

# Adam G Clooney

## List of Publications by Year in descending order

Source: <https://exaly.com/author-pdf/9518347/publications.pdf>

Version: 2024-02-01

17  
papers

2,285  
citations

516215

16  
h-index

887659

17  
g-index

19  
all docs

19  
docs citations

19  
times ranked

3453  
citing authors

| #  | ARTICLE   | IF  | CITATIONS |
|----|---|-----|-----------|
| 1  | Ranking microbiome variance in inflammatory bowel disease: a large longitudinal intercontinental study. <i>Gut</i> , 2021, 70, 499-510.   | 6.1 | 127       |
| 2  | Isolation and characterisation of $\phi$ crAss002, a crAss-like phage from the human gut that infects <i>Bacteroides xylanisolvens</i> . <i>Microbiome</i> , 2021, 9, 89.                   | 4.9 | 48        |
| 3  | Stability of the human gut virome and effect of gluten-free diet. <i>Cell Reports</i> , 2021, 35, 109132.   | 2.9 | 34        |
| 4  | Giant oversights in the human gut virome. <i>Gut</i> , 2020, 69, 1357-1358.   | 6.1 | 23        |
| 5  | Isolation of a Novel Jumbo Bacteriophage Effective Against <i>Klebsiella aerogenes</i> . <i>Frontiers in Medicine</i> , 2020, 7, 67.  | 1.2 | 20        |
| 6  | Oral Administration of Heat-Treated Lactobacilli Modifies the Murine Microbiome and Reduces Citrobacter Induced Colitis. <i>Frontiers in Microbiology</i> , 2020, 11, 69.                   | 1.5 | 18        |
| 7  | The Human Gut Virome Is Highly Diverse, Stable, and Individual Specific. <i>Cell Host and Microbe</i> , 2019, 26, 527-541.e5.   | 5.1 | 449       |
| 8  | Short-term consumption of a high-fat diet increases host susceptibility to <i>Listeria monocytogenes</i> infection. <i>Microbiome</i> , 2019, 7, 7.   | 4.9 | 60        |
| 9  | Evaluation of Phage Therapy in the Context of <i>Enterococcus faecalis</i> and Its Associated Diseases. <i>Viruses</i> , 2019, 11, 366.   | 1.5 | 55        |
| 10 | Whole-Virome Analysis Sheds Light on Viral Dark Matter in Inflammatory Bowel Disease. <i>Cell Host and Microbe</i> , 2019, 26, 764-778.e5.  | 5.1 | 287       |
| 11 | Choice of assembly software has a critical impact on virome characterisation. <i>Microbiome</i> , 2019, 7, 12.  | 4.9 | 106       |
| 12 | Determinants of Reduced Genetic Capacity for Butyrate Synthesis by the Gut Microbiome in Crohn's Disease and Ulcerative Colitis. <i>Journal of Crohn's and Colitis</i> , 2018, 12, 204-216. | 0.6 | 93        |
| 13 | Biology and Taxonomy of crAss-like Bacteriophages, the Most Abundant Virus in the Human Gut. <i>Cell Host and Microbe</i> , 2018, 24, 653-664.e6.   | 5.1 | 233       |
| 14 | A clinician's guide to microbiome analysis. <i>Nature Reviews Gastroenterology and Hepatology</i> , 2017, 14, 585-595.  | 8.2 | 124       |
| 15 | A comparison of the gut microbiome between long-term users and non-users of proton pump inhibitors. <i>Alimentary Pharmacology and Therapeutics</i> , 2016, 43, 974-984.                    | 1.9 | 126       |
| 16 | 16S rRNA gene sequencing of mock microbial populations- impact of DNA extraction method, primer choice and sequencing platform. <i>BMC Microbiology</i> , 2016, 16, 123.                    | 1.3 | 241       |
| 17 | Comparing Apples and Oranges?: Next Generation Sequencing and Its Impact on Microbiome Analysis. <i>PLoS ONE</i> , 2016, 11, e0148028.  | 1.1 | 234       |