List of Publications by Year in descending order

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#	Article	IF	CITATIONS
1	International links between Streptococcus pneumoniae vaccine serotype 4 sequence type (ST) 801 in Northern European shipyard outbreaks of invasive pneumococcal disease. Vaccine, 2022, 40, 1054-1060.	1.7	4
2	Clinical pneumonia in the hospitalised child in Malawi in the post-pneumococcal conjugate vaccine era: a prospective hospital-based observational study. BMJ Open, 2022, 12, e050188.	0.8	2
3	A Streptococcus pneumoniae lineage usually associated with pneumococcal conjugate vaccine (PCV) serotypes is the most common cause of serotype 35B invasive disease in South Africa, following routine use of PCV. Microbial Genomics, 2022, 8, .	1.0	4
4	Comparative Genomics of Disease and Carriage Serotype 1 Pneumococci. Genome Biology and Evolution, 2022, 14, .	1.1	3
5	The role of interspecies recombination in the evolution of antibiotic-resistant pneumococci. ELife, 2021, 10, .	2.8	21
6	Impact and effectiveness of 13-valent pneumococcal conjugate vaccine on population incidence of vaccine and non-vaccine serotype invasive pneumococcal disease in Blantyre, Malawi, 2006–18: prospective observational time-series and case-control studies. The Lancet Global Health, 2021, 9, e989-e998.	2.9	27
7	Streptolysin O concentration and activity is central to in vivo phenotype and disease outcome in Group A Streptococcus infection. Scientific Reports, 2021, 11, 19011.	1.6	1
8	An accessible, efficient and global approach for the large-scale sequencing of bacterial genomes. Genome Biology, 2021, 22, 349.	3.8	20
9	Early Signals of Vaccine-driven Perturbation Seen in Pneumococcal Carriage Population Genomic Data. Clinical Infectious Diseases, 2020, 70, 1294-1303.	2.9	9
10	A mosaic tetracycline resistance gene tet(S/M) detected in an MDR pneumococcal CC230 lineage that underwent capsular switching in South Africa. Journal of Antimicrobial Chemotherapy, 2020, 75, 512-520.	1.3	12
11	Genomic Characteristics of Invasive <i>Streptococcus pneumoniae</i> Serotype 1 in New Caledonia Prior to the Introduction of PCV13. Bioinformatics and Biology Insights, 2020, 14, 117793222096210.	1.0	2
12	Influenza-like illness is associated with high pneumococcal carriage density in Malawian children. Journal of Infection, 2020, 81, 549-556.	1.7	5
13	Bacterial genome-wide association study of hyper-virulent pneumococcal serotype 1 identifies genetic variation associated with neurotropism. Communications Biology, 2020, 3, 559.	2.0	11
14	High residual carriage of vaccine-serotype Streptococcus pneumoniae after introduction of pneumococcal conjugate vaccine in Malawi. Nature Communications, 2020, 11, 2222.	5.8	79
15	Global outbreak research: harmony not hegemony. Lancet Infectious Diseases, The, 2020, 20, 770-772.	4.6	40
16	Increased pathogenicity of pneumococcal serotype 1 is driven by rapid autolysis and release of pneumolysin. Nature Communications, 2020, 11, 1892.	5.8	28
17	Estimating the Economic Impact of Respiratory Syncytial Virus and Other Acute Respiratory Infections Among Infants Receiving Care at a Referral Hospital in Malawi. Journal of the Pediatric Infectious Diseases Society, 2020, 9, 738-745.	0.6	11
18	Visualizing variation within Global Pneumococcal Sequence Clusters (GPSCs) and country population snapshots to contextualize pneumococcal isolates. Microbial Genomics, 2020, 6, .	1.0	25

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19	Evaluation of Pneumococcal Serotyping of Nasopharyngeal-Carriage Isolates by Latex Agglutination, Whole-Genome Sequencing (PneumoCaT), and DNA Microarray in a High-Pneumococcal-Carriage-Prevalence Population in Malawi. Journal of Clinical Microbiology, 2020, 59, .	1.8	8
20	Lower Density and Shorter Duration of Nasopharyngeal Carriage by Pneumococcal Serotype 1 (ST217) May Explain Its Increased Invasiveness over Other Serotypes. MBio, 2020, 11, .	1.8	4
21	Characterization of DNA methylation in Malawian <i>Mycobacterium tuberculosis</i> clinical isolates. PeerJ, 2020, 8, e10432.	0.9	4
22	Impact of Maternal HIV Infection and Placental Malaria on the Transplacental Transfer of Influenza Antibodies in Mother–Infant Pairs in Malawi, 2013–2014. Open Forum Infectious Diseases, 2019, 6, ofz383.	0.4	4
23	Infrequent Transmission of Monovalent Human Rotavirus Vaccine Virus to Household Contacts of Vaccinated Infants in Malawi. Journal of Infectious Diseases, 2019, 219, 1730-1734.	1.9	8
24	Pneumococcal lineages associated with serotype replacement and antibiotic resistance in childhood invasive pneumococcal disease in the post-PCV13 era: an international whole-genome sequencing study. Lancet Infectious Diseases, The, 2019, 19, 759-769.	4.6	165
25	International genomic definition of pneumococcal lineages, to contextualise disease, antibiotic resistance and vaccine impact. EBioMedicine, 2019, 43, 338-346.	2.7	168
26	Genomic analysis of Klebsiella pneumoniae isolates from Malawi reveals acquisition of multiple ESBL determinants across diverse lineages. Journal of Antimicrobial Chemotherapy, 2019, 74, 1223-1232.	1.3	36
27	Pneumococcal pneumonia and carriage in Africa before and after introduction of pneumococcal conjugate vaccines, 2000–2019: protocol for systematic review. BMJ Open, 2019, 9, e030981.	0.8	3
28	Etiology and Risk Factors for Mortality in an Adult Community-acquired Pneumonia Cohort in Malawi. American Journal of Respiratory and Critical Care Medicine, 2019, 200, 359-369.	2.5	51
29	Emerging Resistance to Empiric Antimicrobial Regimens for Pediatric Bloodstream Infections in Malawi (1998–2017). Clinical Infectious Diseases, 2019, 69, 61-68.	2.9	53
30	Putative novel cps loci in a large global collection of pneumococci. Microbial Genomics, 2019, 5, .	1.0	14
31	Impact of Human Immunodeficiency Virus on the Burden and Severity of Influenza Illness in Malawian Adults: A Prospective Cohort and Parallel Case-Control Study. Clinical Infectious Diseases, 2018, 66, 865-876.	2.9	23
32	Global emergence and population dynamics of divergent serotype 3 CC180 pneumococci. PLoS Pathogens, 2018, 14, e1007438.	2.1	74
33	Epidemiology of Severe Acute Respiratory Illness and Risk Factors for Influenza Infection and Clinical Severity among Adults in Malawi, 2011–2013. American Journal of Tropical Medicine and Hygiene, 2018, 99, 772-779.	0.6	11
34	Complexities and dilemmas in community consultation on the design of a research project logo in Malawi. PLoS ONE, 2018, 13, e0205737.	1.1	2
35	Epidemiology of hepatitis B, C and D in Malawi: systematic review. BMC Infectious Diseases, 2018, 18, 516.	1.3	13
36	Global Distribution of Invasive Serotype 35D Streptococcus pneumoniae Isolates following Introduction of 13-Valent Pneumococcal Conjugate Vaccine. Journal of Clinical Microbiology, 2018, 56, .	1.8	12

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37	The global distribution and diversity of protein vaccine candidate antigens in the highly virulent Streptococcus pnuemoniae serotype 1. Vaccine, 2017, 35, 972-980.	1.7	27
38	Epidemiological and Molecular Characterization of an Invasive Group A Streptococcus <i>emm</i> 32.2 Outbreak. Journal of Clinical Microbiology, 2017, 55, 1837-1846.	1.8	12
39	Genomic landscape of extended-spectrum β-lactamase resistance in Escherichia coli from an urban African setting. Journal of Antimicrobial Chemotherapy, 2017, 72, 1602-1609.	1.3	46
40	Comparative Genomic Analysis and In Vivo Modeling of Streptococcus pneumoniae ST3081 and ST618 Isolates Reveal Key Genetic and Phenotypic Differences Contributing to Clonal Replacement of Serotype 1 in The Gambia. Journal of Infectious Diseases, 2017, 216, 1318-1327.	1.9	11
41	Population genetic structure, antibiotic resistance, capsule switching and evolution of invasive pneumococci before conjugate vaccination in Malawi. Vaccine, 2017, 35, 4594-4602.	1.7	27
42	Trends in antimicrobial resistance in bloodstream infection isolates at a large urban hospital in Malawi (1998–2016): a surveillance study. Lancet Infectious Diseases, The, 2017, 17, 1042-1052.	4.6	220
43	Airborne dust and high temperatures are risk factors for invasive bacterial disease. Journal of Allergy and Clinical Immunology, 2017, 139, 977-986.e2.	1.5	59
44	Recombination in Streptococcus pneumoniae Lineages Increase with Carriage Duration and Size of the Polysaccharide Capsule. MBio, 2016, 7, .	1.8	50
45	Respiratory Virus–Associated Severe Acute Respiratory Illness and Viral Clustering in Malawian Children in a Setting With a High Prevalence of HIV Infection, Malaria, and Malnutrition. Journal of Infectious Diseases, 2016, 214, 1700-1711.	1.9	25
46	Distinct Salmonella Enteritidis lineages associated with enterocolitis in high-income settings and invasive disease in low-income settings. Nature Genetics, 2016, 48, 1211-1217.	9.4	191
47	Understanding pneumococcal serotype 1 biology through population genomic analysis. BMC Infectious Diseases, 2016, 16, 649.	1.3	22
48	H3ABioNet, a sustainable pan-African bioinformatics network for human heredity and health in Africa. Genome Research, 2016, 26, 271-277.	2.4	94
49	Phylogenetic Analysis of Invasive Serotype 1 Pneumococcus in South Africa, 1989 to 2013. Journal of Clinical Microbiology, 2016, 54, 1326-1334.	1.8	16
50	Modelling the Contributions of Malaria, HIV, Malnutrition and Rainfall to the Decline in Paediatric Invasive Non-typhoidal Salmonella Disease in Malawi. PLoS Neglected Tropical Diseases, 2015, 9, e0003979.	1.3	48
51	Bioinformatics EducationPerspectives and Challenges out of Africa. Briefings in Bioinformatics, 2015, 16, 355-364.	3.2	61
52	Severe Acute Respiratory Illness Deaths in Sub-Saharan Africa and the Role of Influenza: A Case Series From 8 Countries. Journal of Infectious Diseases, 2015, 212, 853-860.	1.9	43
53	Generic determinants of Streptococcus colonization and infection. Infection, Genetics and Evolution, 2015, 33, 361-370.	1.0	23
54	High multiple carriage and emergence of Streptococcus pneumoniae vaccine serotype variants in Malawian children. BMC Infectious Diseases, 2015, 15, 234.	1.3	56

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55	Mechanisms and impact of genetic recombination in the evolution of Streptococcus pneumoniae. Computational and Structural Biotechnology Journal, 2015, 13, 241-247.	1.9	50
56	Comparative Genomic Analysis of Meningitis- and Bacteremia-Causing Pneumococci Identifies a Common Core Genome. Infection and Immunity, 2015, 83, 4165-4173.	1.0	23
57	Region-specific diversification of the highly virulent serotype 1 Streptococcus pneumoniae. Microbial Genomics, 2015, 1, e000027.	1.0	27
58	A Reduction in Adult Blood Stream Infection and Case Fatality at a Large African Hospital following Antiretroviral Therapy Roll-Out. PLoS ONE, 2014, 9, e92226.	1.1	30
59	Drug Resistance in <i>Salmonella enterica</i> ser. Typhimurium Bloodstream Infection, Malawi. Emerging Infectious Diseases, 2014, 20, 1957-1959.	2.0	56
60	Bacterial Meningitis in Malawian Adults, Adolescents, and Children During the Era of Antiretroviral Scale-up and Haemophilus influenzae Type b Vaccination, 2000–2012. Clinical Infectious Diseases, 2014, 58, e137-e145.	2.9	58
61	Bacterial Meningitis in Malawian Infants <2 Months of Age. Pediatric Infectious Disease Journal, 2014, 33, 560-565.	1.1	32
62	Open source clinical science for emerging infections. Lancet Infectious Diseases, The, 2014, 14, 8-9.	4.6	82
63	Enabling the genomic revolution in Africa. Science, 2014, 344, 1346-1348.	6.0	361
64	Genomic identification of a novel co-trimoxazole resistance genotype and its prevalence amongst Streptococcus pneumoniae in Malawi. Journal of Antimicrobial Chemotherapy, 2014, 69, 368-374.	1.3	31
65	Piliation of Invasive Streptococcus pneumoniae Isolates in the Era before Pneumococcal Conjugate Vaccine Introduction in Malawi. Vaccine Journal, 2013, 20, 1729-1735.	3.2	12
66	Surveillance Programme of IN-patients and Epidemiology (SPINE): Implementation of an Electronic Data Collection Tool within a Large Hospital in Malawi. PLoS Medicine, 2013, 10, e1001400.	3.9	52
67	Defective Pneumococcal-Specific Th1 Responses in HIV-Infected Adults Precedes a Loss of Control of Pneumococcal Colonization. Clinical Infectious Diseases, 2013, 56, 291-299.	2.9	44
68	Bacteraemia in Malawian neonates and young infants 2002–2007: a retrospective audit. BMJ Open, 2012, 2, e000906.	0.8	4
69	Genetic Characterisation of Malawian Pneumococci Prior to the Roll-Out of the PCV13 Vaccine Using a High-Throughput Whole Genome Sequencing Approach. PLoS ONE, 2012, 7, e44250.	1.1	49
70	Invasive <i>Streptococcus pneumoniae</i> in Children, Malawi, 2004–2006. Emerging Infectious Diseases, 2011, 17, 1107-1109.	2.0	15
71	Correlates of HIV-1 Genital Shedding in Tanzanian Women. PLoS ONE, 2011, 6, e17480.	1.1	55
72	Ten Years of Surveillance for Invasive Streptococcus pneumoniae during the Era of Antiretroviral Scale-Up and Cotrimoxazole Prophylaxis in Malawi. PLoS ONE, 2011, 6, e17765.	1.1	64

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73	Invasive <i>Streptococcus pneumoniae</i> in Children, Malawi, 2004–2006. Emerging Infectious Diseases, 2011, 17, 1107-1109.	2.0	21
74	Are Women Who Work in Bars, Guesthouses and Similar Facilities a Suitable Study Population for Vaginal Microbicide Trials in Africa?. PLoS ONE, 2010, 5, e10661.	1.1	17
75	Longâ€Term Impact of Acyclovir Suppressive Therapy on Genital and Plasma HIV RNA in Tanzanian Women: A Randomized Controlled Trial. Journal of Infectious Diseases, 2010, 201, 1285-1297.	1.9	32
76	Association of Schistosomiasis with False-Positive HIV Test Results in an African Adolescent Population. Journal of Clinical Microbiology, 2010, 48, 1570-1577.	1.8	58
77	Suitability of Simple Human Immunodeficiency Virus Rapid Tests in Clinical Trials in Community-Based Clinic Settings. Journal of Clinical Microbiology, 2009, 47, 1058-1062.	1.8	18
78	Bacterial vaginosis in female facility workers in north-western Tanzania: prevalence and risk factors. Sexually Transmitted Infections, 2009, 85, 370-375.	0.8	56
79	Risk factors for HIV incidence in women participating in an HSV suppressive treatment trial in Tanzania. Aids, 2009, 23, 415-422.	1.0	40
80	Effect of Herpes Simplex Suppression on Incidence of HIV among Women in Tanzania. New England Journal of Medicine, 2008, 358, 1560-1571.	13.9	344
81	Risk Factors for Herpes Simplex Virus Type 2 and HIV Among Women at High Risk in Northwestern Tanzania. Journal of Acquired Immune Deficiency Syndromes (1999), 2007, 46, 631-642.	0.9	64
82	Microbicides Development Program, Tanzania—Baseline Characteristics of an Occupational Cohort and Reattendance at 3 Months. Sexually Transmitted Diseases, 2007, 34, 638-643.	0.8	30
83	Biological and behavioural impact of an adolescent sexual health intervention in Tanzania: a community-randomized trial. Aids, 2007, 21, 1943-1955.	1.0	265
84	Low specificity of the Murex fourthâ€generation HIV enzyme immunoassay in Tanzanian adolescents. Tropical Medicine and International Health, 2007, 12, 1323-1326.	1.0	14
85	Impact of Azithromycin Administration for Trachoma Control on the Carriage of Antibiotic-Resistant Streptococcus pneumoniae. Antimicrobial Agents and Chemotherapy, 2003, 47, 2765-2769.	1.4	64
86	A comparison of the bactericidal activity of quinolone antibiotics in a Mycobacterium fortuitum model. Journal of Medical Microbiology, 2001, 50, 565-570.	0.7	37
87	Impact and Effectiveness of 13-Valent Pneumococcal Conjugate Vaccine on Population Incidence of Vaccine and Non-Vaccine Serotype Invasive Pneumococcal Disease in Blantyre, Malawi, 2006-2018: Prospective Observational Time-Series and Case-Control Studies. SSRN Electronic J <u>ournal, 0, , .</u>	0.4	4