

# Dr Luria Leslie Founou

## List of Publications by Year in descending order

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Version: 2024-02-01

38  
papers

1,443  
citations

623734

14  
h-index

414414

32  
g-index

40  
all docs

40  
docs citations

40  
times ranked

2255  
citing authors

#	ARTICLE	IF	CITATIONS
1	Antibiotic Resistance in the Food Chain: A Developing Country-Perspective. <i>Frontiers in Microbiology</i> , 2016, 7, 1881.	3.5	467
2	Clinical and economic impact of antibiotic resistance in developing countries: A systematic review and meta-analysis. <i>PLoS ONE</i> , 2017, 12, e0189621.	2.5	422
3	Antibiotic Resistance in Food Animals in Africa: A Systematic Review and Meta-Analysis. <i>Microbial Drug Resistance</i> , 2018, 24, 648-665.	2.0	48
4	The potential impact of the COVID-19 pandemic on global antimicrobial and biocide resistance: an AMR Insights global perspective. <i>JAC-Antimicrobial Resistance</i> , 2021, 3, dlab038.	2.1	48
5	The Resistome, Mobilome, Virulome and Phylogenomics of Multidrug-Resistant <i>Escherichia coli</i> Clinical Isolates from Pretoria, South Africa. <i>Scientific Reports</i> , 2019, 9, 16457.	3.3	47
6	The COVID-19 pandemic: a threat to antimicrobial resistance containment. <i>Future Science OA</i> , 2021, 7, FSO736.	1.9	41
7	Genome Sequencing of Extended-Spectrum $\beta$ -Lactamase (ESBL)-Producing <i>Klebsiella pneumoniae</i> Isolated from Pigs and Abattoir Workers in Cameroon. <i>Frontiers in Microbiology</i> , 2018, 9, 188.	3.5	38
8	Whole Genome Sequencing of Extended Spectrum $\beta$ -lactamase (ESBL)-producing <i>Klebsiella pneumoniae</i> Isolated from Hospitalized Patients in KwaZulu-Natal, South Africa. <i>Scientific Reports</i> , 2019, 9, 6266.	3.3	36
9	Genomic analysis of methicillin-resistant <i>Staphylococcus aureus</i> isolated from poultry and occupational farm workers in Umgungundlovu District, South Africa. <i>Science of the Total Environment</i> , 2019, 670, 704-716.	8.0	33
10	Genomic analysis of a multidrug-resistant clinical <i>Providencia rettgeri</i> (PR002) strain with the novel integron $\ln$ 1483 and an A/C plasmid replicon. <i>Annals of the New York Academy of Sciences</i> , 2020, 1462, 92-103.	3.8	33
11	Pathogenomics and Evolutionary Epidemiology of Multi-Drug Resistant Clinical <i>Klebsiella pneumoniae</i> Isolated from Pretoria, South Africa. <i>Scientific Reports</i> , 2020, 10, 1232.	3.3	31
12	Extended spectrum beta-lactamase mediated resistance in carriage and clinical gram-negative ESKAPE bacteria: a comparative study between a district and tertiary hospital in South Africa. <i>Antimicrobial Resistance and Infection Control</i> , 2018, 7, 134.	4.1	29
13	Antimicrobial resistance in the farm-to-plate continuum: more than a food safety issue. <i>Future Science OA</i> , 2021, 7, FSO692.	1.9	24
14	Genomic characterisation of <i>Klebsiella michiganensis</i> co-producing OXA-181 and NDM-1 carbapenemases isolated from a cancer patient in uMgungundlovu District, KwaZulu-Natal Province, South Africa. <i>South African Medical Journal</i> , 2018, 109, 7.	0.6	17
15	Emergence and Spread of Extended Spectrum $\beta$ -Lactamase Producing Enterobacteriaceae (ESBL-PE) in Pigs and Exposed Workers: A Multicentre Comparative Study between Cameroon and South Africa. <i>Pathogens</i> , 2019, 8, 10.	2.8	16
16	Mannitol-fermenting methicillin-resistant staphylococci (MRS) in pig abattoirs in Cameroon and South Africa: A serious food safety threat. <i>International Journal of Food Microbiology</i> , 2018, 285, 50-60.	4.7	14
17	Genome analysis of methicillin-resistant <i>Staphylococcus aureus</i> isolated from pigs: Detection of the clonal lineage ST398 in Cameroon and South Africa. <i>Zoonoses and Public Health</i> , 2019, 66, 512-525.	2.2	13
18	Genomic Investigation of Carbapenem-Resistant <i>Klebsiella pneumoniae</i> Colonization in an Intensive Care Unit in South Africa. <i>Genes</i> , 2021, 12, 951.	2.4	11

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19	Bacterial diversity and functional profile of microbial populations on surfaces in public hospital environments in South Africa: A high throughput metagenomic analysis. <i>Science of the Total Environment</i> , 2020, 719, 137360.	8.0	10
20	$\hat{I}^2$ -lactam and fluoroquinolone resistance in Enterobacteriaceae from imported and locally-produced chicken in Mozambique. <i>Journal of Infection in Developing Countries</i> , 2020, 14, 471-478.	1.2	9
21	Extended-spectrum beta-lactamase-producing <i>Escherichia coli</i> harbouring mcr-1 gene isolated from pigs in South Africa. <i>South African Medical Journal</i> , 2018, 108, 796.	0.6	8
22	<i>Enterococcus faecalis</i> ST21 harbouring Tn6009 isolated from a carriage sample in South Africa. <i>South African Medical Journal</i> , 2021, 111, 98.	0.6	7
23	Genome Analysis of ESBL-Producing <i>Escherichia coli</i> Isolated from Pigs. <i>Pathogens</i> , 2022, 11, 776.	2.8	7
24	First report of a clinical multidrug-resistant <i>Pseudomonas aeruginosa</i> ST532 isolate harbouring a ciprofloxacin-modifying enzyme (CrpP) in South Africa. <i>Journal of Global Antimicrobial Resistance</i> , 2020, 22, 145-146.	2.2	6
25	Carbapenem Resistance Determinants Acquired through Novel Chromosomal Integrations in Extensively Drug-Resistant <i>Pseudomonas aeruginosa</i> . <i>Antimicrobial Agents and Chemotherapy</i> , 2021, 65, e0028921.	3.2	6
26	Models for increasing the health workforce. <i>South African Medical Journal</i> , 2012, 102, 830.	0.6	5
27	Draft genome sequence of an extended-spectrum $\hat{I}^2$ -lactamase (CTX-M-15)-producing <i>Escherichia coli</i> ST10 isolated from a nasal sample of an abattoir worker in Cameroon. <i>Journal of Global Antimicrobial Resistance</i> , 2018, 14, 68-69.	2.2	5
28	Multidrug-Resistant and Extended-Spectrum $\hat{I}^2$ -Lactamase (ESBL) - Producing Enterobacterales Isolated from Carriage Samples among HIV Infected Women in YaoundÃ©, Cameroon. <i>Pathogens</i> , 2022, 11, 504.	2.8	4
29	Draft genome sequence of a methicillin-resistant <i>Staphylococcus epidermidis</i> isolate from swine. <i>Journal of Global Antimicrobial Resistance</i> , 2018, 15, 250-251.	2.2	2
30	Draft genome sequences of extended-spectrum $\hat{I}^2$ -lactamase-producing <i>Enterobacter aerogenes</i> isolated from swine and human. <i>Journal of Global Antimicrobial Resistance</i> , 2018, 14, 70-71.	2.2	2
31	Methicillin resistant staphylococci isolated in clinical samples: a 3-yearÂretrospective study analysis. <i>Future Science OA</i> , 2021, 7, FSO681.	1.9	2
32	Genomic characterisation of <i>Staphylococcus aureus</i> ST121 isolated from hospitalised patients in South Africa. <i>South African Medical Journal</i> , 2018, 108, 1007.	0.6	1
33	Draft genome sequence of a clinical <i>Acinetobacter haemolyticus</i> isolate from South Africa. <i>Journal of Global Antimicrobial Resistance</i> , 2020, 20, 16-17.	2.2	0
34	Chronic Lymphocytic Leukemia in a Black African Man: A Cameroonian Case Report. , 2019, 07, .		0
35	Chronic Lymphocytic Leukemia in a Black African Man: A Cameroonian Case Report. , 2019, 07, .		0
36	Extended-spectrum beta-lactamase-producing <i>Klebsiella pneumoniae</i> isolated from an abattoir worker in Cameroon. <i>South African Medical Journal</i> , 2019, 109, 820.	0.6	0

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37	Methicillin-resistant <i>Staphylococcus haemolyticus</i> ST25 isolated from carriage samples in uMgungundlovu district, South Africa. <i>South African Medical Journal</i> , 2020, 110, 959.	0.6	0
38	Extraintestinal pathogenic <i>Escherichia coli</i> (ExPEC) ST221 isolated in the microbiome of swine in South Africa. <i>South African Medical Journal</i> , 2020, 111, 5.	0.6	0