

Geraint B Rogers

List of Publications by Citations

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The third column is the impact factor (IF) of the journal, and the fourth column is the number of citations of the article.

142
papers

5,508
citations

42
h-index

71
g-index

153
ext. papers

6,897
ext. citations

7.9
avg, IF

5.88
L-index

#	Paper	IF	Citations
142	Effect of azithromycin on asthma exacerbations and quality of life in adults with persistent uncontrolled asthma (AMAZES): a randomised, double-blind, placebo-controlled trial. <i>Lancet, The</i> , 2017 , 390, 659-668	4.0	348
141	Sample storage conditions significantly influence faecal microbiome profiles. <i>Scientific Reports</i> , 2015 , 5, 16350	4.9	257
140	characterization of bacterial community diversity in cystic fibrosis lung infections by use of 16s ribosomal DNA terminal restriction fragment length polymorphism profiling. <i>Journal of Clinical Microbiology</i> , 2004 , 42, 5176-83	9.7	251
139	Potentially pathogenic airway bacteria and neutrophilic inflammation in treatment resistant severe asthma. <i>PLoS ONE</i> , 2014 , 9, e100645	3.7	203
138	Partitioning core and satellite taxa from within cystic fibrosis lung bacterial communities. <i>ISME Journal</i> , 2011 , 5, 780-91	11.9	177
137	Bacterial diversity in cases of lung infection in cystic fibrosis patients: 16S ribosomal DNA (rDNA) length heterogeneity PCR and 16S rDNA terminal restriction fragment length polymorphism profiling. <i>Journal of Clinical Microbiology</i> , 2003 , 41, 3548-58	9.7	168
136	Inflammatory phenotypes in patients with severe asthma are associated with distinct airway microbiology. <i>Journal of Allergy and Clinical Immunology</i> , 2018 , 141, 94-103.e15	11.5	159
135	A novel microbiota stratification system predicts future exacerbations in bronchiectasis. <i>Annals of the American Thoracic Society</i> , 2014 , 11, 496-503	4.7	158
134	Clinical measures of disease in adult non-CF bronchiectasis correlate with airway microbiota composition. <i>Thorax</i> , 2013 , 68, 731-7	7.3	149
133	Deriving accurate microbiota profiles from human samples with low bacterial content through post-sequencing processing of Illumina MiSeq data. <i>Microbiome</i> , 2015 , 3, 19	16.6	126
132	Staphylococcus aureus small-colony variants are independently associated with worse lung disease in children with cystic fibrosis. <i>Clinical Infectious Diseases</i> , 2013 , 57, 384-91	11.6	117
131	Use of 16S rRNA gene profiling by terminal restriction fragment length polymorphism analysis to compare bacterial communities in sputum and mouthwash samples from patients with cystic fibrosis. <i>Journal of Clinical Microbiology</i> , 2006 , 44, 2601-4	9.7	115
130	The Diverse Metabolic Roles of Peripheral Serotonin. <i>Endocrinology</i> , 2017 , 158, 1049-1063	4.8	111
129	Long-term cultivation-independent microbial diversity analysis demonstrates that bacterial communities infecting the adult cystic fibrosis lung show stability and resilience. <i>Thorax</i> , 2012 , 67, 867-73	7.3	111
128	The effect of long-term macrolide treatment on respiratory microbiota composition in non-cystic fibrosis bronchiectasis: an analysis from the randomised, double-blind, placebo-controlled BLESS trial. <i>Lancet Respiratory Medicine</i> , 2014 , 2, 988-96	35.1	108
127	The Influence of the Gut Microbiome on Host Metabolism Through the Regulation of Gut Hormone Release. <i>Frontiers in Physiology</i> , 2019 , 10, 428	4.6	107
126	Low-Dose Nitric Oxide as Targeted Anti-biofilm Adjunctive Therapy to Treat Chronic Pseudomonas aeruginosa Infection in Cystic Fibrosis. <i>Molecular Therapy</i> , 2017 , 25, 2104-2116	11.7	106

125	Lean NAFLD: A Distinct Entity Shaped by Differential Metabolic Adaptation. <i>Hepatology</i> , 2020 , 71, 1213-1227	10.4	104
124	Does bacterial density in cystic fibrosis sputum increase prior to pulmonary exacerbation?. <i>Journal of Cystic Fibrosis</i> , 2011 , 10, 357-65	4.1	102
123	Long-Term Azithromycin Reduces and Increases Antibiotic Resistance in Severe Asthma. <i>American Journal of Respiratory and Critical Care Medicine</i> , 2019 , 200, 309-317	10.2	70
122	Three clinically distinct chronic pediatric airway infections share a common core microbiota. <i>Annals of the American Thoracic Society</i> , 2014 , 11, 1039-48	4.7	69
121	Bacterial activity in cystic fibrosis lung infections. <i>Respiratory Research</i> , 2005 , 6, 49	7.3	69
120	Respiratory microbiota resistance and resilience to pulmonary exacerbation and subsequent antimicrobial intervention. <i>ISME Journal</i> , 2016 , 10, 1081-91	11.9	68
119	Assessing the diagnostic importance of nonviable bacterial cells in respiratory infections. <i>Diagnostic Microbiology and Infectious Disease</i> , 2008 , 62, 133-41	2.9	67
118	Reducing bias in bacterial community analysis of lower respiratory infections. <i>ISME Journal</i> , 2013 , 7, 697-706	11.0	66
117	Studying bacterial infections through culture-independent approaches. <i>Journal of Medical Microbiology</i> , 2009 , 58, 1401-1418	3.2	60
116	Respiratory microbiota: addressing clinical questions, informing clinical practice. <i>Thorax</i> , 2015 , 70, 74-81	7.3	59
115	The Microbiota-Inflammasome Hypothesis of Major Depression. <i>BioEssays</i> , 2018 , 40, e1800027	4.1	58
114	Matrix metalloproteinases vary with airway microbiota composition and lung function in non-cystic fibrosis bronchiectasis. <i>Annals of the American Thoracic Society</i> , 2015 , 12, 701-7	4.7	57
113	Revealing the dynamics of polymicrobial infections: implications for antibiotic therapy. <i>Trends in Microbiology</i> , 2010 , 18, 357-64	12.4	56
112	Analysis of the bacterial communities present in lungs of patients with cystic fibrosis from American and British centers. <i>Journal of Clinical Microbiology</i> , 2011 , 49, 281-91	9.7	54
111	Interpreting infective microbiota: the importance of an ecological perspective. <i>Trends in Microbiology</i> , 2013 , 21, 271-6	12.4	53
110	The exclusion of dead bacterial cells is essential for accurate molecular analysis of clinical samples. <i>Clinical Microbiology and Infection</i> , 2010 , 16, 1656-8	9.5	53
109	Siblings of patients with Crohn's disease exhibit a biologically relevant dysbiosis in mucosal microbial metacommunities. <i>Gut</i> , 2016 , 65, 944-53	19.2	49
108	The gut microbiome regulates host glucose homeostasis via peripheral serotonin. <i>Proceedings of the National Academy of Sciences of the United States of America</i> , 2019 , 116, 19802-19804	11.5	49

107	Bacterial viability in faecal transplants: Which bacteria survive?. <i>EBioMedicine</i> , 2019 , 41, 509-516	8.8	49
106	Studying bacteria in respiratory specimens by using conventional and molecular microbiological approaches. <i>BMC Pulmonary Medicine</i> , 2009 , 9, 14	3.5	48
105	Impact of antibiotic treatment for pulmonary exacerbations on bacterial diversity in cystic fibrosis. <i>Journal of Cystic Fibrosis</i> , 2013 , 12, 22-8	4.1	47
104	Characterization of bacterial community diversity in chronic rhinosinusitis infections using novel culture-independent techniques. <i>American Journal of Rhinology and Allergy</i> , 2011 , 25, e133-40	2.4	46
103	Lung function and microbiota diversity in cystic fibrosis. <i>Microbiome</i> , 2020 , 8, 45	16.6	44
102	Predominant pathogen competition and core microbiota divergence in chronic airway infection. <i>ISME Journal</i> , 2015 , 9, 217-25	11.9	43
101	Determining cystic fibrosis-affected lung microbiology: comparison of spontaneous and serially induced sputum samples by use of terminal restriction fragment length polymorphism profiling. <i>Journal of Clinical Microbiology</i> , 2010 , 48, 78-86	9.7	43
100	The microbiome of otitis media with effusion in Indigenous Australian children. <i>International Journal of Pediatric Otorhinolaryngology</i> , 2015 , 79, 1548-55	1.7	40
99	DNA extraction approaches substantially influence the assessment of the human breast milk microbiome. <i>Scientific Reports</i> , 2020 , 10, 123	4.9	40
98	Comparing the microbiota of the cystic fibrosis lung and human gut. <i>Gut Microbes</i> , 2010 , 1, 85-93	8.8	40
97	Neuroimmunomodulation in Major Depressive Disorder: Focus on Caspase 1, Inducible Nitric Oxide Synthase, and Interferon-Gamma. <i>Molecular Neurobiology</i> , 2019 , 56, 4288-4305	6.2	37
96	Soluble fibre supplementation with and without a probiotic in adults with asthma: A 7-day randomised, double blind, three way cross-over trial. <i>EBioMedicine</i> , 2019 , 46, 473-485	8.8	34
95	Complexity, temporal stability, and clinical correlates of airway bacterial community composition in primary ciliary dyskinesia. <i>Journal of Clinical Microbiology</i> , 2013 , 51, 4029-35	9.7	33
94	Lung infections in cystic fibrosis: deriving clinical insight from microbial complexity. <i>Expert Review of Molecular Diagnostics</i> , 2010 , 10, 187-96	3.8	33
93	Not Just Antibiotics: Is Cancer Chemotherapy Driving Antimicrobial Resistance?. <i>Trends in Microbiology</i> , 2018 , 26, 393-400	12.4	33
92	Combined systems approaches reveal highly plastic responses to antimicrobial peptide challenge in <i>Escherichia coli</i> . <i>PLoS Pathogens</i> , 2014 , 10, e1004104	7.6	30
91	Next-generation sequencing in the analysis of human microbiota: essential considerations for clinical application. <i>Molecular Diagnosis and Therapy</i> , 2010 , 14, 343-50	4.5	30
90	Host-microbiome interactions in acute and chronic respiratory infections. <i>Cellular Microbiology</i> , 2016 , 18, 652-62	3.9	29

89	Impact of Long-Term Erythromycin Therapy on the Oropharyngeal Microbiome and Resistance Gene Reservoir in Non-Cystic Fibrosis Bronchiectasis. <i>MSphere</i> , 2018 , 3,	5	28
88	Infection 5 Sweet Tooth: How Glycans Mediate Infection and Disease Susceptibility. <i>Trends in Microbiology</i> , 2018 , 26, 92-101	12.4	27
87	Time between collection and storage significantly influences bacterial sequence composition in sputum samples from cystic fibrosis respiratory infections. <i>Journal of Clinical Microbiology</i> , 2014 , 52, 3011-6	9.7	27
86	Antibiotic exposure and interpersonal variance mask the effect of ivacaftor on respiratory microbiota composition. <i>Journal of Cystic Fibrosis</i> , 2018 , 17, 50-56	4.1	26
85	Ascites bacterial burden and immune cell profile are associated with poor clinical outcomes in the absence of overt infection. <i>PLoS ONE</i> , 2015 , 10, e0120642	3.7	25
84	Ascitic microbiota composition is correlated with clinical severity in cirrhosis with portal hypertension. <i>PLoS ONE</i> , 2013 , 8, e74884	3.7	25
83	genotype influences lung function, exacerbation frequency and airway microbiota in non-CF bronchiectasis. <i>Thorax</i> , 2017 , 72, 304-310	7.3	24
82	Precision respiratory medicine and the microbiome. <i>Lancet Respiratory Medicine</i> , 2016 , 4, 73-82	35.1	24
81	Characterisation of bacteria in ascites--reporting the potential of culture-independent, molecular analysis. <i>European Journal of Clinical Microbiology and Infectious Diseases</i> , 2010 , 29, 533-41	5.3	24
80	Effects of almond consumption on metabolic function and liver fat in overweight and obese adults with elevated fasting blood glucose: A randomised controlled trial. <i>Clinical Nutrition ESPEN</i> , 2019 , 30, 10-18	1.3	23
79	Enhancing the utility of existing antibiotics by targeting bacterial behaviour?. <i>British Journal of Pharmacology</i> , 2012 , 165, 845-57	8.6	23
78	Novel concepts in evaluating antimicrobial therapy for bacterial lung infections in patients with cystic fibrosis. <i>Journal of Cystic Fibrosis</i> , 2011 , 10, 387-400	4.1	23
77	Clinical and symptom scores are significantly correlated with fecal microbiota features in patients with symptomatic uncomplicated diverticular disease: a pilot study. <i>European Journal of Gastroenterology and Hepatology</i> , 2018 , 30, 107-112	2.2	23
76	Molecular detection of CF lung pathogens: current status and future potential. <i>Journal of Cystic Fibrosis</i> , 2013 , 12, 194-205	4.1	22
75	Neutrophil extracellular traps, disease severity, and antibiotic response in bronchiectasis: an international, observational, multicohort study. <i>Lancet Respiratory Medicine</i> , 2021 , 9, 873-884	35.1	22
74	Germs and joints: the contribution of the human microbiome to rheumatoid arthritis. <i>Nature Medicine</i> , 2015 , 21, 839-41	50.5	21
73	Opportunistic bacteria confer the ability to ferment prebiotic starch in the adult cystic fibrosis gut. <i>Gut Microbes</i> , 2019 , 10, 367-381	8.8	21
72	Implications of multiple freeze-thawing on respiratory samples for culture-independent analyses. <i>Journal of Cystic Fibrosis</i> , 2015 , 14, 464-7	4.1	20

71	Macrolide Treatment Inhibits <i>Pseudomonas aeruginosa</i> Quorum Sensing in Non-Cystic Fibrosis Bronchiectasis. An Analysis from the Bronchiectasis and Low-Dose Erythromycin Study Trial. <i>Annals of the American Thoracic Society</i> , 2016 , 13, 1697-1703	4.7	20
70	Divergent Relationships between Fecal Microbiota and Metabolome following Distinct Antibiotic-Induced Disruptions. <i>MSphere</i> , 2017 , 2,	5	18
69	Role of Dietary Flavonoid Compounds in Driving Patterns of Microbial Community Assembly. <i>MBio</i> , 2019 , 10,	7.8	18
68	A relationship between growth behaviour and cystic fibrosis patient lung function identified in a metabolomic investigation. <i>Metabolomics</i> , 2013 , 9, 1262	4.7	17
67	The use of culture-independent tools to characterize bacteria in endo-tracheal aspirates from pre-term infants at risk of bronchopulmonary dysplasia. <i>Journal of Perinatal Medicine</i> , 2010 , 38, 333-7	2.7	17
66	Adult non-cystic fibrosis bronchiectasis is characterised by airway luminal Th17 pathway activation. <i>PLoS ONE</i> , 2015 , 10, e0119325	3.7	17
65	Airway abundance of predicts response to azithromycin in adults with persistent uncontrolled asthma. <i>European Respiratory Journal</i> , 2020 , 56,	13.6	16
64	B Part of It study: a longitudinal study to assess carriage of <i>Neisseria meningitidis</i> in first year university students in South Australia. <i>Human Vaccines and Immunotherapeutics</i> , 2019 , 15, 987-994	4.4	16
63	Inbred Mouse Populations Exhibit Intergenerational Changes in Intestinal Microbiota Composition and Function Following Introduction to a Facility. <i>Frontiers in Microbiology</i> , 2017 , 8, 608	5.7	15
62	Using bacterial biomarkers to identify early indicators of cystic fibrosis pulmonary exacerbation onset. <i>Expert Review of Molecular Diagnostics</i> , 2011 , 11, 197-206	3.8	15
61	The CF gastrointestinal microbiome: Structure and clinical impact. <i>Pediatric Pulmonology</i> , 2016 , 51, S35-S44	5.4	15
60	The impact of CFTR modulator therapies on CF airway microbiology. <i>Journal of Cystic Fibrosis</i> , 2020 , 19, 359-364	4.1	15
59	Neutrophils in asthma: the good, the bad and the bacteria. <i>Thorax</i> , 2021 ,	7.3	15
58	Examining the Evidence for an Adult Healthy Middle Ear Microbiome. <i>MSphere</i> , 2019 , 4,	5	14
57	How can the cystic fibrosis respiratory microbiome influence our clinical decision-making?. <i>Current Opinion in Pulmonary Medicine</i> , 2017 , 23, 536-543	3	12
56	Bacterial community diversity in cultures derived from healthy and inflamed ileal pouches after restorative proctocolectomy. <i>Inflammatory Bowel Diseases</i> , 2009 , 15, 1803-11	4.5	12
55	Improving Risk-Benefit in Faecal Transplantation through Microbiome Screening. <i>Trends in Microbiology</i> , 2020 , 28, 331-339	12.4	11
54	Understanding the impact of antibiotic therapies on the respiratory tract resistome: a novel pooled-template metagenomic sequencing strategy. <i>Multidisciplinary Respiratory Medicine</i> , 2018 , 13, 30	3	11

53	Changes in the Composition of the Gut Microbiota and the Blood Transcriptome in Preterm Infants at Less than 29 Weeks Gestation Diagnosed with Bronchopulmonary Dysplasia. <i>MSystems</i> , 2019 , 4,	7.6	11
52	is an anti-inflammatory bacterium in the respiratory tract of patients with chronic lung disease. <i>European Respiratory Journal</i> , 2021 ,	13.6	11
51	Acute Colitis Drives Tolerance by Persistently Altering the Epithelial Barrier and Innate and Adaptive Immunity. <i>Inflammatory Bowel Diseases</i> , 2019 , 25, 1196-1207	4.5	9
50	Challenges and opportunities for faecal microbiota transplantation therapy. <i>Epidemiology and Infection</i> , 2013 , 141, 2235-42	4.3	9
49	Inclusivity and equity in human microbiome research. <i>Lancet, The</i> , 2019 , 393, 728-729	4.0	8
48	Burkholderia lata Infections from Intrinsically Contaminated Chlorhexidine Mouthwash, Australia, 2016. <i>Emerging Infectious Diseases</i> , 2018 , 24, 2109-2111	10.2	8
47	Mice lacking Casp1, Ifngr and Nos2 genes exhibit altered depressive- and anxiety-like behaviour, and gut microbiome composition. <i>Scientific Reports</i> , 2019 , 9, 6456	4.9	7
46	Culture-Independent Detection of Nontuberculous Mycobacteria in Clinical Respiratory Samples. <i>Journal of Clinical Microbiology</i> , 2016 , 54, 2395-8	9.7	7
45	Almond consumption affects fecal microbiota composition, stool pH, and stool moisture in overweight and obese adults with elevated fasting blood glucose: A randomized controlled trial. <i>Nutrition Research</i> , 2021 , 85, 47-59	4	7
44	The human microbiome: opportunities and challenges for clinical care. <i>Internal Medicine Journal</i> , 2015 , 45, 889-98	1.6	6
43	Total bacterial load, inflammation, and structural lung disease in paediatric cystic fibrosis. <i>Journal of Cystic Fibrosis</i> , 2020 , 19, 923-930	4.1	6
42	The Capacity of the Fecal Microbiota From Malawian Infants to Ferment Resistant Starch. <i>Frontiers in Microbiology</i> , 2019 , 10, 1459	5.7	6
41	Multi-centre ethics and research governance review can impede non-interventional clinical research. <i>Internal Medicine Journal</i> , 2019 , 49, 722-728	1.6	6
40	The contribution of respiratory microbiome analysis to a treatable traits model of care. <i>Respirology</i> , 2019 , 24, 19-28	3.6	6
39	Optimisation of a propidium monoazide based method to determine the viability of microbes in faecal slurries for transplantation. <i>Journal of Microbiological Methods</i> , 2019 , 156, 40-45	2.8	5
38	The composition of the gut microbiota following early-life antibiotic exposure affects host health and longevity in later life. <i>Cell Reports</i> , 2021 , 36, 109564	10.6	5
37	Is it time to rethink syphilis control?. <i>Clinical Infectious Diseases</i> , 2015 , 60, 325-6	11.6	4
36	National Trends in Antibiotic Use in Australian Residential Aged Care Facilities, 2005-2016. <i>Clinical Infectious Diseases</i> , 2021 , 72, 2167-2174	11.6	4

35	Safety and Efficacy of Using Nuts to Improve Bowel Health in Hemodialysis Patients. <i>Journal of Renal Nutrition</i> , 2020 , 30, 462-469	3	4
34	Exploring the parallel development of microbial systems in neonates with cystic fibrosis. <i>MBio</i> , 2012 , 3, e00408-12	7.8	4
33	Add-on azithromycin reduces sputum cytokines in non-eosinophilic asthma: an AMAZES substudy. <i>Thorax</i> , 2021 , 76, 733-736	7.3	4
32	The lung microbiome. <i>Emerging Topics in Life Sciences</i> , 2017 , 1, 313-324	3.5	3
31	Intestinal Microbiota Composition in Sudden Infant Death Syndrome and Age-Matched Controls. <i>Journal of Pediatrics</i> , 2017 , 191, 63-68.e1	3.6	3
30	Antibiotic stewardship in aged care facilities. <i>Lancet Infectious Diseases</i> , 2018 , 18, 1061-1063	25.5	3
29	Conventional myelosuppressive chemotherapy for non-haematological malignancy disrupts the intestinal microbiome. <i>BMC Cancer</i> , 2021 , 21, 591	4.8	2
28	The influence of early-life microbial exposures on long-term respiratory health. <i>Paediatric Respiratory Reviews</i> , 2021 , 40, 15-23	4.8	2
27	The effects of increasing fruit and vegetable intake in children with asthma: A randomized controlled trial. <i>Clinical and Experimental Allergy</i> , 2021 , 51, 1144-1156	4.1	2
26	Cellular Regulation of Peripheral Serotonin 2019 , 137-153		2
25	Establishment of murine gut microbiota in gnotobiotic mice. <i>iScience</i> , 2021 , 24, 102049	6.1	2
24	Draft Genome Sequence of a Non-O1/O139 Vibrio cholerae Strain Isolated from a Patient Presenting with Dysuria. <i>Microbiology Resource Announcements</i> , 2018 , 7,	1.3	2
23	Do we really understand how faecal microbiota transplantation works? AuthorsTreply. <i>EBioMedicine</i> , 2019 , 42, 40	8.8	1
22	Protect commensal gut bacteria to improve antimicrobial stewardship. <i>Clinical Microbiology and Infection</i> , 2020 , 26, 814-815	9.5	1
21	PPAR α s reduced in the airways of non-CF bronchiectasis subjects and is inversely correlated with the presence of Pseudomonas aeruginosa. <i>PLoS ONE</i> , 2018 , 13, e0202296	3.7	1
20	Draft Genome Sequences of Two subsp. Strains Isolated from Australian Hematology Patients with Bacteremia. <i>Genome Announcements</i> , 2017 , 5,		1
19	Gut Microbiome Regulation of Autophagic Flux and Neurodegenerative Disease Risks.. <i>Frontiers in Microbiology</i> , 2021 , 12, 817433	5.7	1
18	Intestinal microbiology shapes population health impacts of diet and lifestyle risk exposures in Torres Strait Islander communities. <i>ELife</i> , 2020 , 9,	8.9	1

17	Study protocol for a 9-month randomised controlled trial assessing the effects of almonds versus carbohydrate-rich snack foods on weight loss and weight maintenance. <i>BMJ Open</i> , 2020 , 10, e036542	3	1
16	The Structure of Relationships between the Human Exposome and Cardiometabolic Health: The Million Veteran Program. <i>Nutrients</i> , 2021 , 13,	6.7	1
15	Dietary yogurt is distinct from other dairy foods in its association with circulating lipid profile: Findings from the Million Veteran Program. <i>Clinical Nutrition ESPEN</i> , 2021 , 43, 456-463	1.3	1
14	A High Amylose Wheat Diet Improves Gastrointestinal Health Parameters and Gut Microbiota in Male and Female Mice. <i>Foods</i> , 2021 , 10,	4.9	1
13	Contribution of facility level factors to variation in antibiotic use in long-term care facilities: a national cohort study. <i>Journal of Antimicrobial Chemotherapy</i> , 2021 , 76, 1339-1348	5.1	1
12	Case report: Identification of intra-laboratory blood culture contamination with <i>Staphylococcus aureus</i> by whole genome sequencing. <i>Diagnostic Microbiology and Infectious Disease</i> , 2019 , 94, 331-333	2.9	0
11	The nasopharyngeal microbiome and LRTIs in infants. <i>Lancet Respiratory Medicine</i> , 2019 , 7, 369-371	35.1	0
10	Republished: Respiratory microbiota: addressing clinical questions, informing clinical practice. <i>Postgraduate Medical Journal</i> , 2015 , 91, 463-70	2	0
9	Chromosomal genes conferring tolerance to heavy metal (Ag) toxicity. <i>The Environmentalist</i> , 2009 , 29, 85-92		0
8	Ear microbiota and middle ear disease: a longitudinal pilot study of Aboriginal children in a remote south Australian setting.. <i>BMC Microbiology</i> , 2022 , 22, 24	4.5	0
7	The lung microbiome in chronic suppurative lung disease: cystic fibrosis and non-cystic fibrosis bronchiectasis 2019 , 158-172		0
6	Investigating potential transmission of antimicrobial resistance in an open-plan hospital ward: a cross-sectional metagenomic study of resistome dispersion in a lower middle-income setting. <i>Antimicrobial Resistance and Infection Control</i> , 2021 , 10, 56	6.2	0
5	Preservation of Gastrointestinal Mucosal Barrier Function and Microbiome in Patients With Controlled HIV Infection. <i>Frontiers in Immunology</i> , 2021 , 12, 688886	8.4	0
4	The cystic fibrosis gut as a potential source of multidrug resistant pathogens. <i>Journal of Cystic Fibrosis</i> , 2021 , 20, 413-420	4.1	0
3	Gut microbiota transplantation for colonization of germ-free mice. <i>STAR Protocols</i> , 2021 , 2, 100610	1.4	0
2	Assessment of Long-Term Macrolide Exposure on the Oropharyngeal Microbiome and Macrolide Resistance in Healthy Adults and Consequences for Onward Transmission of Resistance.. <i>Antimicrobial Agents and Chemotherapy</i> , 2022 , e0224621	5.9	0
1	The bronchiectasis microbiome	82-98	