Clinical Microbiology, 2012, 50, 2366-2372.	3.9	46
62	Molecular Characterization of Salmonella enterica Serotype Enteritidis Isolates from Humans by Antimicrobial Resistance, Virulence Genes, and Pulsed-Field Gel Electrophoresis. Foodborne Pathogens and Disease, 2012, 9, 232-238.	1.8	54
63	Phylogenetic Analysis Reveals Common Antimicrobial Resistant Campylobacter coli Population in Antimicrobial-Free (ABF) and Commercial Swine Systems. PLoS ONE, 2012, 7, e44662.	2.5	10
64	Prevalence and Antimicrobial Resistance Profile of <i>Campylobacter </i> Spp. Isolated from Conventional and Antimicrobial-Free Swine Production Systems from Different U.S. Regions. Foodborne Pathogens and Disease, 2011, 8, 367-374.	1.8	31
65	Detection of Clostridium difficile and Salmonella in Feral Swine Population in North Carolina. Journal of Wildlife Diseases, 2011, 47, 774-776.	0.8	33
66	Prevalence and Distribution of <i>Salmonella</i> in Organic and Conventional Broiler Poultry Farms. Foodborne Pathogens and Disease, 2010, 7, 1363-1371.	1.8	102
67	Prevalence of antimicrobial resistance and association with toxin genes in Clostridium difficile in commercial swine. American Journal of Veterinary Research, 2010, 71, 1189-1194.	0.6	33
68	Antimicrobial Resistance, Virulence, and Genotypic Profile Comparison of <i>Campylobacter jejuni</i> and <i>Campylobacter coli</i> lsolated from Humans and Retail Meats. Foodborne Pathogens and Disease, 2010, 7, 835-844.	1.8	56
69	Longitudinal Study of <i>Salmonella</i> Dispersion and the Role of Environmental Contamination in Commercial Swine Production Systems. Applied and Environmental Microbiology, 2009, 75, 1478-1486.	3.1	48
70	Occurrence of <i>spvA</i> Virulence Gene and Clinical Significance for Multidrug-Resistant <i>Salmonella</i> Strains. Journal of Clinical Microbiology, 2009, 47, 777-780.	3.9	31
71	Molecular Epidemiologic Investigation of <i>Campylobacter coli</i> in Swine Production Systems, Using Multilocus Sequence Typing. Applied and Environmental Microbiology, 2008, 74, 342-342.	3.1	7
72	Occurrence of multidrug resistant Salmonella in antimicrobial-free (ABF) swine production systems. Veterinary Microbiology, 2007, 125, 362-367.	1.9	26

#	Article	lF	Citations
73	Comparison of Prevalence, Antimicrobial Resistance, and Occurrence of Multidrug-Resistant Salmonella in Antimicrobial-Free and Conventional Pig Production. Journal of Food Protection, 2006, 69, 743-748.	1.7	58
74	Molecular Epidemiologic Investigation of Campylobacter coli in Swine Production Systems, Using Multilocus Sequence Typing. Applied and Environmental Microbiology, 2006, 72, 5666-5669.	3.1	38
75	Prevalence and Antimicrobial Resistance of Campylobacter in Antimicrobial-Free and Conventional Pig Production Systems. Journal of Food Protection, 2005, 68, 2402-2410.	1.7	73
76	Multidrug-Resistant Salmonella enterica Serovar Muenchen from Pigs and Humans and Potential Interserovar Transfer of Antimicrobial Resistance. Antimicrobial Agents and Chemotherapy, 2005, 49, 503-511.	3.2	84
77	Campylobacter coli in Swine Production: Antimicrobial Resistance Mechanisms and Molecular Epidemiology. Journal of Clinical Microbiology, 2005, 43, 5705-5714.	3.9	57
78	Campylobacter coli: prevalence and antimicrobial resistance in antimicrobial-free (ABF) swine production systems. Journal of Antimicrobial Chemotherapy, 2005, 56, 765-768.	3.0	50
79	Characterization of Antimicrobial-Resistant Phenotypes and Genotypes among Salmonella enterica Recovered from Pigs on Farms, from Transport Trucks, and from Pigs after Slaughter. Journal of Food Protection, 2004, 67, 698-705.	1.7	44
80	Trends in antimicrobial resistance, phage types and integrons among Salmonella serotypes from pigs, 1997-2000. Journal of Antimicrobial Chemotherapy, 2004, 53, 997-1003.	3.0	84
81	Preharvest Farming Practices Impacting Fresh Produce Safety. , 0, , 19-46.		2
82	Nuts and Grains: Microbiology and Preharvest Contamination Risks., 0,, 105-121.		12
83	Risks Associated with Fish and Seafood. , 0, , 123-142.		4
84	Water for Agriculture: the Convergence of Sustainability and Safety., 0,, 143-157.		7
85	Importance of Soil Amendments: Survival of Bacterial Pathogens in Manure and Compost Used as Organic Fertilizers., 0,, 159-175.		9
86	Prevalence and Antimicrobial Resistance of Salmonella, E. coli, and Campylobacter in Pigs from Swine Producing States in the United States. , $0$ , , .		3
87	Understanding the Complexities of Food Safety Using a "One Health―Approach. , 0, , 401-411.		3
88	Reducing Foodborne Pathogen Persistence and Transmission in Animal Production Environments: Challenges and Opportunities., 0,, 177-203.		1
89	Molecular Tools To Study Preharvest Food Safety Challenges. , 0, , 361-382.		0
90	Current Status of the Preharvest Application of Pro- and Prebiotics to Farm Animals to Enhance the Microbial Safety of Animal Products., 0,, 349-360.		1

#	Article	IF	CITATIONS
91	Preharvest Food Safety Under the Influence of a Changing Climate. , 0, , 261-271.		O
92	Introduction to Preharvest Food Safety., 0,, 1-17.		O
93	Toxoplasma gondii as a Parasite in Food: Analysis and Control. , 0, , 227-247.		1
94	Regulatory Issues Associated with Preharvest Food Safety: European Union Perspective., 0,, 325-347.		0
95	Preharvest Food Safety Challenges in Beef and Dairy Production. , 0, , 47-68.		O
96	Preharvest Food Safety in Broiler Chicken Production. , 0, , 69-86.		2
97	Mathematical Modeling Tools to Study Preharvest Food Safety. , 0, , 383-400.		O
98	Egg Safety in the Realm of Preharvest Food Safety. , 0, , 87-104.		0
99	Local Food Systems Food Safety Concerns. , 0, , 249-260.		2
100	Phage Therapy Approaches to Reducing Pathogen Persistence and Transmission in Animal Production Environments: Opportunities and Challenges., 0,, 289-308.		1
101	Potential for Meta-Analysis in the Realm of Preharvest Food Safety. , 0, , 273-287.		O