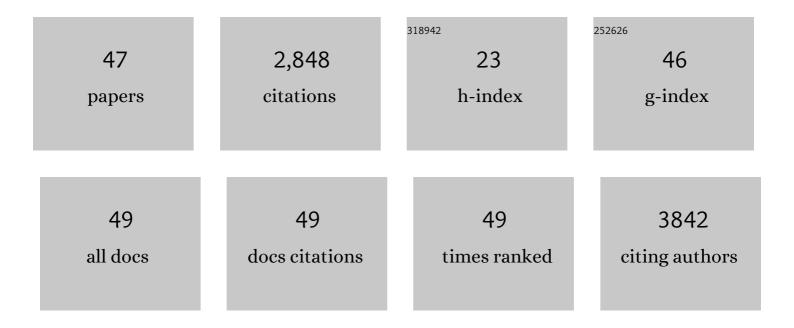
## Mignon du Plessis

List of Publications by Year in descending order

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#	Article	IF	CITATIONS
1	Early assessment of the clinical severity of the SARS-CoV-2 omicron variant in South Africa: a data linkage study. Lancet, The, 2022, 399, 437-446.	6.3	818
2	SARS-CoV-2 incidence, transmission, and reinfection in a rural and an urban setting: results of the PHIRST-C cohort study, South Africa, 2020–21. Lancet Infectious Diseases, The, 2022, 22, 821-834.	4.6	74
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	SARS-CoV-2 transmission, persistence of immunity, and estimates of Omicron's impact in South African population cohorts. Science Translational Medicine, 2022, 14, .	5.8	36
5	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
6	Visualizing variation within Global Pneumococcal Sequence Clusters (GPSCs) and country population snapshots to contextualize pneumococcal isolates. Microbial Genomics, 2020, 6, .	1.0	25
7	Pediatric Bacterial Meningitis Surveillance in the World Health Organization African Region Using the Invasive Bacterial Vaccine-Preventable Disease Surveillance Network, 2011–2016. Clinical Infectious Diseases, 2019, 69, S49-S57.	2.9	25
8	Joint sequencing of human and pathogen genomes reveals the genetics of pneumococcal meningitis. Nature Communications, 2019, 10, 2176.	5.8	83
9	International genomic definition of pneumococcal lineages, to contextualise disease, antibiotic resistance and vaccine impact. EBioMedicine, 2019, 43, 338-346.	2.7	168
10	Declining Incidence of Invasive Meningococcal Disease in South Africa: 2003–2016. Clinical Infectious Diseases, 2019, 69, 495-504.	2.9	10
11	Putative novel cps loci in a large global collection of pneumococci. Microbial Genomics, 2019, 5, .	1.0	14
12	Genomic differences among carriage and invasive nontypeable pneumococci circulating in South Africa. Microbial Genomics, 2019, 5, .	1.0	0
13	Invasive Disease Caused Simultaneously by Dual Serotypes of Streptococcus pneumoniae. Journal of Clinical Microbiology, 2018, 56, .	1.8	13
14	Global emergence and population dynamics of divergent serotype 3 CC180 pneumococci. PLoS Pathogens, 2018, 14, e1007438.	2.1	74
15	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
16	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
17	Two cases of serotypeable and non-serotypeable variants of Streptococcus pneumoniae detected simultaneously during invasive disease. BMC Microbiology, 2016, 16, 126.	1.3	2
18	Understanding pneumococcal serotype 1 biology through population genomic analysis. BMC Infectious Diseases, 2016, 16, 649.	1.3	22

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19	Phylogenetic Analysis of Invasive Serotype 1 Pneumococcus in South Africa, 1989 to 2013. Journal of Clinical Microbiology, 2016, 54, 1326-1334.	1.8	16
20	Genomic resolution of an aggressive, widespread, diverse and expanding meningococcal serogroup B, C and W lineage. Journal of Infection, 2015, 71, 544-552.	1.7	185
21	Region-specific diversification of the highly virulent serotype 1 Streptococcus pneumoniae. Microbial Genomics, 2015, 1, e000027.	1.0	27
22	Population Snapshot of Streptococcus pneumoniae Causing Invasive Disease in South Africa Prior to Introduction of Pneumococcal Conjugate Vaccines. PLoS ONE, 2014, 9, e107666.	1.1	18
23	High Nasopharyngeal Pneumococcal Density, Increased by Viral Coinfection, Is Associated With Invasive Pneumococcal Pneumonia. Journal of Infectious Diseases, 2014, 210, 1649-1657.	1.9	163
24	HIV and Influenza Virus Infections Are Associated With Increased Blood Pneumococcal Load: A Prospective, Hospital-Based Observational Study in South Africa, 2009-2011. Journal of Infectious Diseases, 2014, 209, 56-65.	1.9	30
25	Challenges of Using Molecular Serotyping for Surveillance of Pneumococcal Disease. Journal of Clinical Microbiology, 2014, 52, 3271-3276.	1.8	25
26	Meningococcal serogroup Y lpxL1 variants from South Africa are associated with clonal complex 23 among young adults. Journal of Infection, 2014, 68, 455-461.	1.7	6
27	Sequential Triplex Real-Time PCR Assay for Detecting 21 Pneumococcal Capsular Serotypes That Account for a High Global Disease Burden. Journal of Clinical Microbiology, 2013, 51, 647-652.	1.8	124
28	Population Snapshot of Invasive Serogroup B Meningococci in South Africa from 2005 to 2008. Journal of Clinical Microbiology, 2012, 50, 2577-2584.	1.8	6
29	Clinical Validation of Multiplex Real-Time PCR Assays for Detection of Bacterial Meningitis Pathogens. Journal of Clinical Microbiology, 2012, 50, 702-708.	1.8	116
30	Clonal Analysis of Neisseria meningitidis Serogroup B Strains in South Africa, 2002 to 2006: Emergence of New Clone ST-4240/6688. Journal of Clinical Microbiology, 2012, 50, 3678-3686.	1.8	5
31	Quadriplex real-time polymerase chain reaction (lytA, mef, erm, pbp2bwt) for pneumococcal detection and assessment of antibiotic susceptibility. Diagnostic Microbiology and Infectious Disease, 2011, 71, 453-456.	0.8	14
32	Distribution of factor H binding protein beyond serogroup B: Variation among five serogroups of invasive Neisseria meningitidis in South Africa. Vaccine, 2011, 29, 2187-2192.	1.7	21
33	Invasive Neisseria meningitidis with decreased susceptibility to fluoroquinolones in South Africa, 2009. Journal of Antimicrobial Chemotherapy, 2010, 65, 2258-2260.	1.3	15
34	An Unusual Pneumococcal Sequence Type Is the Predominant Cause of Serotype 3 Invasive Disease in South Africa. Journal of Clinical Microbiology, 2010, 48, 184-191.	1.8	17
35	High Prevalence of Dihydropteroate Synthase Mutations in <i>Pneumocystis jirovecii</i> Isolated from Patients with <i>Pneumocystis</i> Pneumonia in South Africa. Journal of Clinical Microbiology, 2010, 48, 2016-2021.	1.8	36
36	In Vitro Evaluation of the Antimicrobial Activity of Ceftaroline against Cephalosporin-Resistant Isolates of <i>Streptococcus pneumoniae</i> . Antimicrobial Agents and Chemotherapy, 2009, 53, 552-556.	1.4	65

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37	Sequence Diversity of the Factor H Binding Protein Vaccine Candidate in Epidemiologically Relevant Strains of Serogroup B <i>Neisseria meningitidis</i> . Journal of Infectious Diseases, 2009, 200, 379-389.	1.9	180
38	Molecular Characterization of Emerging Non-Levofloxacin-Susceptible Pneumococci Isolated from Children in South Africa. Journal of Clinical Microbiology, 2009, 47, 1319-1324.	1.8	17
39	Molecular basis and clonal nature of increasing pneumococcal macrolide resistance in South Africa, 2000–2005. International Journal of Antimicrobial Agents, 2008, 32, 62-67.	1.1	17
40	Serotype 6C is associated with penicillin-susceptible meningeal infections in human immunodeficiency virus (HIV)-infected adults among invasive pneumococcal isolates previously identified as serotype 6A in South Africa. International Journal of Antimicrobial Agents, 2008, 32, S66-S70.	1.1	23
41	<i>Neisseria meningitidis</i> Intermediately Resistant to Penicillin and Causing Invasive Disease in South Africa in 2001 to 2005. Journal of Clinical Microbiology, 2008, 46, 3208-3214.	1.8	29
42	Emergence of levofloxacin-non-susceptible Streptococcus pneumoniae and treatment for multidrug-resistant tuberculosis in children in South Africa: a cohort observational surveillance study. Lancet, The, 2008, 371, 1108-1113.	6.3	57
43	Emergence of Endemic Serogroup W135 Meningococcal Disease Associated with a High Mortality Rate in South Africa. Clinical Infectious Diseases, 2008, 46, 377-386.	2.9	88
44	Meningococcal Disease in South Africa, 1999–2002. Emerging Infectious Diseases, 2007, 13, 273-281.	2.0	37
45	Prevalence of DHPS Polymorphisms Associated with Sulfa Resistance in South African Pneumocystis jirovecii Strains. Journal of Eukaryotic Microbiology, 2006, 53, S110-S111.	0.8	5
46	Analysis of Penicillin-Binding Protein Genes of Clinical Isolates of Streptococcus pneumoniae with Reduced Susceptibility to Amoxicillin. Antimicrobial Agents and Chemotherapy, 2002, 46, 2349-2357.	1.4	75
47	Rapid discrimination between BRO β-lactamases from clinical isolates of Moraxella catarrhalis using restriction endonuclease analysis. Diagnostic Microbiology and Infectious Disease, 2001, 39, 65-67.	0.8	8