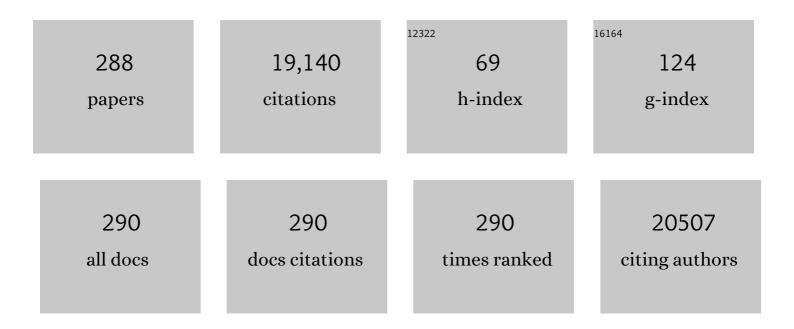
Trinad Chakraborty

List of Publications by Year in descending order

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#	Article	IF	CITATIONS
1	Prediction of antimicrobial resistance based on whole-genome sequencing and machine learning. Bioinformatics, 2022, 38, 325-334.	1.8	54
2	Ecology of <i>Listeria monocytogenes</i> and <i>Listeria</i> species in India: the occurrence, resistance to biocides, genomic landscape and biocontrol. Environmental Microbiology, 2022, 24, 2759-2780.	1.8	4
3	New Delhi Metallo-β-Lactamase-Producing Enterobacterales Bacteria. Emerging Infectious Diseases, 2022, 28, 265-265.	2.0	1
4	Development and validation of BLOOMY prediction scores for 14-day and 6-month mortality in hospitalised adults with bloodstream infections: a multicentre, prospective, cohort study. Lancet Infectious Diseases, The, 2022, 22, 731-741.	4.6	15
5	Fosfomycin as a salvage therapy for treating urinary tract infections due to multidrug-resistant Escherichia coli. European Journal of Clinical Microbiology and Infectious Diseases, 2022, 41, 689-690.	1.3	1
6	Multi-label classification for multi-drug resistance prediction of Escherichia coli. Computational and Structural Biotechnology Journal, 2022, 20, 1264-1270.	1.9	8
7	Porphyromonas gingivalis W83 Membrane Components Induce Distinct Profiles of Metabolic Genes in Oral Squamous Carcinoma Cells. International Journal of Molecular Sciences, 2022, 23, 3442.	1.8	3
8	Characterisation of new anti-O157 bacteriophages of bovine origin representing three genera. Archives of Microbiology, 2022, 204, 231.	1.0	1
9	Comparison of Urine Flow Cytometry on the UF-1000i System and Urine Culture of Urine Samples from Urological Patients. Urologia Internationalis, 2022, 106, 858-868.	0.6	1
10	Recombinant <i>Porphyromonas gingivalis</i> W83 FimA alters immune response and metabolic gene expression in oral squamous carcinoma cells. Clinical and Experimental Dental Research, 2022, 8, 976-987.	0.8	3
11	Dual Role of Hydrogen Peroxide as an Oxidant in Pneumococcal Pneumonia. Antioxidants and Redox Signaling, 2021, 34, 962-978.	2.5	13
12	Uropathogenic <i>Escherichia coli</i> Virulence Factor α-Hemolysin Reduces Histone Acetylation to Inhibit Expression of Proinflammatory Cytokine Genes. Journal of Infectious Diseases, 2021, 223, 1040-1051.	1.9	4
13	Streptococcus pneumoniae and Its Virulence Factors H2O2 and Pneumolysin Are Potent Mediators of the Acute Chest Syndrome in Sickle Cell Disease. Toxins, 2021, 13, 157.	1.5	10
14	Cross-Border Emergence of Escherichia coli Producing the Carbapenemase NDM-5 in Switzerland and Germany. Journal of Clinical Microbiology, 2021, 59, .	1.8	35
15	Molecular (real-time reverse transcription polymerase chain reaction) diagnosis of SARS-CoV-2 infections: complexity and challenges. Journal of Laboratory Medicine, 2021, 45, 135-142.	1.1	7
16	The Genus Listeria. , 2021, , 411-442.		8
17	<i>Porphyromonas gingivalis</i> induced upâ€regulation of PDâ€L1 in colon carcinoma cells. Molecular Oral Microbiology, 2021, 36, 172-181.	1.3	13
18	Changing epidemiology of vancomycin-resistant Enterococcus faecium: Results of a genome-based study at a regional neurological acute hospital with intensive care and early rehabilitation treatment. Infection Prevention in Practice, 2021, 3, 100138.	0.6	3

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19	Recent Emergence of Aztreonam-Avibactam Resistance in NDM and OXA-48 Carbapenemase-Producing Escherichia coli in Germany. Antimicrobial Agents and Chemotherapy, 2021, 65, e0109021.	1.4	14
20	Carbapenem-Resistant Citrobacter spp. as an Emerging Concern in the Hospital-Setting: Results From a Genome-Based Regional Surveillance Study. Frontiers in Cellular and Infection Microbiology, 2021, 11, 744431.	1.8	23
21	Enabling One Health solutions through genomics. Indian Journal of Medical Research, 2021, 153, 273.	0.4	2
22	Dichotomous Role of Tumor Necrosis Factor in Pulmonary Barrier Function and Alveolar Fluid Clearance. Frontiers in Physiology, 2021, 12, 793251.	1.3	16
23	Genetic Diversity of <i>Listeria monocytogenes</i> Isolates from Invasive Listeriosis in China. Foodborne Pathogens and Disease, 2020, 17, 215-227.	0.8	12
24	Cross-border emergence of clonal lineages of ST38 Escherichia coli producing the OXA-48-like carbapenemase OXA-244 in Germany and Switzerland. International Journal of Antimicrobial Agents, 2020, 56, 106157.	1.1	18
25	Bacterial Colonization within the First Six Weeks of Life and Pulmonary Outcome in Preterm Infants <1000 g. Journal of Clinical Medicine, 2020, 9, 2240.	1.0	15
26	Detection of blaCTX-M-27-encoding Escherichia coli ST206 in Nigerian poultry stocks. Journal of Antimicrobial Chemotherapy, 2020, 75, 3070-3072.	1.3	7
27	ResFinder 4.0 for predictions of phenotypes from genotypes. Journal of Antimicrobial Chemotherapy, 2020, 75, 3491-3500.	1.3	1,523
28	ASA3P: An automatic and scalable pipeline for the assembly, annotation and higher-level analysis of closely related bacterial isolates. PLoS Computational Biology, 2020, 16, e1007134.	1.5	59
29	Phosphocholine Antagonizes Listeriolysin O-Induced Host Cell Responses of Listeria monocytogenes. Journal of Infectious Diseases, 2020, 222, 1505-1516.	1.9	8
30	Ongoing dissemination of OXA-244 carbapenemase-producing Escherichia coli in Switzerland and their detection. Diagnostic Microbiology and Infectious Disease, 2020, 97, 115059.	0.8	12
31	Impact of Bacterial Toxins in the Lungs. Toxins, 2020, 12, 223.	1.5	21
32	Near-ubiquitous presence of a vancomycin-resistant Enterococcus faecium ST117/CT71/vanB –clone in the Rhine-Main metropolitan area of Germany. Antimicrobial Resistance and Infection Control, 2019, 8, 128.	1.5	33
33	Complete Genome Sequence of the Plant Growth-Promoting Bacterium <i>Hartmannibacter diazotrophicus</i> Strain E19 ^T . International Journal of Genomics, 2019, 2019, 1-12.	0.8	17
34	Whole-Genome Sequences of Clinical Enterobacter bugandensis Isolates from Germany. Microbiology Resource Announcements, 2019, 8, .	0.3	8
35	A hybrid sub-lineage of Listeria monocytogenes comprising hypervirulent isolates. Nature Communications, 2019, 10, 4283.	5.8	76
36	Bacterial Outer Membrane Vesicles (OMVs)-Based Dual Vaccine for Influenza A H1N1 Virus and MERS-CoV. Vaccines, 2019, 7, 46.	2.1	38

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37	Draft Genome Sequence of Listeria monocytogenes CIIMS-NV-3, a Strain Isolated from Vaginal Discharge of a Woman from Central India. Microbiology Resource Announcements, 2019, 8, .	0.3	2
38	Multidrug-Resistant and Clinically Relevant Gram-Negative Bacteria Are Present in German Surface Waters. Frontiers in Microbiology, 2019, 10, 2779.	1.5	38
39	Complete genome sequence of C130_2, a novel myovirus infecting pathogenic Escherichia coli and Shigella strains. Archives of Virology, 2019, 164, 321-324.	0.9	5
40	Genome-based analyses indicate that Serratia marcescens subsp. marcescens and Serratia marcescens subsp. sakuensis do not merit separation to subspecies status. International Journal of Systematic and Evolutionary Microbiology, 2019, 69, 3924-3926.	0.8	6
41	In Reply. Deutsches Ärzteblatt International, 2019, 116, 115.	0.6	0
42	An Improved Medium for Colistin Susceptibility Testing. Journal of Clinical Microbiology, 2018, 56, .	1.8	21
43	Draft Genome Sequence of <i>Listeria monocytogenes</i> Strain CIIMS-PH-1, a Serovar 4b Isolate from Infant Septicemia. Genome Announcements, 2018, 6, .	0.8	2
44	Enterobacter bugandensis: a novel enterobacterial species associated with severe clinical infection. Scientific Reports, 2018, 8, 5392.	1.6	61
45	Missing elimination via membrane vesicle shedding contributes to the diminished calcium sensitivity of listeriolysin O. Scientific Reports, 2018, 8, 15846.	1.6	9
46	Genome Sequence of Listeria monocytogenes 2542, a Serotype 4b Strain from a Cheese-Related Outbreak in Portugal. Genome Announcements, 2018, 6, .	0.8	2
47	Listeriolysin O Causes ENaC Dysfunction in Human Airway Epithelial Cells. Toxins, 2018, 10, 79.	1.5	5
48	Treatment Options for Carbapenem-Resistant Gram-Negative Infections. Deutsches Ärzteblatt International, 2018, 115, 345-352.	0.6	68
49	<i>Escherichia coli</i> Sequence Type 410 Is Causing New International High-Risk Clones. MSphere, 2018, 3, .	1.3	183
50	Identification and characterization of new broad host-range rV5-like coliphages C203 and P206 directed against enterobacteria. Infection, Genetics and Evolution, 2018, 64, 254-261.	1.0	14
51	Hsp70 Suppresses Mitochondrial Reactive Oxygen Species and Preserves Pulmonary Microvascular Barrier Integrity Following Exposure to Bacterial Toxins. Frontiers in Immunology, 2018, 9, 1309.	2.2	33
52	Identification and Characterization of T5-Like Bacteriophages Representing Two Novel Subgroups from Food Products. Frontiers in Microbiology, 2018, 9, 202.	1.5	39
53	Retrospective Analysis of Bacterial Cultures Sampled in German Chicken-Fattening Farms During the Years 2011–2012 Revealed Additional VIM-1 Carbapenemase-Producing Escherichia coli and a Serologically Rough Salmonella enterica Serovar Infantis. Frontiers in Microbiology, 2018, 9, 538.	1.5	14
54	Predictors of the extended-spectrum-beta lactamases producing Enterobacteriaceae neonatal sepsis at a tertiary hospital, Tanzania. International Journal of Medical Microbiology, 2018, 308, 803-811.	1.5	56

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55	Uropathogenic <i>Escherichia coli</i> virulence factor hemolysin A causes programmed cell necrosis by altering mitochondrial dynamics. FASEB Journal, 2018, 32, 4107-4120.	0.2	25
56	Diversity of CTX-M-1-producing E. coli from German food samples and genetic diversity of the bla CTX-M-1 region on Incl1 ST3 plasmids. Veterinary Microbiology, 2018, 221, 98-104.	0.8	54
57	Listeria goaensis sp. nov International Journal of Systematic and Evolutionary Microbiology, 2018, 68, 3285-3291.	0.8	38
58	High resistance to tetracycline and ciprofloxacin in bacteria isolated from poultry farms in Ibadan, Nigeria. Journal of Infection in Developing Countries, 2018, 12, 462-470.	0.5	18
59	Autophagy: A Potential Antibacterial Therapeutic Target. , 2018, , 203-214.		Ο
60	Comparative genome analysis of IncHI2 VIM-1 carbapenemase-encoding plasmids of Escherichia coli and Salmonella enterica isolated from a livestock farm in Germany. Veterinary Microbiology, 2017, 200, 114-117.	0.8	55
61	Genome Analysis of the Carbapenem- and Colistin-Resistant Escherichia coli Isolate NRZ14408 Reveals Horizontal Gene Transfer Pathways towards Panresistance and Enhanced Virulence. Antimicrobial Agents and Chemotherapy, 2017, 61, .	1.4	9
62	Evaluation of a Loop-Mediated Isothermal Amplification-Based Assay for the Rapid Detection of Plasmid-Encoded Colistin Resistance Gene <i>mcr-1</i> in Enterobacteriaceae Isolates. Antimicrobial Agents and Chemotherapy, 2017, 61, .	1.4	18
63	Draft Genome Sequences of Pandrug-Resistant Serratia marcescens Clinical Isolates Harboring <i> bla _{NDM-1} </i> . Genome Announcements, 2017, 5, .	0.8	2
64	Requirement of the RNA-binding protein SmpB during intracellular growth of Listeria monocytogenes. International Journal of Medical Microbiology, 2017, 307, 166-173.	1.5	8
65	Characterization of the Micro-Environment of the Testis that Shapes the Phenotype and Function of Testicular Macrophages. Journal of Immunology, 2017, 198, 4327-4340.	0.4	86
66	Environmental emission of multiresistant <i>Escherichia coli</i> carrying the colistin resistance gene <i>mcr-1</i> from German swine farms. Journal of Antimicrobial Chemotherapy, 2017, 72, dkw585.	1.3	77
67	Synthesis of a biological active β-hairpin peptide by addition of two structural motifs. Bioorganic and Medicinal Chemistry, 2017, 25, 603-608.	1.4	9
68	A Comparative Evaluation of Antimicrobial Effect of Thymus capitatus Ethanolic Extract on the Different Respiratory Tract Infections Isolates. BioNanoScience, 2017, 7, 644-647.	1.5	3
69	Complete Genome Sequence of <i>bla</i> _{CTX-M-27} -Encoding Escherichia coli Strain H105 of Sequence Type 131 Lineage C1/H30R. Genome Announcements, 2017, 5, .	0.8	15
70	Gene expression profiling at birth characterizing the preterm infant with early onset infection. Journal of Molecular Medicine, 2017, 95, 169-180.	1.7	7
71	Distinct Neurotoxicity Profile of Listeriolysin O from Listeria monocytogenes. Toxins, 2017, 9, 34.	1.5	6
72	Listeriolysin O Regulates the Expression of Optineurin, an Autophagy Adaptor That Inhibits the Growth of Listeria monocytogenes. Toxins, 2017, 9, 273.	1.5	16

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73	Epithelial Sodium Channel-α Mediates the Protective Effect of the TNF-Derived TIP Peptide in Pneumolysin-Induced Endothelial Barrier Dysfunction. Frontiers in Immunology, 2017, 8, 842.	2.2	35
74	Insights into a Novel blaKPC-2-Encoding IncP-6 Plasmid Reveal Carbapenem-Resistance Circulation in Several Enterobacteriaceae Species from Wastewater and a Hospital Source in Spain. Frontiers in Microbiology, 2017, 8, 1143.	1.5	50
75	CTX-M-15-Producing E. coli Isolates from Food Products in Germany Are Mainly Associated with an IncF-Type Plasmid and Belong to Two Predominant Clonal E. coli Lineages. Frontiers in Microbiology, 2017, 8, 2318.	1.5	62
76	Listeria monocytogenes Induces a Virulence-Dependent microRNA Signature That Regulates the Immune Response in Galleria mellonella. Frontiers in Microbiology, 2017, 8, 2463.	1.5	17
77	<i>bla</i> _{CTX-M-27} –Encoding <i>Escherichia coli</i> Sequence Type 131 Lineage C1-M27 Clone in Clinical Isolates, Germany. Emerging Infectious Diseases, 2017, 23, 1754-1756.	2.0	48
78	Chromosomal Locations ofmcr-1andblaCTX-M-15in Fluoroquinolone-ResistantEscherichia coliST410. Emerging Infectious Diseases, 2016, 22, 1689-1691.	2.0	70
79	Multiple ESBL-Producing Escherichia coli Sequence Types Carrying Quinolone and Aminoglycoside Resistance Genes Circulating in Companion and Domestic Farm Animals in Mwanza, Tanzania, Harbor Commonly Occurring Plasmids. Frontiers in Microbiology, 2016, 7, 142.	1.5	63
80	Predominance of CTX-M-15 among ESBL Producers from Environment and Fish Gut from the Shores of Lake Victoria in Mwanza, Tanzania. Frontiers in Microbiology, 2016, 7, 1862.	1.5	68
81	TLR9 mediates S. aureus killing inside osteoblasts via induction of oxidative stress. BMC Microbiology, 2016, 16, 230.	1.3	29
82	Detection of translocatable units in a blaCTX-M-15 extended-spectrum β-lactamase-producing ST131 Escherichia coli isolate using a hybrid sequencing approach. International Journal of Antimicrobial Agents, 2016, 47, 245-247.	1.1	9
83	Varibaculum anthropi sp. nov. represented by three genetically different genomovars isolated from clinical material and emended description of the genus Varibaculum. Systematic and Applied Microbiology, 2016, 39, 546-552.	1.2	15
84	ActA of Listeria monocytogenes and Its Manifold Activities as an Important Listerial Virulence Factor. Current Topics in Microbiology and Immunology, 2016, 399, 113-132.	0.7	26
85	A multiplex PCR for detection of Listeria monocytogenes and its lineages. Journal of Microbiological Methods, 2016, 130, 144-147.	0.7	23
86	The Lectin-like Domain of TNF Increases ENaC Open Probability through a Novel Site at the Interface between the Second Transmembrane and C-terminal Domains of the I±-Subunit. Journal of Biological Chemistry, 2016, 291, 23440-23451.	1.6	20
87	Modern diagnostic methods for urinary tract infections. Expert Review of Anti-Infective Therapy, 2016, 14, 1047-1063.	2.0	25
88	Tracking bacterial virulence: global modulators as indicators. Scientific Reports, 2016, 6, 25973.	1.6	12
89	Diverse roles of endoplasmic reticulum stress sensors in bacterial infection. Molecular and Cellular Pediatrics, 2016, 3, 9.	1.0	27
90	Circulation of clonal populations of fluoroquinolone-resistant CTX-M-15-producing Escherichia coli ST410 in humans and animals in Germany. International Journal of Antimicrobial Agents, 2016, 47, 457-465.	1.1	107

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91	Predictors of blaCTX-M-15 in varieties of Escherichia coli genotypes from humans in community settings in Mwanza, Tanzania. BMC Infectious Diseases, 2016, 16, 187.	1.3	69
92	Presence of a widely disseminated <i>Listeria monocytogenes</i> serotype 4b clone in India. Emerging Microbes and Infections, 2016, 5, 1-4.	3.0	17
93	Colistin resistance gene mcr-1 in extended-spectrum β-lactamase-producing and carbapenemase-producing Gram-negative bacteria in Germany. Lancet Infectious Diseases, The, 2016, 16, 282-283.	4.6	271
94	Uropathogenic <i>Escherichia coli</i> Epigenetically Manipulate Host Cell Death Pathways. Journal of Infectious Diseases, 2016, 213, 1198-1207.	1.9	14
95	Kosakonia pseudosacchari sp. nov., an endophyte of Zea mays. Systematic and Applied Microbiology, 2016, 39, 1-7.	1.2	30
96	Fisetin Protects AgainstListeria monocytogenesInfection by Reducing the Production of Listeriolysin O. Journal of Infectious Diseases, 2016, 213, 684-685.	1.9	3
97	The status of the species Enterobacter siamensis Khunthongpan et al. 2014 . Request for an Opinion. International Journal of Systematic and Evolutionary Microbiology, 2016, 66, 524-525.	0.8	8
98	Enterobacter bugandensis sp. nov., isolated from neonatal blood. International Journal of Systematic and Evolutionary Microbiology, 2016, 66, 968-974.	0.8	61
99	Taxonomic reassessment of the genus Elizabethkingia using whole-genome sequencing: Elizabethkingia endophytica K¤pfer et al. 2015 is a later subjective synonym of Elizabethkingia anophelis K¤pfer et al. 2011. International Journal of Systematic and Evolutionary Microbiology, 2016, 66, 4555-4559.	0.8	36
100	A detailed view of the intracellular transcriptome of Listeria monocytogenes in murine macrophages using RNA-seq. Frontiers in Microbiology, 2015, 6, 1199.	1.5	36
101	Biofilm-Forming Abilities of Listeria monocytogenes Serotypes Isolated from Different Sources. PLoS ONE, 2015, 10, e0137046.	1.1	120
102	Pneumococcal Hydrogen Peroxide–Induced Stress Signaling Regulates Inflammatory Genes. Journal of Infectious Diseases, 2015, 211, 306-316.	1.9	31
103	Impact of prophylactic CpG Oligodeoxynucleotide application on implant-associated Staphylococcus aureus bone infection. Bone, 2015, 78, 194-202.	1.4	13
104	Comprehensive molecular, genomic and phenotypic analysis of a major clone of Enterococcus faecalis MLST ST40. BMC Genomics, 2015, 16, 175.	1.2	33
105	Differential Activation of Inflammatory Pathways in Testicular Macrophages Provides a Rationale for Their Subdued Inflammatory Capacity. Journal of Immunology, 2015, 194, 5455-5464.	0.4	64
106	Cell-autonomous responses in <i>Listeria monocytogenes</i> infection. Future Microbiology, 2015, 10, 583-597.	1.0	12
107	Global Transcriptome and Mutagenic Analyses of the Acid Tolerance Response of Salmonella enterica Serovar Typhimurium. Applied and Environmental Microbiology, 2015, 81, 8054-8065.	1.4	60
108	Molecular epidemiology and characterization of an outbreak causing Klebsiella pneumoniae clone carrying chromosomally located bla CTX-M-15 at a German University-Hospital. BMC Microbiology, 2015, 15, 122.	1.3	29

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109	Phylogenomic grouping of Listeria monocytogenes. Canadian Journal of Microbiology, 2015, 61, 637-646.	0.8	12
110	Acute Epididymitis Revisited: Impact of Molecular Diagnostics on Etiology and Contemporary Guideline Recommendations. European Urology, 2015, 68, 428-435.	0.9	97
111	Listeria. , 2015, , 653-690.		Ο
112	PKC-Dependent Phosphorylation of eNOS at T495 Regulates eNOS Coupling and Endothelial Barrier Function in Response to G+ -Toxins. PLoS ONE, 2014, 9, e99823.	1.1	46
113	Detection of Very Long Antisense Transcripts by Whole Transcriptome RNA-Seq Analysis of Listeria monocytogenes by Semiconductor Sequencing Technology. PLoS ONE, 2014, 9, e108639.	1.1	23
114	Uropathogenic Escherichia coli Modulates Innate Immunity To Suppress Th1-Mediated Inflammatory Responses during Infectious Epididymitis. Infection and Immunity, 2014, 82, 1104-1111.	1.0	19
115	Protective effect of Growth Hormone-Releasing Hormone agonist in bacterial toxin-induced pulmonary barrier dysfunction. Frontiers in Physiology, 2014, 5, 259.	1.3	18
116	The TIR Domain Containing Locus of <i>Enterococcus faecalis</i> Is Predominant among Urinary Tract Infection Isolates and Downregulates Host Inflammatory Response. International Journal of Microbiology, 2014, 2014, 1-9.	0.9	14
117	Comparison of Widely Used Listeria monocytogenes Strains EGD, 10403S, and EGD-e Highlights Genomic Differences Underlying Variations in Pathogenicity. MBio, 2014, 5, e00969-14.	1.8	201
118	A Systematic Proteomic Analysis of Listeria monocytogenes House-keeping Protein Secretion Systems. Molecular and Cellular Proteomics, 2014, 13, 3063-3081.	2.5	23
119	A Novel Tumor Necrosis Factor–mediated Mechanism of Direct Epithelial Sodium Channel Activation. American Journal of Respiratory and Critical Care Medicine, 2014, 190, 522-532.	2.5	49
120	The IDO1-induced kynurenines play a major role in the antimicrobial effect of human myeloid cells against <i>Listeria monocytogenes</i> . Innate Immunity, 2014, 20, 401-411.	1.1	33
121	Multiresistant extended-spectrum β-lactamase-producing Enterobacteriaceae from humans, companion animals and horses in central Hesse, Germany. BMC Microbiology, 2014, 14, 187.	1.3	144
122	Crystal structure of listeriolysin O reveals molecular details of oligomerization and pore formation. Nature Communications, 2014, 5, 3690.	5.8	116
123	Complete Nucleotide Sequence of a Citrobacter freundii Plasmid Carrying KPC-2 in a Unique Genetic Environment. Genome Announcements, 2014, 2, .	0.8	12
124	A Î²â€Łactoneâ€Based Antivirulence Drug Ameliorates <i>Staphylococcus aureus</i> Skin Infections in Mice. ChemMedChem, 2014, 9, 710-713.	1.6	35
125	Subgrouping of ESBL-producing Escherichia coli from animal and human sources: An approach to quantify the distribution of ESBL types between different reservoirs. International Journal of Medical Microbiology, 2014, 304, 805-816.	1.5	119
126	Complete Genome Sequence of Phage-Like Plasmid pECOH89, Encoding CTX-M-15. Genome Announcements, 2014, 2, .	0.8	21

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127	Resistance plasmids in ESBL-encoding Escherichia coli isolates from humans, dogs and cats. Berliner Und Munchener Tierarztliche Wochenschrift, 2014, 127, 458-63.	0.7	4
128	A relevant experimental model for human bronchiolitis obliterans syndrome. Journal of Heart and Lung Transplantation, 2013, 32, 1131-1139.	0.3	16
129	Predominance of Klebsiella pneumoniaeST14 carrying CTX-M-15 causing neonatal sepsis in Tanzania. BMC Infectious Diseases, 2013, 13, 466.	1.3	90
130	Occurrence of mutations impairing sigma factor B (SigB) function upon inactivation of Listeria monocytogenes genes encoding surface proteins. Microbiology (United Kingdom), 2013, 159, 1328-1339.	0.7	12
131	Listeria. , 2013, , 219-235.		1
132	Complete Genome Sequence of the Probiotic Enterococcus faecalis Symbioflor 1 Clone DSM 16431. Genome Announcements, 2013, 1, .	0.8	33
133	Mini-Review: Novel Therapeutic Strategies to Blunt Actions of Pneumolysin in the Lungs. Toxins, 2013, 5, 1244-1260.	1.5	26
134	Brain infection and activation of neuronal repair mechanisms by the human pathogen <i>Listeria monocytogenes</i> in the lepidopteran model host <i>Galleria mellonella</i> . Virulence, 2013, 4, 324-332.	1.8	49
135	Arginase 1: An Unexpected Mediator of Pulmonary Capillary Barrier Dysfunction in Models of Acute Lung Injury. Frontiers in Immunology, 2013, 4, 228.	2.2	27
136	Clinical Application of Volatile Organic Compound Analysis for Detecting Infectious Diseases. Clinical Microbiology Reviews, 2013, 26, 462-475.	5.7	251
137	Necrosis Is the Dominant Cell Death Pathway in Uropathogenic Escherichia coli Elicited Epididymo-Orchitis and Is Responsible for Damage of Rat Testis. PLoS ONE, 2013, 8, e52919.	1.1	48
138	microRNA Response to Listeria monocytogenes Infection in Epithelial Cells. International Journal of Molecular Sciences, 2012, 13, 1173-1185.	1.8	57
139	Complete Sequences of Plasmids from the Hemolytic-Uremic Syndrome-Associated Escherichia coli Strain HUSEC41. Journal of Bacteriology, 2012, 194, 532-533.	1.0	26
140	Protein Kinase C-α and Arginase I Mediate Pneumolysin-Induced Pulmonary Endothelial Hyperpermeability. American Journal of Respiratory Cell and Molecular Biology, 2012, 47, 445-453.	1.4	60
141	Agonist of growth hormone-releasing hormone reduces pneumolysin-induced pulmonary permeability edema. Proceedings of the National Academy of Sciences of the United States of America, 2012, 109, 2084-2089.	3.3	50
142	Sertoli-cell-specific knockout of connexin 43 leads to multiple alterations in testicular gene expression in prepubertal mice. DMM Disease Models and Mechanisms, 2012, 5, 895-913.	1.2	37
143	RIG-I detects infection with live <i>Listeria</i> by sensing secreted bacterial nucleic acids. EMBO Journal, 2012, 31, 4153-4164.	3.5	153
144	Comparative genomics and transcriptomics of lineages I, II, and III strains of Listeria monocytogenes. BMC Genomics, 2012, 13, 144.	1.2	88

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145	Lack of PPARÎ ³ in Myeloid Cells Confers Resistance to Listeria monocytogenes Infection. PLoS ONE, 2012, 7, e37349.	1.1	27
146	Protective Immunity to Listeria Monocytogenes Infection Mediated by Recombinant Listeria innocua Harboring the VGC Locus. PLoS ONE, 2012, 7, e35503.	1.1	13
147	Activation of the unfolded protein response by Listeria monocytogenes. Cellular Microbiology, 2012, 14, 949-964.	1.1	107
148	A Novel Approach to the Use of Subgingival Controlled-Release Chlorhexidine Delivery in Chronic Periodontitis: A Randomized Clinical Trial. Journal of Periodontology, 2011, 82, 1131-1139.	1.7	18
149	Outbreak of a novel Enterobacter sp. carrying blaCTX-M-15 in a neonatal unit of a tertiary care hospital in Tanzania. International Journal of Antimicrobial Agents, 2011, 38, 265-9.	1.1	46
150	Role of TLR- / NLR-signaling and the associated cytokines involved in recruitment of neutrophils in murine models of <i>Staphylococcus aureus</i> infection. Virulence, 2011, 2, 316-328.	1.8	22
151	Incidence and genetic variability of Listeria species from three milk processing plants. Food Control, 2011, 22, 1900-1904.	2.8	27
152	Adaptation of Listeria monocytogenes to oxidative and nitrosative stress in IFN-Î ³ -activated macrophages. International Journal of Medical Microbiology, 2011, 301, 547-555.	1.5	10
153	B7-H1 and B7-DC receptors of oral squamous carcinoma cells are upregulated by Porphyromonas gingivalis. Immunobiology, 2011, 216, 1302-1310.	0.8	95
154	Quantification of cell infection caused by Listeria monocytogenes invasion. Journal of Biotechnology, 2011, 154, 76-83.	1.9	4
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