Blake M Hanson

List of Publications by Year in descending order

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236925 175258 3,774 57 25 52 h-index citations g-index papers 62 62 62 5805 all docs docs citations times ranked citing authors

#	Article	IF	CITATIONS
1	Emergence of Clinical <i>Clostridioides difficile</i> Isolates With Decreased Susceptibility to Vancomycin. Clinical Infectious Diseases, 2022, 74, 120-126.	5.8	23
2	Clinical outcomes and bacterial characteristics of carbapenem-resistant Klebsiella pneumoniae complex among patients from different global regions (CRACKLE-2): a prospective, multicentre, cohort study. Lancet Infectious Diseases, The, 2022, 22, 401-412.	9.1	122
3	Efficacy, Safety, Pharmacokinetics, and Microbiome Changes of Ibezapolstat in Adults with <i>Clostridioides difficile</i> Infection: A Phase 2a Multicenter Clinical Trial. Clinical Infectious Diseases, 2022, 75, 1164-1170.	5.8	17
4	Accessory Genomes Drive Independent Spread of Carbapenem-Resistant Klebsiella pneumoniae Clonal Groups 258 and 307 in Houston, TX. MBio, 2022, 13, e0049722.	4.1	17
5	Contemporary Clinical and Molecular Epidemiology of Vancomycin-Resistant Enterococcal Bacteremia: A Prospective Multicenter Cohort Study (VENOUS I). Open Forum Infectious Diseases, 2022, 9, ofab616.	0.9	18
6	Unraveling complex transposable elements surrounding blaGES- 16 in a Pseudomonas aeruginosa ExoU strain. Journal of Global Antimicrobial Resistance, 2022, , .	2.2	O
7	Prevalence and Characterization of the Cefazolin Inoculum Effect in North American Methicillin-Susceptible Staphylococcus aureus Isolates. Journal of Clinical Microbiology, 2022, 60, e0249521.	3.9	7
8	IS <i>>26</i> -mediated amplification of <i>bla</i> OXA-1 and <i>bla</i> CTX-M-15 with concurrent outer membrane porin disruption associated with <i>de novo</i> carbapenem resistance in a recurrent bacteraemia cohort. Journal of Antimicrobial Chemotherapy, 2021, 76, 385-395.	3.0	29
9	Selective digestive decontamination with oral colistin plus gentamicin for persistent bacteraemia caused by non-carbapenemase-producing carbapenem-resistant Klebsiella pneumoniae in a neutropenic patient. JAC-Antimicrobial Resistance, 2021, 3, dlab079.	2.1	2
10	Development and Characterization of High-Throughput Caenorhabditis elegans – Enterococcus faecium Infection Model. Frontiers in Cellular and Infection Microbiology, 2021, 11, 667327.	3.9	5
11	Genomic analysis of carbapenem-resistant Pseudomonas aeruginosa ST143 clone showing susceptibility to broad-spectrum cephalosporins. Journal of Global Antimicrobial Resistance, 2021, 26, 177-179.	2.2	1
12	<i>Candida auris</i> Invasive Infections during a COVID-19 Case Surge. Antimicrobial Agents and Chemotherapy, 2021, 65, e0114621.	3.2	42
13	Impact of Bicarbonate-Î ² -Lactam Exposures on Methicillin-Resistant Staphylococcus aureus (MRSA) Gene Expression in Bicarbonate-Î ² -Lactam-Responsive vs. Non-Responsive Strains. Genes, 2021, 12, 1650.	2.4	7
14	Characterization of the Type I Restriction Modification System Broadly Conserved among Group A Streptococci. MSphere, 2021, 6, e0079921.	2.9	14
15	Commentary on: Optimizing Breast Pocket Irrigation: The Breast Implant–Associated Anaplastic Large Cell Lymphoma (BIA-ALCL) Era. Aesthetic Surgery Journal, 2020, 40, 626-629.	1.6	4
16	Dynamics of <i>bla</i> _{KPC-2} Dissemination from Non-CG258 <i>Klebsiella pneumoniae</i> to Other <i>Enterobacterales</i> via IncN Plasmids in an Area of High Endemicity. Antimicrobial Agents and Chemotherapy, 2020, 64, .	3.2	27
17	Molecular and clinical epidemiology of carbapenem-resistant Enterobacterales in the USA (CRACKLE-2): a prospective cohort study. Lancet Infectious Diseases, The, 2020, 20, 731-741.	9.1	174
18	Simultaneous Infection with <i>Enterobacteriaceae</i> and Pseudomonas aeruginosa Harboring Multiple Carbapenemases in a Returning Traveler Colonized with Candida auris. Antimicrobial Agents and Chemotherapy, 2020, 64, .	3.2	23

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19	Tools for Analysis of the Microbiome. Digestive Diseases and Sciences, 2020, 65, 674-685.	2.3	70
20	Comparison of methodological approaches to human gut microbiota changes in response to metabolic and bariatric surgery: A systematic review. Obesity Reviews, 2020, 21, e13025.	6.5	26
21	Long-Term Compassionate Use of Cefiderocol To Treat Chronic Osteomyelitis Caused by Extensively Drug-Resistant Pseudomonas aeruginosa and Extended-Spectrum-β-Lactamase-Producing Klebsiella pneumoniae in a Pediatric Patient. Antimicrobial Agents and Chemotherapy, 2020, 64, .	3.2	44
22	Insights into the Microbiome of Breast Implants and Periprosthetic Tissue in Breast Implant-Associated Anaplastic Large Cell Lymphoma. Scientific Reports, 2019, 9, 10393.	3.3	76
23	Extensively Drug-Resistant Pseudomonas aeruginosa ST309 Harboring Tandem Guiana Extended Spectrum Î ² -Lactamase Enzymes: A Newly Emerging Threat in the United States. Open Forum Infectious Diseases, 2019, 6, ofz273.	0.9	36
24	Evaluation of 16S rRNA gene sequencing for species and strain-level microbiome analysis. Nature Communications, 2019, 10, 5029.	12.8	1,007
25	Analysis of Sinonasal Microbiota in Exacerbations of Chronic Rhinosinusitis Subgroups. OTO Open, 2019, 3, 2473974X1987510.	1.4	10
26	Longitudinal multi-omics of host–microbe dynamics in prediabetes. Nature, 2019, 569, 663-671.	27.8	391
27	622. The Accessory Genome in Enterococcal Bacteremia: Results from the Vancomycin-Resistant Enterococcal Bacteremia Outcomes Study (VENOUS). Open Forum Infectious Diseases, 2019, 6, S289-S289.	0.9	0
28	626. Mobile Genetic Element Dynamics of Co-Circulating Klebsiella pneumoniae Sequence Types Carrying blaKPC in Houston, Texas. Open Forum Infectious Diseases, 2019, 6, S290-S291.	0.9	0
29	New statistical method identifies cytokines that distinguish stool microbiomes. Scientific Reports, 2019, 9, 20082.	3.3	5
30	Unexpected relationships between frequency of antimicrobial resistance, disease phenotype and emm type in group A Streptococcus. Microbial Genomics, 2019, 5, .	2.0	18
31	Extensive Gene Amplification as a Mechanism for Piperacillin-Tazobactam Resistance in Escherichia coli. MBio, 2018, 9, .	4.1	54
32	Integrative Personal Omics Profiles during Periods of Weight Gain and Loss. Cell Systems, 2018, 6, 157-170.e8.	6.2	183
33	An Analysis of the Epidemic of Klebsiella pneumoniae Carbapenemase-Producing K. pneumoniae: Convergence of Two Evolutionary Mechanisms Creates the "Perfect Storm― Journal of Infectious Diseases, 2018, 217, 82-92.	4.0	70
34	Phylogenomic Classification and the Evolution of Clonal Complex 5 Methicillin-Resistant Staphylococcus aureus in the Western Hemisphere. Frontiers in Microbiology, 2018, 9, 1901.	3.5	84
35	Prevalence and molecular characterization of Staphylococcus aureus in commercially available meat over a one-year period in Iowa, USA. Food Microbiology, 2017, 65, 122-129.	4.2	57
36	Rapid replacement by non-vaccine pneumococcal serotypes may mitigate the impact of the pneumococcal conjugate vaccine on nasopharyngeal bacterial ecology. Scientific Reports, 2017, 7, 8127.	3.3	49

3

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37	Assessing the potential for raw meat to influence human colonization with Staphylococcus aureus. Scientific Reports, 2017, 7, 10848.	3.3	14
38	A Novel Methicillin-ResistantStaphylococcus aureust11469 and a Poultry Endemic Strain t002 (ST5) Are Present in Chicken in Ebonyi State, Nigeria. BioMed Research International, 2017, 2017, 1-5.	1.9	17
39	Community characteristics of the gut microbiomes of competitive cyclists. Microbiome, 2017, 5, 98.	11.1	219
40	Resistance to Ceftazidime-Avibactam Is Due to Transposition of KPC in a Porin-Deficient Strain of Klebsiella pneumoniae with Increased Efflux Activity. Antimicrobial Agents and Chemotherapy, 2017, 61,	3.2	121
41	Abstract 2252: Microbial profiling of the head and neck tumor microenvironment as a biomarker of clinical response to chemoradiation. , 2017, , .		0
42	Characterization of the bacterial and fungal microbiome in indoor dust and outdoor air samples: a pilot study. Environmental Sciences: Processes and Impacts, 2016, 18, 713-724.	3.5	74
43	The importance of the microbiome in epidemiologic research. Annals of Epidemiology, 2016, 26, 301-305.	1.9	35
44	Detection of Airborne Methicillin-Resistant <i>Staphylococcus aureus</i> Inside and Downwind of a Swine Building, and in Animal Feed: Potential Occupational, Animal Health, and Environmental Implications. Journal of Agromedicine, 2016, 21, 149-153.	1.5	28
45	Zoonotic Diseases of Swine: Food-borne and Occupational Aspects of Infection., 2015,, 23-68.		0
46	Molecular characteristics of Staphylococcus aureus isolated from employees, children, and environmental surfaces in Iowa child daycare facilities. American Journal of Infection Control, 2015, 43, 482-488.	2.3	14
47	Swine Farming Is a Risk Factor for Infection With and High Prevalence of Carriage of Multidrug-Resistant Staphylococcus aureus. Clinical Infectious Diseases, 2015, 61, 59-66.	5 . 8	68
48	Detection of livestock-associated methicillin-resistant Staphylococcus aureus among swine workers in Romania. Journal of Infection and Public Health, 2014, 7, 323-332.	4.1	19
49	Isolation and Characterization of Methicillin-Resistant Staphylococcus aureus from Pork Farms and Visiting Veterinary Students. PLoS ONE, 2013, 8, e53738.	2.5	143
50	Antimicrobial resistance and molecular epidemiology of <i>Staphylococcus aureus </i> from Ulaanbaatar, Mongolia. Peerl, 2013, 1, e176.	2.0	7
51	Molecular and epidemiologic predictors of Staphylococcus aureus colonization site in a population with limited nosocomial exposure. American Journal of Infection Control, 2012, 40, 992-996.	2.3	17
52	MRSA in Conventional and Alternative Retail Pork Products. PLoS ONE, 2012, 7, e30092.	2.5	133
53	Methicillin-Susceptible <i>Staphylococcus aureus</i> ST398, New York and New Jersey, USA. Emerging Infectious Diseases, 2012, 18, 700-702.	4.3	55
54	Emerging Swine Zoonoses. Vector-Borne and Zoonotic Diseases, 2011, 11, 1225-1234.	1.5	41

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55	Methicillin-Resistant Staphylococcus aureus in Pork Production Shower Facilities. Applied and Environmental Microbiology, 2011, 77, 696-698.	3.1	17
56	Design Lessons for Older Adult Personal Health Records Software from Older Adults. Lecture Notes in Computer Science, 2011, , 176-185.	1.3	10
57	Reply to Lutgring et al. Clinical Infectious Diseases, 0, , .	5.8	O