Geraint B Rogers

List of Publications by Year
in descending order

150 papers	7,932 citations	46918 47 h-index	56606 83 g-index
153 all docs	153 docs citations	153 times ranked	9995 citing authors

#	Article	IF	Citations
1	Effect of azithromycin on asthma exacerbations and quality of life in adults with persistent uncontrolled asthma (AMAZES): a randomised, double-blind, placebo-controlled trial. Lancet, The, 2017, 390, 659-668.	6.3	489
2	Sample storage conditions significantly influence faecal microbiome profiles. Scientific Reports, 2015, 5, 16350.	1.6	350
3	Characterization of Bacterial Community Diversity in Cystic Fibrosis Lung Infections by Use of 16S Ribosomal DNA Terminal Restriction Fragment Length Polymorphism Profiling. Journal of Clinical Microbiology, 2004, 42, 5176-5183.	1.8	289
4	Potentially Pathogenic Airway Bacteria and Neutrophilic Inflammation in Treatment Resistant Severe Asthma. PLoS ONE, 2014, 9, e100645.	1.1	258
5	Inflammatory phenotypes in patients with severe asthma are associated with distinct airway microbiology. Journal of Allergy and Clinical Immunology, 2018, 141, 94-103.e15.	1.5	233
6	The Influence of the Gut Microbiome on Host Metabolism Through the Regulation of Gut Hormone Release. Frontiers in Physiology, 2019, 10, 428.	1.3	228
7	Partitioning core and satellite taxa from within cystic fibrosis lung bacterial communities. ISME Journal, 2011, 5, 780-791.	4.4	222
8	Lean NAFLD: A Distinct Entity Shaped by Differential Metabolic Adaptation. Hepatology, 2020, 71, 1213-1227.	3.6	209
9	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. Journal of Clinical Microbiology, 2003, 41, 3548-3558.	1.8	196
10	A Novel Microbiota Stratification System Predicts Future Exacerbations in Bronchiectasis. Annals of the American Thoracic Society, 2014, 11, 496-503.	1.5	183
11	Deriving accurate microbiota profiles from human samples with low bacterial content through post-sequencing processing of Illumina MiSeq data. Microbiome, 2015, 3, 19.	4.9	179
12	Clinical measures of disease in adult non-CF bronchiectasis correlate with airway microbiota composition. Thorax, 2013, 68, 731-737.	2.7	173
13	The Diverse Metabolic Roles of Peripheral Serotonin. Endocrinology, 2017, 158, 1049-1063.	1.4	164
14	Staphylococcus aureus Small-Colony Variants Are Independently Associated With Worse Lung Disease in Children With Cystic Fibrosis. Clinical Infectious Diseases, 2013, 57, 384-391.	2.9	153
15	Low-Dose Nitric Oxide as Targeted Anti-biofilm Adjunctive Therapy to Treat Chronic Pseudomonas aeruginosa Infection in Cystic Fibrosis. Molecular Therapy, 2017, 25, 2104-2116.	3.7	149
16	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. Lancet Respiratory Medicine,the, 2014, 2, 988-996.	5.2	146
17	Lung function and microbiota diversity in cystic fibrosis. Microbiome, 2020, 8, 45.	4.9	138
18	Long-term cultivation-independent microbial diversity analysis demonstrates that bacterial communities infecting the adult cystic fibrosis lung show stability and resilience. Thorax, 2012, 67, 867-873.	2.7	136

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19	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. Journal of Clinical Microbiology, 2006, 44, 2601-2604.	1.8	129
20	Does bacterial density in cystic fibrosis sputum increase prior to pulmonary exacerbation?. Journal of Cystic Fibrosis, 2011, 10, 357-365.	0.3	123
21	Long-Term Azithromycin Reduces Haemophilus influenzae and Increases Antibiotic Resistance in Severe Asthma. American Journal of Respiratory and Critical Care Medicine, 2019, 200, 309-317.	2.5	121
22	Neutrophil extracellular traps, disease severity, and antibiotic response in bronchiectasis: an international, observational, multicohort study. Lancet Respiratory Medicine,the, 2021, 9, 873-884.	5.2	99
23	Three Clinically Distinct Chronic Pediatric Airway Infections Share a Common Core Microbiota. Annals of the American Thoracic Society, 2014, 11, 1039-1048.	1.5	93
24	Respiratory microbiota resistance and resilience to pulmonary exacerbation and subsequent antimicrobial intervention. ISME Journal, 2016, 10, 1081-1091.	4.4	92
25	The Microbiotaâ€Inflammasome Hypothesis of Major Depression. BioEssays, 2018, 40, e1800027.	1.2	91
26	Bacterial activity in cystic fibrosis lung infections. Respiratory Research, 2005, 6, 49.	1.4	87
27	The gut microbiome regulates host glucose homeostasis via peripheral serotonin. Proceedings of the National Academy of Sciences of the United States of America, 2019, 116, 19802-19804.	3.3	84
28	Bacterial viability in faecal transplants: Which bacteria survive?. EBioMedicine, 2019, 41, 509-516.	2.7	84
29	Reducing bias in bacterial community analysis of lower respiratory infections. ISME Journal, 2013, 7, 697-706.	4.4	80
30	Matrix Metalloproteinases Vary with Airway Microbiota Composition and Lung Function in Nonâ€“Cystic Fibrosis Bronchiectasis. Annals of the American Thoracic Society, 2015, 12, 701-707.	1.5	77
31	Respiratory microbiota: addressing clinical questions, informing clinical practice. Thorax, 2015, 70, 74-81.	2.7	75
32	Assessing the diagnostic importance of nonviable bacterial cells in respiratory infections. Diagnostic Microbiology and Infectious Disease, 2008, 62, 133-141.	0.8	72
33	The exclusion of dead bacterial cells is essential for accurate molecular analysis of clinical samples. Clinical Microbiology and Infection, 2010, 16, 1656-1658.	2.8	69
34	Interpreting infective microbiota: the importance of an ecological perspective. Trends in Microbiology, 2013, 21, 271-276.	3.5	69
35	Revealing the dynamics of polymicrobial infections: implications for antibiotic therapy. Trends in Microbiology, 2010, 18, 357-364.	3.5	68
36	Studying bacterial infections through culture-independent approaches. Journal of Medical Microbiology, 2009, 58, 1401-1418.	0.7	67

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37	Soluble fibre supplementation with and without a probiotic in adults with asthma: A 7-day randomised, double blind, three way cross-over trial. EBioMedicine, 2019, 46, 473-485.	2.7	67
38	Neuroimmunomodulation in Major Depressive Disorder: Focus on Caspase 1, Inducible Nitric Oxide Synthase, and Interferon-Gamma. Molecular Neurobiology, 2019, 56, 4288-4305.	1.9	62
39	DNA extraction approaches substantially influence the assessment of the human breast milk microbiome. Scientific Reports, 2020, 10, 123.	1.6	62
40	Rothia mucilaginosa is an anti-inflammatory bacterium in the respiratory tract of patients with chronic lung disease. European Respiratory Journal, 2022, 59, 2101293.	3.1	60
41	Analysis of the Bacterial Communities Present in Lungs of Patients with Cystic Fibrosis from American and British Centers. Journal of Clinical Microbiology, 2011, 49, 281-291.	1.8	58
42	Impact of Long-Term Erythromycin Therapy on the Oropharyngeal Microbiome and Resistance Gene Reservoir in Non-Cystic Fibrosis Bronchiectasis. MSphere, 2018, 3, .	1.3	58
43	Neutrophils in asthma: the good, the bad and the bacteria. Thorax, 2021, 76, 835-844.	2.7	58
44	Siblings of patients with Crohnâ€™s disease exhibit a biologically relevant dysbiosis in mucosal microbial metacommunities. Gut, 2016, 65, 944-953.	6.1	56
45	Studying bacteria in respiratory specimens by using conventional and molecular microbiological approaches. BMC Pulmonary Medicine, 2009, 9, 14.	0.8	55
46	Predominant pathogen competition and core microbiota divergence in chronic airway infection. ISME Journal, 2015, 9, 217-225.	4.4	53
47	Characterization of Bacterial Community Diversity in Chronic Rhinosinusitis Infections Using Novel Culture-independent Techniques. American Journal of Rhinology and Allergy, 2011, 25, e133-e140.	1.0	52
48	The microbiome of otitis media with effusion in Indigenous Australian children. International Journal of Pediatric Otorhinolaryngology, 2015, 79, 1548-1555.	0.4	52
49	Impact of antibiotic treatment for pulmonary exacerbations on bacterial diversity in cystic fibrosis. Journal of Cystic Fibrosis, 2013, 12, 22-28.	0.3	50
50	Determining Cystic Fibrosis-Affected Lung Microbiology: Comparison of Spontaneous and Serially Induced Sputum Samples by Use of Terminal Restriction Fragment Length Polymorphism Profiling. Journal of Clinical Microbiology, 2010, 48, 78-86.	1.8	49
51	Comparing the microbiota of the cystic fibrosis lung and human gut. Gut Microbes, 2010, 1, 85-93.	4.3	47
52	Infectionâ€™s Sweet Tooth: How Glycans Mediate Infection and Disease Susceptibility. Trends in Microbiology, 2018, 26, 92-101.	3.5	47
53	Complexity, Temporal Stability, and Clinical Correlates of Airway Bacterial Community Composition in Primary Ciliary Dyskinesia. Journal of Clinical Microbiology, 2013, 51, 4029-4035.	1.8	46
54	Not Just Antibiotics: Is Cancer Chemotherapy Driving Antimicrobial Resistance?. Trends in Microbiology, 2018, 26, 393-400.	3.5	46

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55	Time between Collection and Storage Significantly Influences Bacterial Sequence Composition in Sputum Samples from Cystic Fibrosis Respiratory Infections. Journal of Clinical Microbiology, 2014, 52, 3011-3016.	1.8	43
56	Opportunistic bacteria confer the ability to ferment prebiotic starch in the adult cystic fibrosis gut. Gut Microbes, 2019, 10, 367-381.	4.3	39
57	The gut microbiome and mental health: advances in research and emerging priorities. Molecular Psychiatry, 2022, 27, 1908-1919.	4.1	39
58	Lung infections in cystic fibrosis: deriving clinical insight from microbial complexity. Expert Review of Molecular Diagnostics, 2010, 10, 187-196.	1.5	38
59	Ascites Bacterial Burden and Immune Cell Profile Are Associated with Poor Clinical Outcomes in the Absence of Overt Infection. PLoS ONE, 2015, 10, e0120642.	1.1	38
60	Combined Systems Approaches Reveal Highly Plastic Responses to Antimicrobial Peptide Challenge in Escherichia coli. PLoS Pathogens, 2014, 10, e1004104.	2.1	37
61	Antibiotic exposure and interpersonal variance mask the effect of ivacaftor on respiratory microbiota composition. Journal of Cystic Fibrosis, 2018, 17, 50-56.	0.3	37
62	Effects of almond consumption on metabolic function and liver fat in overweight and obese adults with elevated fasting blood glucose: A randomised controlled trial. Clinical Nutrition ESPEN, 2019, 30, 10-18.	0.5	36
63	Ascitic Microbiota Composition Is Correlated with Clinical Severity in Cirrhosis with Portal Hypertension. PLoS ONE, 2013, 8, e74884.	1.1	36
64	Next-Generation Sequencing in the Analysis of Human Microbiota. Molecular Diagnosis and Therapy, 2010, 14, 343-350.	1.6	35
65	Host-microbiome interactions in acute and chronic respiratory infections. Cellular Microbiology, 2016, 18, 652-662.	1.1	35
66	FUT2genotype influences lung function, exacerbation frequency and airway microbiota in non-CF bronchiectasis. Thorax, 2017, 72, 304-310.	2.7	35
67	Clinical and symptom scores are significantly correlated with fecal microbiota features in patients with symptomatic uncomplicated diverticular disease. European Journal of Gastroenterology and Hepatology, 2018, 30, 107-112.	0.8	33
68	The impact of CFTR modulator therapies on CF airway microbiology. Journal of Cystic Fibrosis, 2020, 19, 359-364.	0.3	33
69	Divergent Relationships between Fecal Microbiota and Metabolome following Distinct Antibiotic-Induced Disruptions. MSphere, 2017, 2, .	1.3	31
70	Airway abundance of Haemophilus influenzae predicts response to azithromycin in adults with persistent uncontrolled asthma. European Respiratory Journal, 2020, 56, 2000194.	3.1	31
71	The composition of the gut microbiota following early-life antibiotic exposure affects host health and longevity in later life. Cell Reports, 2021, 36, 109564.	2.9	31
72	Implications of multiple freeze-thawing on respiratory samples for culture-independent analyses. Journal of Cystic Fibrosis, 2015, 14, 464-467.	0.3	29

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73	Germs and joints: the contribution of the human microbiome to rheumatoid arthritis. Nature Medicine, 2015, 21, 839-841.	15.2	29
74	Characterisation of bacteria in ascitesâ€"reporting the potential of culture-independent, molecular analysis. European Journal of Clinical Microbiology and Infectious Diseases, 2010, 29, 533-541.	1.3	28
75	Enhancing the utility of existing antibiotics by targeting bacterial behaviour?. British Journal of Pharmacology, 2012, 165, 845-857.	2.7	28
76	B Part of It study: a longitudinal study to assess carriage of Neisseria meningitidis in first year university students in South Australia. Human Vaccines and Immunotherapeutics, 2019, 15, 987-994.	1.4	28
77	The CF gastrointestinal microbiome: Structure and clinical impact. Pediatric Pulmonology, 2016, 51, S35-S44.	1.0	27
78	Precision respiratory medicine and the microbiome. Lancet Respiratory Medicine,the, 2016, 4, 73-82.	5.2	27
79	Role of Dietary Flavonoid Compounds in Driving Patterns of Microbial Community Assembly. MBio, 2019, 10, .	1.8	27
80	Macrolide Treatment InhibitsPseudomonas aeruginosaQuorum Sensing in Non-CF Bronchiectasis: An Analysis from the BLESS Trial. Annals of the American Thoracic Society, 2016, 13, 1697-1703.	1.5	26
81	Molecular detection of CF lung pathogens: Current status and future potential. Journal of Cystic Fibrosis, 2013, 12, 194-205.	0.3	25
82	Novel concepts in evaluating antimicrobial therapy for bacterial lung infections in patients with cystic fibrosis. Journal of Cystic Fibrosis, 2011, 10, 387-400.	0.3	23
83	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. MSystems, 2019, 4, .	1.7	23
84	The use of culture-independent tools to characterize bacteria in endo-tracheal aspirates from pre-term infants at risk of bronchopulmonary dysplasia. Journal of Perinatal Medicine, 2010, 38, 333-7.	0.6	22
85	Examining the Evidence for an Adult Healthy Middle Ear Microbiome. MSphere, 2019, 4, .	1.3	22
86	Inbred Mouse Populations Exhibit Intergenerational Changes in Intestinal Microbiota Composition and Function Following Introduction to a Facility. Frontiers in Microbiology, 2017, 8, 608.	1.5	21
87	Improving Riskâ€"Benefit in Faecal Transplantation through Microbiome Screening. Trends in Microbiology, 2020, 28, 331-339.	3.5	19
88	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. Nutrition Research, 2021, 85, 47-59.	1.3	19
89	A relationship between Pseudomonal growth behaviour and cystic fibrosis patient lung function identified in a metabolomic investigation. Metabolomics, 2013, 9, 1262-1273.	1.4	18
90	Adult Non-Cystic Fibrosis Bronchiectasis Is Characterised by Airway Luminal Th17 Pathway Activation. PLoS ONE, 2015, 10, e0119325.	1.1	18

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91	Understanding the impact of antibiotic therapies on the respiratory tract resistome: a novel pooled-template metagenomic sequencing strategy. Multidisciplinary Respiratory Medicine, 2018, 13, 30.	0.6	17
92	Using bacterial biomarkers to identify early indicators of cystic fibrosis pulmonary exacerbation onset. Expert Review of Molecular Diagnostics, 2011, 11, 197-206.	1.5	16
93	Burkholderia lata Infections from Intrinsically Contaminated Chlorhexidine Mouthwash, Australia, 2016. Emerging Infectious Diseases, 2018, 24, 2109-2111.	2.0	16
94	Add-on azithromycin reduces sputum cytokines in non-eosinophilic asthma: an AMAZES substudy. Thorax, 2021, 76, 733-736.	2.7	16
95	The effects of increasing fruit and vegetable intake in children with asthma: A randomized controlled trial. Clinical and Experimental Allergy, 2021, 51, 1144-1156.	1.4	16
96	Mice lacking Casp1, Ifngr and Nos2 genes exhibit altered depressive- and anxiety-like behaviour, and gut microbiome composition. Scientific Reports, 2019, 9, 6456.	1.6	15
97	Optimisation of a propidium monoazide based method to determine the viability of microbes in faecal slurries for transplantation. Journal of Microbiological Methods, 2019, 156, 40-45.	0.7	15
98	Total bacterial load, inflammation, and structural lung disease in paediatric cystic fibrosis. Journal of Cystic Fibrosis, 2020, 19, 923-930.	0.3	15
99	Bacterial community diversity in cultures derived from healthy and inflamed ileal pouches after restorative proctocolectomy. Inflammatory Bowel Diseases, 2009, 15, 1803-1811.	0.9	13
100	How can the cystic fibrosis respiratory microbiome influence our clinical decision-making?. Current Opinion in Pulmonary Medicine, 2017, 23, 536-543.	1.2	13
101	Inclusivity and equity in human microbiome research. Lancet, The, 2019, 393, 728-729.	6.3	13
102	The cystic fibrosis gut as a potential source of multidrug resistant pathogens. Journal of Cystic Fibrosis, 2021, 20, 413-420.	0.3	13
103	Establishment of murine gut microbiota in gnotobiotic mice. IScience, 2021, 24, 102049.	1.9	13
104	National Trends in Antibiotic Use in Australian Residential Aged Care Facilities, 2005â€“2016. Clinical Infectious Diseases, 2021, 72, 2167-2174.	2.9	12
105	Culture-Independent Detection of Nontuberculous Mycobacteria in Clinical Respiratory Samples. Journal of Clinical Microbiology, 2016, 54, 2395-2398.	1.8	11
106	The Capacity of the Fecal Microbiota From Malawian Infants to Ferment Resistant Starch. Frontiers in Microbiology, 2019, 10, 1459.	1.5	11
107	Multiâ€centre ethics and research governance review can impede nonâ€interventional clinical research. Internal Medicine Journal, 2019, 49, 722-728.	0.5	11
108	Conventional myelosuppressive chemotherapy for non-haematological malignancy disrupts the intestinal microbiome. BMC Cancer, 2021, 21, 591.	1.1	11

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109	Challenges and opportunities for faecal microbiota transplantation therapy. Epidemiology and Infection, 2013, 141, 2235-2242.	1.0	10
110	Acute Colitis Drives Tolerance by Persistently Altering the Epithelial Barrier and Innate and Adaptive Immunity. Inflammatory Bowel Diseases, 2019, 25, 1196-1207.	0.9	10
111	Safety and Efficacy of Using Nuts to Improve Bowel Health in Hemodialysis Patients. , 2020, 30, 462-469.		10
112	Preservation of Gastrointestinal Mucosal Barrier Function and Microbiome in Patients With Controlled HIV Infection. Frontiers in Immunology, 2021, 12, 688886.	2.2	9
113	Intestinal Microbiota Composition in Sudden Infant Death Syndrome and Age-Matched Controls. Journal of Pediatrics, 2017, 191, 63-68.e1.	0.9	8
114	The contribution of respiratory microbiome analysis to a treatable traits model of care. Respirology, 2019, 24, 19-28.	1.3	8
115	The human microbiome: opportunities and challenges for clinical care. Internal Medicine Journal, 2015, 45, 889-898.	0.5	7
116	A High Amylose Wheat Diet Improves Gastrointestinal Health Parameters and Gut Microbiota in Male and Female Mice. Foods, 2021, 10, 220.	1.9	7
117	Contribution of facility level factors to variation in antibiotic use in long-term care facilities: a national cohort study. Journal of Antimicrobial Chemotherapy, 2021, 76, 1339-1348.	1.3	7
118	Gut Microbiome Regulation of Autophagic Flux and Neurodegenerative Disease Risks. Frontiers in Microbiology, 2021, 12, 817433.	1.5	7
119	Assessment of Long-Term Macrolide Exposure on the Oropharyngeal Microbiome and Macrolide Resistance in Healthy Adults and Consequences for Onward Transmission of Resistance. Antimicrobial Agents and Chemotherapy, 2022, 66, e0224621.	1.4	6
120	Exploring the Parallel Development of Microbial Systems in Neonates with Cystic Fibrosis. MBio, 2012, 3, e00408-12.	1.8	5
121	Is It Time to Rethink Syphilis Control?. Clinical Infectious Diseases, 2015, 60, 325-326.	2.9	5
122	The lung microbiome. Emerging Topics in Life Sciences, 2017, 1, 313-324.	1.1	5
123	Gut microbiota transplantation for colonization of germ-free mice. STAR Protocols, 2021, 2, 100610.	0.5	5
124	Environmental dynamics of hospital microbiome upon transfer from a major hospital to a new facility. Journal of Infection, 2021, 83, 637-643.	1.7	5
125	Intestinal microbiology shapes population health impacts of diet and lifestyle risk exposures in Torres Strait Islander communities. ELife, 2020, 9, .	2.8	5
126	Ear microbiota and middle ear disease: a longitudinal pilot study of Aboriginal children in a remote south Australian setting. BMC Microbiology, 2022, 22, 24.	1.3	5

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127	Antibiotic stewardship in aged care facilities. Lancet Infectious Diseases, The, 2018, 18, 1061-1063.	4.6	4
128	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. BMJ Open, 2020, 10, e036542.	0.8	4
129	The Structure of Relationships between the Human Exposome and Cardiometabolic Health: The Million Veteran Program. Nutrients, 2021, 13, 1364.	1.7	4
130	The influence of early-life microbial exposures on long-term respiratory health. Paediatric Respiratory Reviews, 2021, 40, 15-23.	1.2	4
131	Draft Genome Sequence of a Non-O1/O139 Vibrio cholerae Strain Isolated from a Patient Presenting with Dysuria. Microbiology Resource Announcements, 2018, 7, .	0.3	3
132	Cellular Regulation of Peripheral Serotonin. , 2019, , 137-153.		3
133	Dietary yogurt is distinct from other dairy foods in its association with circulating lipid profile: Findings from the Million Veteran Program. Clinical Nutrition ESPEN, 2021, 43, 456-463.	0.5	3
134	Draft Genome Sequences of Two Enterobacter cloacae subsp. cloacae Strains Isolated from Australian Hematology Patients with Bacteremia. Genome Announcements, 2017, 5, .	0.8	2
135	Inflammation, age and changing microbiology: the search for causation in the cystic fibrosis airways. European Respiratory Journal, 2017, 50, 1701935.	3.1	2
136	Case report: Identification of intra-laboratory blood culture contamination with Staphylococcus aureus by whole genome sequencing. Diagnostic Microbiology and Infectious Disease, 2019, 94, 331-333.	0.8	2
137	The nasopharyngeal microbiome and LRTIs in infants. Lancet Respiratory Medicine,the, 2019, 7, 369-371.	5.2	2
138	Protect commensal gut bacteria to improve antimicrobial stewardship. Clinical Microbiology and Infection, 2020, 26, 814-815.	2.8	2
139	Chromosomal genes conferring tolerance to heavy metal (Ag) toxicity. The Environmentalist, 2009, 29, 85-92.	0.7	1
140	Republished: Respiratory microbiota: addressing clinical questions, informing clinical practice. Postgraduate Medical Journal, 2015, 91, 463-470.	0.9	1
141	PPARÎ³ is reduced in the airways of non-CF bronchiectasis subjects and is inversely correlated with the presence of Pseudomonas aeruginosa. PLoS ONE, 2018, 13, e0202296.	1.1	1
142	Do we really understand how faecal microbiota transplantation works? Authors' reply. EBioMedicine, 2019, 42, 40.	2.7	1
143	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. Antimicrobial Resistance and Infection Control, 2021, 10, 56.	1.5	1
144	The lung microbiome in chronic suppurative lung disease: cystic fibrosis and non-cystic fibrosis bronchiectasis. , 2019, , 158-172.		1

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145	Cystic Fibrosis Pulmonary Exacerbations: The Role Of Bacterial Community Structure Beyond Pseudomonas Aeruginosa. , 2010, , .		0
146	150 Siblings of Crohn's Disease Patients Exhibit a Pathologically Relevant Dysbiosis: Examination of Mucosal Microbiota Communities Using 16S rRNA Gene Pyrosequencing. Gastroenterology, 2014, 146, S-42.	0.6	0
147	Microbiology review series: CF microbiology â€" Towards 2020 and beyond. Journal of Cystic Fibrosis, 2015, 14, 289-290.	0.3	0
148	Airway microbiome studies challenge simplistic models of inhaled tobramycin benefit. Thorax, 2020, 75, 1031-1032.	2.7	0
149	The bronchiectasis microbiome. , 0, , 82-98.		0
150	Carriage and Transmission of Macrolide Resistance Genes in Patients With Chronic Respiratory Conditions and Their Close Contacts. Chest, 2022, 162, 56-65.	0.4	0