

# Erik L Clarke

## List of Publications by Year in descending order

Source: <https://exaly.com/author-pdf/3368482/publications.pdf>

Version: 2024-02-01

13  
papers

1,016  
citations

933264

10  
h-index

1199470

12  
g-index

14  
all docs

14  
docs citations

14  
times ranked

2081  
citing authors

#	ARTICLE	IF	CITATIONS
1	Respiratory Microbiome Disruption and Risk for Ventilator-Associated Lower Respiratory Tract Infection. <i>Clinical Infectious Diseases</i> , 2021, , .	2.9	1
2	Upper Respiratory Dysbiosis with a Facultative-dominated Ecotype in Advanced Lung Disease and Dynamic Change after Lung Transplant. <i>Annals of the American Thoracic Society</i> , 2019, 16, 1383-1391.	1.5	16
3	Sunbeam: an extensible pipeline for analyzing metagenomic sequencing experiments. <i>Microbiome</i> , 2019, 7, 46.	4.9	134
4	Bidirectional transfer of Anelloviridae lineages between graft and host during lung transplantation. <i>American Journal of Transplantation</i> , 2019, 19, 1086-1097.	2.6	30
5	Molecular analysis of bacterial contamination on stethoscopes in an intensive care unit. <i>Infection Control and Hospital Epidemiology</i> , 2019, 40, 171-177.	1.0	25
6	Microbial Lineages in Sarcoidosis. A Metagenomic Analysis Tailored for Lowâ€ Microbial Content Samples. <i>American Journal of Respiratory and Critical Care Medicine</i> , 2018, 197, 225-234.	2.5	59
7	614. Modeling Changes in Gastrointestinal and Respiratory Tract Bacterial Community Diversity Attributable to Common Antibiotic Exposures During Long-Term Acute Care. <i>Open Forum Infectious Diseases</i> , 2018, 5, S224-S224.	0.4	0
8	T cell dynamics and response of the microbiota after gene therapy to treat X-linked severe combined immunodeficiency. <i>Genome Medicine</i> , 2018, 10, 70.	3.6	28
9	Lack of detection of a human placenta microbiome in samples from preterm and term deliveries. <i>Microbiome</i> , 2018, 6, 196.	4.9	221
10	Evaluation of a therapy for Idiopathic Chronic Enterocolitis in rhesus macaques (<i>Macaca</i> Tj ETQq0 0 0 rgBT /Overlock 10 Tf 50 382 T	0.9	5
11	swga: a primer design toolkit for selective whole genome amplification. <i>Bioinformatics</i> , 2017, 33, 2071-2077.	1.8	27
12	Optimizing methods and dodging pitfalls in microbiome research. <i>Microbiome</i> , 2017, 5, 52.	4.9	420
13	Modeling altered T-cell development with induced pluripotent stem cells from patients with RAG1-dependent immune deficiencies. <i>Blood</i> , 2016, 128, 783-793.	0.6	45