Geraint B Rogers

List of Publications by Year in Descending Order

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	Ear microbiota and middle ear disease: a longitudinal pilot study of Aboriginal children in a remote south Australian setting.. *BMC Microbiology*, **2022**, 22, 24	4.5	0
141	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**, e0224621	5.9	0
140	National Trends in Antibiotic Use in Australian Residential Aged Care Facilities, 2005-2016. *Clinical Infectious Diseases*, **2021**, 72, 2167-2174	11.6	4
139	Gut Microbiome Regulation of Autophagic Flux and Neurodegenerative Disease Risks.. *Frontiers in Microbiology*, **2021**, 12, 817433	5.7	1
138	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	6.2	0
137	The Structure of Relationships between the Human Exposome and Cardiometabolic Health: The Million Veteran Program. *Nutrients*, **2021**, 13,	6.7	1
136	Conventional myelosuppressive chemotherapy for non-haematological malignancy disrupts the intestinal microbiome. *BMC Cancer*, **2021**, 21, 591	4.8	2
135	Preservation of Gastrointestinal Mucosal Barrier Function and Microbiome in Patients With Controlled HIV Infection. *Frontiers in Immunology*, **2021**, 12, 688886	8.4	0
134	The influence of early-life microbial exposures on long-term respiratory health. *Paediatric Respiratory Reviews*, **2021**, 40, 15-23	4.8	2
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	1.3	1
132	The effects of increasing fruit and vegetable intake in children with asthma: A randomized controlled trial. *Clinical and Experimental Allergy*, **2021**, 51, 1144-1156	4.1	2
131	The cystic fibrosis gut as a potential source of multidrug resistant pathogens. *Journal of Cystic Fibrosis*, **2021**, 20, 413-420	4.1	0
130	A High Amylose Wheat Diet Improves Gastrointestinal Health Parameters and Gut Microbiota in Male and Female Mice. *Foods*, **2021**, 10,	4.9	1
129	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	4	7
128	Add-on azithromycin reduces sputum cytokines in non-eosinophilic asthma: an AMAZES substudy. *Thorax*, **2021**, 76, 733-736	7.3	4
127	Neutrophils in asthma: the good, the bad and the bacteria. *Thorax*, **2021**,	7.3	15
126	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	5.1	1

125	Establishment of murine gut microbiota in gnotobiotic mice. *IScience*, **2021**, 24, 102049	6.1	2
124	The composition of the gut microbiota following early-life antibiotic exposure affects host health and longevity in later life. *Cell Reports*, **2021**, 36, 109564	10.6	5
123	Neutrophil extracellular traps, disease severity, and antibiotic response in bronchiectasis: an international, observational, multicohort study. *Lancet Respiratory Medicine,the*, **2021**, 9, 873-884	35.1	22
122	is an anti-inflammatory bacterium in the respiratory tract of patients with chronic lung disease. *European Respiratory Journal*, **2021**,	13.6	11
121	Gut microbiota transplantation for colonization of germ-free mice. *STAR Protocols*, **2021**, 2, 100610	1.4	0
120	Airway abundance of predicts response to azithromycin in adults with persistent uncontrolled asthma. *European Respiratory Journal*, **2020**, 56,	13.6	16
119	Total bacterial load, inflammation, and structural lung disease in paediatric cystic fibrosis. *Journal of Cystic Fibrosis*, **2020**, 19, 923-930	4.1	6
118	Lung function and microbiota diversity in cystic fibrosis. *Microbiome*, **2020**, 8, 45	16.6	44
117	Protect commensal gut bacteria to improve antimicrobial stewardship. *Clinical Microbiology and Infection*, **2020**, 26, 814-815	9.5	1
116	Improving Risk-Benefit in Faecal Transplantation through Microbiome Screening. *Trends in Microbiology*, **2020**, 28, 331-339	12.4	11
115	DNA extraction approaches substantially influence the assessment of the human breast milk microbiome. *Scientific Reports*, **2020**, 10, 123	4.9	40
114	Safety and Efficacy of Using Nuts to Improve Bowel Health in Hemodialysis Patients. *Journal of Renal Nutrition*, **2020**, 30, 462-469	3	4
113	Intestinal microbiology shapes population health impacts of diet and lifestyle risk exposures in Torres Strait Islander communities. *ELife*, **2020**, 9,	8.9	1
112	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	3	1
111	The impact of CFTR modulator therapies on CF airway microbiology. *Journal of Cystic Fibrosis*, **2020**, 19, 359-364	4.1	15
110	Lean NAFLD: A Distinct Entity Shaped by Differential Metabolic Adaptation. *Hepatology*, **2020**, 71, 1213-1227	11.2	104
109	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	11.5	49
108	Mice lacking Casp1, Ifngr and Nos2 genes exhibit altered depressive- and anxiety-like behaviour, and gut microbiome composition. *Scientific Reports*, **2019**, 9, 6456	4.9	7

107	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	2.9	0
106	The nasopharyngeal microbiome and LRTIs in infants. *Lancet Respiratory Medicine,the*, **2019**, 7, 369-371	35.1	0
105	Long-Term Azithromycin Reduces and Increases Antibiotic Resistance in Severe Asthma. *American Journal of Respiratory and Critical Care Medicine*, **2019**, 200, 309-317	10.2	70
104	Do we really understand how faecal microbiota transplantation works? AuthorsTreply. *EBioMedicine* , **2019**, 42, 40	8.8	1
103	The Influence of the Gut Microbiome on Host Metabolism Through the Regulation of Gut Hormone Release. *Frontiers in Physiology*, **2019**, 10, 428	4.6	107
102	Inclusivity and equity in human microbiome research. *Lancet, The*, **2019**, 393, 728-729	40	8
101	Bacterial viability in faecal transplants: Which bacteria survive?. *EBioMedicine*, **2019**, 41, 509-516	8.8	49
100	Acute Colitis Drives Tolerance by Persistently Altering the Epithelial Barrier and Innate and Adaptive Immunity. *Inflammatory Bowel Diseases*, **2019**, 25, 1196-1207	4.5	9
99	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	1.3	23
98	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	8.8	34
97	The Capacity of the Fecal Microbiota From Malawian Infants to Ferment Resistant Starch. *Frontiers in Microbiology*, **2019**, 10, 1459	5.7	6
96	Role of Dietary Flavonoid Compounds in Driving Patterns of Microbial Community Assembly. *MBio*, **2019**, 10,	7.8	18
95	The lung microbiome in chronic suppurative lung disease: cystic fibrosis and non-cystic fibrosis bronchiectasis **2019**, 158-172		0
94	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,	7.6	11
93	Examining the Evidence for an Adult Healthy Middle Ear Microbiome. *MSphere*, **2019**, 4,	5	14
92	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	2.8	5
91	Multi-centre ethics and research governance review can impede non-interventional clinical research. *Internal Medicine Journal*, **2019**, 49, 722-728	1.6	6
90	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	4.4	16

89	Opportunistic bacteria confer the ability to ferment prebiotic starch in the adult cystic fibrosis gut. *Gut Microbes*, **2019**, 10, 367-381	8.8	21
88	Cellular Regulation of Peripheral Serotonin **2019**, 137-153		2
87	Neuroimmunomodulation in Major Depressive Disorder: Focus on Caspase 1, Inducible Nitric Oxide Synthase, and Interferon-Gamma. *Molecular Neurobiology*, **2019**, 56, 4288-4305	6.2	37
86	The contribution of respiratory microbiome analysis to a treatable traits model of care. *Respirology*, **2019**, 24, 19-28	3.6	6
85	Impact of Long-Term Erythromycin Therapy on the Oropharyngeal Microbiome and Resistance Gene Reservoir in Non-Cystic Fibrosis Bronchiectasis. *MSphere*, **2018**, 3,	5	28
84	Inflammatory phenotypes in patients with severe asthma are associated with distinct airway microbiology. *Journal of Allergy and Clinical Immunology*, **2018**, 141, 94-103.e15	11.5	159
83	Antibiotic exposure and interpersonal variance mask the effect of ivacaftor on respiratory microbiota composition. *Journal of Cystic Fibrosis*, **2018**, 17, 50-56	4.1	26
82	Infection Ts Sweet Tooth: How Glycans Mediate Infection and Disease Susceptibility. *Trends in Microbiology*, **2018**, 26, 92-101	12.4	27
81	The Microbiota-Inflammasome Hypothesis of Major Depression. *BioEssays*, **2018**, 40, e1800027	4.1	58
80	PPAR s reduced in the airways of non-CF bronchiectasis subjects and is inversely correlated with the presence of Pseudomonas aeruginosa. *PLoS ONE*, **2018**, 13, e0202296	3.7	1
79	Understanding the impact of antibiotic therapies on the respiratory tract resistome: a novel pooled-template metagenomic sequencing strategy. *Multidisciplinary Respiratory Medicine*, **2018**, 13, 30	3	11
78	Clinical and symptom scores are significantly correlated with fecal microbiota features in patients with symptomatic uncomplicated diverticular disease: a pilot study. *European Journal of Gastroenterology and Hepatology*, **2018**, 30, 107-112	2.2	23
77	Not Just Antibiotics: Is Cancer Chemotherapy Driving Antimicrobial Resistance?. *Trends in Microbiology*, **2018**, 26, 393-400	12.4	33
76	Antibiotic stewardship in aged care facilities. *Lancet Infectious Diseases, The*, **2018**, 18, 1061-1063	25.5	3
75	Burkholderia lata Infections from Intrinsically Contaminated Chlorhexidine Mouthwash, Australia, 2016. *Emerging Infectious Diseases*, **2018**, 24, 2109-2111	10.2	8
74	Draft Genome Sequence of a Non-O1/O139 Vibrio cholerae Strain Isolated from a Patient Presenting with Dysuria. *Microbiology Resource Announcements*, **2018**, 7,	1.3	2
73	genotype influences lung function, exacerbation frequency and airway microbiota in non-CF bronchiectasis. *Thorax*, **2017**, 72, 304-310	7.3	24
72	Divergent Relationships between Fecal Microbiota and Metabolome following Distinct Antibiotic-Induced Disruptions. *MSphere*, **2017**, 2,	5	18

71	The Diverse Metabolic Roles of Peripheral Serotonin. *Endocrinology*, **2017**, 158, 1049-1063	4.8	111
70	The lung microbiome. *Emerging Topics in Life Sciences*, **2017**, 1, 313-324	3.5	3
69	Draft Genome Sequences of Two subsp. Strains Isolated from Australian Hematology Patients with Bacteremia. *Genome Announcements*, **2017**, 5,		1
68	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	11.7	106
67	How can the cystic fibrosis respiratory microbiome influence our clinical decision-making?. *Current Opinion in Pulmonary Medicine*, **2017**, 23, 536-543	3	12
66	Intestinal Microbiota Composition in Sudden Infant Death Syndrome and Age-Matched Controls. *Journal of Pediatrics*, **2017**, 191, 63-68.e1	3.6	3
65	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	40	348
64	Inbred Mouse Populations Exhibit Intergenerational Changes in Intestinal Microbiota Composition and Function Following Introduction to a Facility. *Frontiers in Microbiology*, **2017**, 8, 608	5.7	15
63	Siblings of patients with Crohn's disease exhibit a biologically relevant dysbiosis in mucosal microbial metacommunities. *Gut*, **2016**, 65, 944-53	19.2	49
62	Culture-Independent Detection of Nontuberculous Mycobacteria in Clinical Respiratory Samples. *Journal of Clinical Microbiology*, **2016**, 54, 2395-8	9.7	7
61	Precision respiratory medicine and the microbiome. *Lancet Respiratory Medicine,the*, **2016**, 4, 73-82	35.1	24
60	Host-microbiome interactions in acute and chronic respiratory infections. *Cellular Microbiology*, **2016**, 18, 652-62	3.9	29
59	Respiratory microbiota resistance and resilience to pulmonary exacerbation and subsequent antimicrobial intervention. *ISME Journal*, **2016**, 10, 1081-91	11.9	68
58	The CF gastrointestinal microbiome: Structure and clinical impact. *Pediatric Pulmonology*, **2016**, 51, S35-S44	3.5	15
57	Macrolide Treatment Inhibits Pseudomonas aeruginosa Quorum Sensing in Non-Cystic Fibrosis Bronchiectasis. An Analysis from the Bronchiectasis and Low-Dose Erythromycin Study Trial. *Annals of the American Thoracic Society*, **2016**, 13, 1697-1703	4.7	20
56	Respiratory microbiota: addressing clinical questions, informing clinical practice. *Thorax*, **2015**, 70, 74-81	7.3	59
55	The human microbiome: opportunities and challenges for clinical care. *Internal Medicine Journal*, **2015**, 45, 889-98	1.6	6
54	Germs and joints: the contribution of the human microbiome to rheumatoid arthritis. *Nature Medicine*, **2015**, 21, 839-41	50.5	21

53	Matrix metalloproteinases vary with airway microbiota composition and lung function in non-cystic fibrosis bronchiectasis. *Annals of the American Thoracic Society*, **2015**, 12, 701-7	4.7	57
52	Is it time to rethink syphilis control?. *Clinical Infectious Diseases*, **2015**, 60, 325-6	11.6	4
51	Republished: Respiratory microbiota: addressing clinical questions, informing clinical practice. *Postgraduate Medical Journal*, **2015**, 91, 463-70	2	0
50	The microbiome of otitis media with effusion in Indigenous Australian children. *International Journal of Pediatric Otorhinolaryngology*, **2015**, 79, 1548-55	1.7	40
49	Predominant pathogen competition and core microbiota divergence in chronic airway infection. *ISME Journal*, **2015**, 9, 217-25	11.9	43
48	Sample storage conditions significantly influence faecal microbiome profiles. *Scientific Reports*, **2015**, 5, 16350	4.9	257
47	Ascites bacterial burden and immune cell profile are associated with poor clinical outcomes in the absence of overt infection. *PLoS ONE*, **2015**, 10, e0120642	3.7	25
46	Deriving accurate microbiota profiles from human samples with low bacterial content through post-sequencing processing of Illumina MiSeq data. *Microbiome*, **2015**, 3, 19	16.6	126
45	Implications of multiple freeze-thawing on respiratory samples for culture-independent analyses. *Journal of Cystic Fibrosis*, **2015**, 14, 464-7	4.1	20
44	Adult non-cystic fibrosis bronchiectasis is characterised by airway luminal Th17 pathway activation. *PLoS ONE*, **2015**, 10, e0119325	3.7	17
43	Three clinically distinct chronic pediatric airway infections share a common core microbiota. *Annals of the American Thoracic Society*, **2014**, 11, 1039-48	4.7	69
42	Time between collection and storage significantly influences bacterial sequence composition in sputum samples from cystic fibrosis respiratory infections. *Journal of Clinical Microbiology*, **2014**, 52, 3011-6	9.7	27
41	A novel microbiota stratification system predicts future exacerbations in bronchiectasis. *Annals of the American Thoracic Society*, **2014**, 11, 496-503	4.7	158
40	Combined systems approaches reveal highly plastic responses to antimicrobial peptide challenge in Escherichia coli. *PLoS Pathogens*, **2014**, 10, e1004104	7.6	30
39	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, 988-96	35.1	108
38	Potentially pathogenic airway bacteria and neutrophilic inflammation in treatment resistant severe asthma. *PLoS ONE*, **2014**, 9, e100645	3.7	203
37	Interpreting infective microbiota: the importance of an ecological perspective. *Trends in Microbiology*, **2013**, 21, 271-6	12.4	53
36	A relationship between growth behaviour and cystic fibrosis patient lung function identified in a metabolomic investigation. *Metabolomics*, **2013**, 9, 1262	4.7	17

35	Molecular detection of CF lung pathogens: current status and future potential. *Journal of Cystic Fibrosis*, **2013**, 12, 194-205	4.1	22
34	Clinical measures of disease in adult non-CF bronchiectasis correlate with airway microbiota composition. *Thorax*, **2013**, 68, 731-7	7.3	149
33	Staphylococcus aureus small-colony variants are independently associated with worse lung disease in children with cystic fibrosis. *Clinical Infectious Diseases*, **2013**, 57, 384-91	11.6	117
32	Reducing bias in bacterial community analysis of lower respiratory infections. *ISME Journal*, **2013**, 7, 697-706	11.9	66
31	Challenges and opportunities for faecal microbiota transplantation therapy. *Epidemiology and Infection*, **2013**, 141, 2235-42	4.3	9
30	Impact of antibiotic treatment for pulmonary exacerbations on bacterial diversity in cystic fibrosis. *Journal of Cystic Fibrosis*, **2013**, 12, 22-8	4.1	47
29	Complexity, temporal stability, and clinical correlates of airway bacterial community composition in primary ciliary dyskinesia. *Journal of Clinical Microbiology*, **2013**, 51, 4029-35	9.7	33
28	Ascitic microbiota composition is correlated with clinical severity in cirrhosis with portal hypertension. *PLoS ONE*, **2013**, 8, e74884	3.7	25
27	Exploring the parallel development of microbial systems in neonates with cystic fibrosis. *MBio*, **2012**, 3, e00408-12	7.8	4
26	Long-term cultivation-independent microbial diversity analysis demonstrates that bacterial communities infecting the adult cystic fibrosis lung show stability and resilience. *Thorax*, **2012**, 67, 867-73	7.3	111
25	Enhancing the utility of existing antibiotics by targeting bacterial behaviour?. *British Journal of Pharmacology*, **2012**, 165, 845-57	8.6	23
24	Does bacterial density in cystic fibrosis sputum increase prior to pulmonary exacerbation?. *Journal of Cystic Fibrosis*, **2011**, 10, 357-65	4.1	102
23	Novel concepts in evaluating antimicrobial therapy for bacterial lung infections in patients with cystic fibrosis. *Journal of Cystic Fibrosis*, **2011**, 10, 387-400	4.1	23
22	Characterization of bacterial community diversity in chronic rhinosinusitis infections using novel culture-independent techniques. *American Journal of Rhinology and Allergy*, **2011**, 25, e133-40	2.4	46
21	Partitioning core and satellite taxa from within cystic fibrosis lung bacterial communities. *ISME Journal*, **2011**, 5, 780-91	11.9	177
20	Analysis of the bacterial communities present in lungs of patients with cystic fibrosis from American and British centers. *Journal of Clinical Microbiology*, **2011**, 49, 281-91	9.7	54
19	Using bacterial biomarkers to identify early indicators of cystic fibrosis pulmonary exacerbation onset. *Expert Review of Molecular Diagnostics*, **2011**, 11, 197-206	3.8	15
18	The exclusion of dead bacterial cells is essential for accurate molecular analysis of clinical samples. *Clinical Microbiology and Infection*, **2010**, 16, 1656-8	9.5	53

17	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	2.7	17
16	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	9.7	43
15	Lung infections in cystic fibrosis: deriving clinical insight from microbial complexity. *Expert Review of Molecular Diagnostics*, **2010**, 10, 187-96	3.8	33
14	Comparing the microbiota of the cystic fibrosis lung and human gut. *Gut Microbes*, **2010**, 1, 85-93	8.8	40
13	Next-generation sequencing in the analysis of human microbiota: essential considerations for clinical application. *Molecular Diagnosis and Therapy*, **2010**, 14, 343-50	4.5	30
12	Revealing the dynamics of polymicrobial infections: implications for antibiotic therapy. *Trends in Microbiology*, **2010**, 18, 357-64	12.4	56
11	Characterisation of bacteria in ascites--reporting the potential of culture-independent, molecular analysis. *European Journal of Clinical Microbiology and Infectious Diseases*, **2010**, 29, 533-41	5.3	24
10	Studying bacterial infections through culture-independent approaches. *Journal of Medical Microbiology*, **2009**, 58, 1401-1418	3.2	60
9	Bacterial community diversity in cultures derived from healthy and inflamed ileal pouches after restorative proctocolectomy. *Inflammatory Bowel Diseases*, **2009**, 15, 1803-11	4.5	12
8	Chromosomal genes conferring tolerance to heavy metal (Ag) toxicity. *The Environmentalist*, **2009**, 29, 85-92		0
7	Studying bacteria in respiratory specimens by using conventional and molecular microbiological approaches. *BMC Pulmonary Medicine*, **2009**, 9, 14	3.5	48
6	Assessing the diagnostic importance of nonviable bacterial cells in respiratory infections. *Diagnostic Microbiology and Infectious Disease*, **2008**, 62, 133-41	2.9	67
5	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, 2601-4	9.7	115
4	Bacterial activity in cystic fibrosis lung infections. *Respiratory Research*, **2005**, 6, 49	7.3	69
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, 5176-83	9.7	251
2	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, 3548-58	9.7	168
1	The bronchiectasis microbiome82-98		