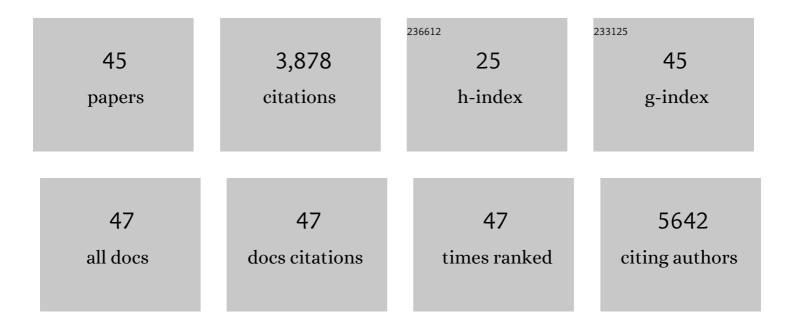
Jonathan Harris

List of Publications by Year in descending order

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#	Article	IF	CITATIONS
1	Effects of ivacaftor on systemic inflammation and the plasma proteome in people with CF and G551D. Journal of Cystic Fibrosis, 2022, 21, 950-958.	0.3	9
2	Temporal airway microbiome changes related to ventilator-associated pneumonia in children. European Respiratory Journal, 2021, 57, 2001829.	3.1	16
3	Modified PCR protocol to increase sensitivity for determination of bacterial community composition. Microbiome, 2021, 9, 90.	4.9	2
4	Influence of Acid Blockade on the Aerodigestive Tract Microbiome in Children With Cystic Fibrosis. Journal of Pediatric Gastroenterology and Nutrition, 2021, 72, 520-527.	0.9	3
5	Microbial and Biogeochemical Indicators of Methane in Groundwater Aquifers of the Denver Basin, Colorado. Environmental Science & Technology, 2021, 55, 292-303.	4.6	7
6	Bacterial Signatures of Paediatric Respiratory Disease: An Individual Participant Data Meta-Analysis. Frontiers in Microbiology, 2021, 12, 711134.	1.5	5
7	Changes in Airway Microbiome and Inflammation with Ivacaftor Treatment in Patients with Cystic Fibrosis and the G551D Mutation. Annals of the American Thoracic Society, 2020, 17, 212-220.	1.5	113
8	Nanodiagnostics to monitor biofilm oxygen metabolism for antibiotic susceptibility testing. Analyst, The, 2020, 145, 3996-4003.	1.7	5
9	Pulmonary Aptamer Signatures in Children's Interstitial and Diffuse Lung Disease. American Journal of Respiratory and Critical Care Medicine, 2019, 200, 1496-1504.	2.5	21
10	Luminescent Nanosensors for Ratiometric Monitoring of Three-Dimensional Oxygen Gradients in Laboratory and Clinical Pseudomonas aeruginosa Biofilms. Applied and Environmental Microbiology, 2019, 85, .	1.4	18
11	On the Use of Diversity Measures in Longitudinal Sequencing Studies of Microbial Communities. Frontiers in Microbiology, 2018, 9, 1037.	1.5	135
12	Succession of toxicity and microbiota in hydraulic fracturing flowback and produced water in the Denver–Julesburg Basin. Science of the Total Environment, 2018, 644, 183-192.	3.9	35
13	Airway microbiome and responses to corticosteroids in corticosteroid-resistant asthma patients treated with acid suppression medications. Journal of Allergy and Clinical Immunology, 2017, 140, 860-862.e1.	1.5	11
14	Longitudinal and Source-to-Tap New Orleans, LA, U.S.A. Drinking Water Microbiology. Environmental Science & Technology, 2017, 51, 4220-4229.	4.6	48
15	Airway microbiota across age and disease spectrum in cystic fibrosis. European Respiratory Journal, 2017, 50, 1700832.	3.1	193
16	Molecular analysis of endotracheal tube biofilms and tracheal aspirates in the pediatric intensive care unit. Advances in Pediatric Research, 2017, 4, .	2.0	1
17	Factors Influencing Bacterial Diversity and Community Composition in Municipal Drinking Waters in the Ohio River Basin, USA. PLoS ONE, 2016, 11, e0157966.	1.1	70
18	Assessing the Airway Microbiota in Cystic Fibrosis. Clinical Microbiology Newsletter, 2016, 38, 179-184.	0.4	2

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#	Article	IF	CITATIONS
19	Molecular Identification of Staphylococcus aureus in Airway Samples from Children with Cystic Fibrosis. PLoS ONE, 2016, 11, e0147643.	1.1	15
20	Esophageal Microbiome in Eosinophilic Esophagitis. PLoS ONE, 2015, 10, e0128346.	1.1	134
21	Molecular analysis of single room humidifier bacteriology. Water Research, 2015, 69, 318-327.	5.3	9
22	Assessment of Airway Microbiota and Inflammation in Cystic Fibrosis Using Multiple Sampling Methods. Annals of the American Thoracic Society, 2015, 12, 221-229.	1.5	128
23	High-Resolution Microbial Community Succession of Microbially Induced Concrete Corrosion in Working Sanitary Manholes. PLoS ONE, 2015, 10, e0116400.	1.1	30
24	Effect of monochloramine treatment on the microbial ecology of Legionella and associated bacterial populations in a hospital hot water system. Systematic and Applied Microbiology, 2015, 38, 198-205.	1.2	17
25	Novosphingobium and Its Potential Role in Chronic Obstructive Pulmonary Diseases: Insights from Microbiome Studies. PLoS ONE, 2014, 9, e111150.	1.1	23
26	Molecular analysis of point-of-use municipal drinking water microbiology. Water Research, 2014, 49, 225-235.	5.3	107
27	Explicet: graphical user interface software for metadata-driven management, analysis and visualization of microbiome data. Bioinformatics, 2013, 29, 3100-3101.	1.8	261
28	Microbiome of the Lower Airways Alters Corticosteroid Responsiveness in Asthma. Journal of Allergy and Clinical Immunology, 2013, 131, AB138.	1.5	1
29	Phylogenetic stratigraphy in the Guerrero Negro hypersaline microbial mat. ISME Journal, 2013, 7, 50-60.	4.4	198
30	The Effects of Airway Microbiome on Corticosteroid Responsiveness in Asthma. American Journal of Respiratory and Critical Care Medicine, 2013, 188, 1193-1201.	2.5	310
31	Bacterial identification and analytic challenges in clinical microbiome studies. Journal of Allergy and Clinical Immunology, 2012, 129, 441-442.	1.5	13
32	Application of Two-Part Statistics for Comparison of Sequence Variant Counts. PLoS ONE, 2011, 6, e20296.	1.1	55
33	A comparative molecular analysis of waterâ€filled limestone sinkholes in northâ€eastern Mexico. Environmental Microbiology, 2011, 13, 226-240.	1.8	23
34	Reliability of Quantitative Real-Time PCR for Bacterial Detection in Cystic Fibrosis Airway Specimens. PLoS ONE, 2010, 5, e15101.	1.1	71
35	Comparison of Normalization Methods for Construction of Large, Multiplex Amplicon Pools for Next-Generation Sequencing. Applied and Environmental Microbiology, 2010, 76, 3863-3868.	1.4	71
36	Measuring and improving respiratory outcomes in cystic fibrosis lung disease: Opportunities and challenges to therapy. Journal of Cystic Fibrosis, 2010, 9, 1-16.	0.3	93

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37	Novel Microbial Diversity Retrieved by Autonomous Robotic Exploration of the World's Deepest Vertical Phreatic Sinkhole. Astrobiology, 2010, 10, 201-213.	1.5	39
38	Diversity and Stratification of Archaea in a Hypersaline Microbial Mat. Applied and Environmental Microbiology, 2009, 75, 1801-1810.	1.4	102
39	Molecular identification of bacteria in bronchoalveolar lavage fluid from children with cystic fibrosis. Proceedings of the National Academy of Sciences of the United States of America, 2007, 104, 20529-20533.	3.3	339
40	Unexpected Diversity and Complexity of the Guerrero Negro Hypersaline Microbial Mat. Applied and Environmental Microbiology, 2006, 72, 3685-3695.	1.4	435
41	Phylogenetic diversity and ecology of environmental Archaea. Current Opinion in Microbiology, 2005, 8, 638-642.	2.3	120
42	New Perspective on Uncultured Bacterial Phylogenetic Division OP11. Applied and Environmental Microbiology, 2004, 70, 845-849.	1.4	190
43	The Genetic Core of the Universal Ancestor. Genome Research, 2003, 13, 407-412.	2.4	253
44	Evaluation and refinement of tmRNA structure using gene sequences from natural microbial communities. Rna, 2001, 7, 1310-1316.	1.6	11
45	Expanding the Known Diversity and Environmental Distribution of an Uncultured Phylogenetic Division of Bacteria, Applied and Environmental Microbiology, 2000, 66, 1617-1621.	1.4	135