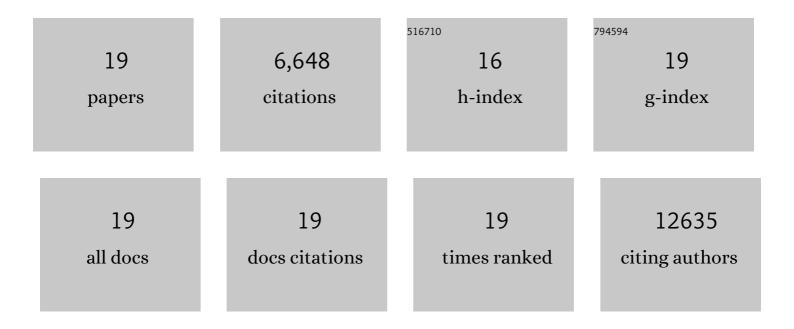
Kelvin Li

List of Publications by Year in descending order

Source: https://exaly.com/author-pdf/10756393/publications.pdf Version: 2024-02-01



KEIMINTI

#	Article	IF	CITATIONS
1	Dynamics of hepatic steatosis resolution and changes in gut microbiome with weight loss in nonalcoholic fatty liver disease. Obesity Science and Practice, 2021, 7, 217-225.	1.9	5
2	"Longitudinal Fecal Microbiome Study of Total Body Irradiated Mice Treated With Radiation Mitigators Identifies Bacterial Associations With Survival― Frontiers in Cellular and Infection Microbiology, 2021, 11, 715396.	3.9	5
3	Respiratory Tract Dysbiosis Is Associated with Worse Outcomes in Mechanically Ventilated Patients. American Journal of Respiratory and Critical Care Medicine, 2020, 202, 1666-1677.	5.6	49
4	Respiratory Microbiome Profiling for Etiologic Diagnosis of Pneumonia in Mechanically Ventilated Patients. Frontiers in Microbiology, 2018, 9, 1413.	3.5	61
5	Interactions between HIV-1 Gag and Viral RNA Genome Enhance Virion Assembly. Journal of Virology, 2017, 91, .	3.4	28
6	Community differentiation of the cutaneous microbiota in psoriasis. Microbiome, 2013, 1, 31.	11.1	353
7	Analyses of the Stability and Core Taxonomic Memberships of the Human Microbiome. PLoS ONE, 2013, 8, e63139.	2.5	184
8	Antibiotics in early life alter the murine colonic microbiome and adiposity. Nature, 2012, 488, 621-626.	27.8	1,358
9	Analyses of the Microbial Diversity across the Human Microbiome. PLoS ONE, 2012, 7, e32118.	2.5	203
10	VariantClassifier: A hierarchical variant classifier for annotated genomes. BMC Research Notes, 2010, 3, 191.	1.4	16
11	ANDES: Statistical tools for the ANalyses of DEep Sequencing. BMC Research Notes, 2010, 3, 199.	1.4	21
12	Infidelity of SARS-CoV Nsp14-Exonuclease Mutant Virus Replication Is Revealed by Complete Genome Sequencing. PLoS Pathogens, 2010, 6, e1000896.	4.7	381
13	The minimum information about a genome sequence (MICS) specification. Nature Biotechnology, 2008, 26, 541-547.	17.5	1,069
14	Novel computational methods for increasing PCR primer design effectiveness in directed sequencing. BMC Bioinformatics, 2008, 9, 191.	2.6	30
15	Genetic Variation in an Individual Human Exome. PLoS Genetics, 2008, 4, e1000160.	3.5	248
16	Aggressive assembly of pyrosequencing reads with mates. Bioinformatics, 2008, 24, 2818-2824.	4.1	498
17	The Sorcerer II Global Ocean Sampling Expedition: Northwest Atlantic through Eastern Tropical Pacific. PLoS Biology, 2007, 5, e77.	5.6	1,757
18	A Sanger/pyrosequencing hybrid approach for the generation of high-quality draft assemblies of marine microbial genomes. Proceedings of the National Academy of Sciences of the United States of America, 2006, 103, 11240-11245.	7.1	243

#	Article	IF	CITATIONS
19	Sequence survey of receptor tyrosine kinases reveals mutations in glioblastomas. Proceedings of the National Academy of Sciences of the United States of America, 2005, 102, 14344-14349.	7.1	139