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 1,016 10 12 papers citations h-index g-index

14 14 2081 all docs docs citations times ranked citing authors

#	Article	IF	CITATIONS
1	Optimizing methods and dodging pitfalls in microbiome research. Microbiome, 2017, 5, 52.	4.9	420
2	Lack of detection of a human placenta microbiome in samples from preterm and term deliveries. Microbiome, 2018, 6, 196.	4.9	221
3	Sunbeam: an extensible pipeline for analyzing metagenomic sequencing experiments. Microbiome, 2019, 7, 46.	4.9	134
4	Microbial Lineages in Sarcoidosis. A Metagenomic Analysis Tailored for Low–Microbial Content Samples. American Journal of Respiratory and Critical Care Medicine, 2018, 197, 225-234.	2.5	59
5	Modeling altered T-cell development with induced pluripotent stem cells from patients with RAG1-dependent immune deficiencies. Blood, 2016, 128, 783-793.	0.6	45
6	Bidirectional transfer of Anelloviridae lineages between graft and host during lung transplantation. American Journal of Transplantation, 2019, 19, 1086-1097.	2.6	30
7	T cell dynamics and response of the microbiota after gene therapy to treat X-linked severe combined immunodeficiency. Genome Medicine, 2018, 10, 70.	3.6	28
8	swga: a primer design toolkit for selective whole genome amplification. Bioinformatics, 2017, 33, 2071-2077.	1.8	27
9	Molecular analysis of bacterial contamination on stethoscopes in an intensive care unit. Infection Control and Hospital Epidemiology, 2019, 40, 171-177.	1.0	25
10	Upper Respiratory Dysbiosis with a Facultative-dominated Ecotype in Advanced Lung Disease and Dynamic Change after Lung Transplant. Annals of the American Thoracic Society, 2019, 16, 1383-1391.	1.5	16
11	Evaluation of a therapy for Idiopathic Chronic Enterocolitis in rhesus macaques (<i>Macaca) Tj ETQq1 1 0.78431</i>	4 rgBT /Ov	verjock 10 Tf
12	Respiratory Microbiome Disruption and Risk for Ventilator-Associated Lower Respiratory Tract Infection. Clinical Infectious Diseases, 2021, , .	2.9	1
13	614. Modeling Changes in Gastrointestinal and Respiratory Tract Bacterial Community Diversity Attributable to Common Antibiotic Exposures During Long-Term Acute Care. Open Forum Infectious Diseases, 2018, 5, S224-S224.	0.4	O