

Kelvin Li

List of Publications by Year in descending order

Source: <https://exaly.com/author-pdf/10756393/publications.pdf>

Version: 2024-02-01

19
papers

6,648
citations

516710

16
h-index

794594

19
g-index

19
all docs

19
docs citations

19
times ranked

12635
citing authors

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