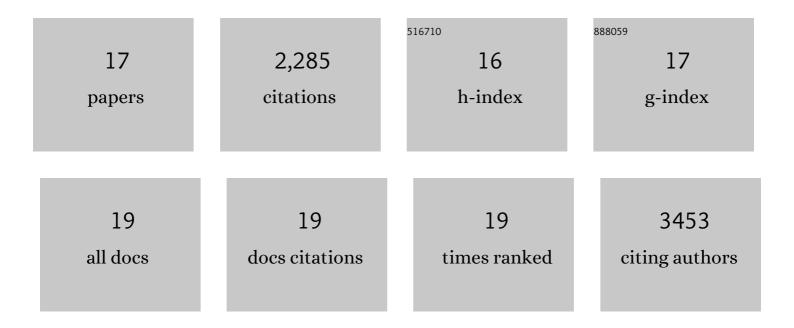
## Adam G Clooney

List of Publications by Year in descending order

Source: https://exaly.com/author-pdf/9518347/publications.pdf Version: 2024-02-01



#	Article	IF	CITATIONS
1	Ranking microbiome variance in inflammatory bowel disease: a large longitudinal intercontinental study. Gut, 2021, 70, 499-510.	12.1	127
2	Isolation and characterisation of î¦crAss002, a crAss-like phage from the human gut that infects Bacteroides xylanisolvens. Microbiome, 2021, 9, 89.	11.1	48
3	Stability of the human gut virome and effect of gluten-free diet. Cell Reports, 2021, 35, 109132.	6.4	34
4	Giant oversights in the human gut virome. Gut, 2020, 69, 1357-1358.	12.1	23
5	Isolation of a Novel Jumbo Bacteriophage Effective Against Klebsiella aerogenes. Frontiers in Medicine, 2020, 7, 67.	2.6	20
6	Oral Administration of Heat-Treated Lactobacilli Modifies the Murine Microbiome and Reduces Citrobacter Induced Colitis. Frontiers in Microbiology, 2020, 11, 69.	3.5	18
7	The Human Gut Virome Is Highly Diverse, Stable, and Individual Specific. Cell Host and Microbe, 2019, 26, 527-541.e5.	11.0	449
8	Short-term consumption of a high-fat diet increases host susceptibility to Listeria monocytogenes infection. Microbiome, 2019, 7, 7.	11.1	60
9	Evaluation of Phage Therapy in the Context of Enterococcus faecalis and Its Associated Diseases. Viruses, 2019, 11, 366.	3.3	55
10	Whole-Virome Analysis Sheds Light on Viral Dark Matter in Inflammatory Bowel Disease. Cell Host and Microbe, 2019, 26, 764-778.e5.	11.0	287
11	Choice of assembly software has a critical impact on virome characterisation. Microbiome, 2019, 7, 12.	11.1	106
12	Determinants of Reduced Genetic Capacity for Butyrate Synthesis by the Gut Microbiome in Crohn's Disease and Ulcerative Colitis. Journal of Crohn's and Colitis, 2018, 12, 204-216.	1.3	93
13	Biology and Taxonomy of crAss-like Bacteriophages, the Most Abundant Virus in the Human Gut. Cell Host and Microbe, 2018, 24, 653-664.e6.	11.0	233
14	A clinician's guide to microbiome analysis. Nature Reviews Gastroenterology and Hepatology, 2017, 14, 585-595.	17.8	124
15	A comparison of the gut microbiome between long-term users and non-users of proton pump inhibitors. Alimentary Pharmacology and Therapeutics, 2016, 43, 974-984.	3.7	126
16	16S rRNA gene sequencing of mock microbial populations- impact of DNA extraction method, primer choice and sequencing platform. BMC Microbiology, 2016, 16, 123.	3.3	241
17	Comparing Apples and Oranges?: Next Generation Sequencing and Its Impact on Microbiome Analysis. PLoS ONE, 2016, 11, e0148028.	2.5	234