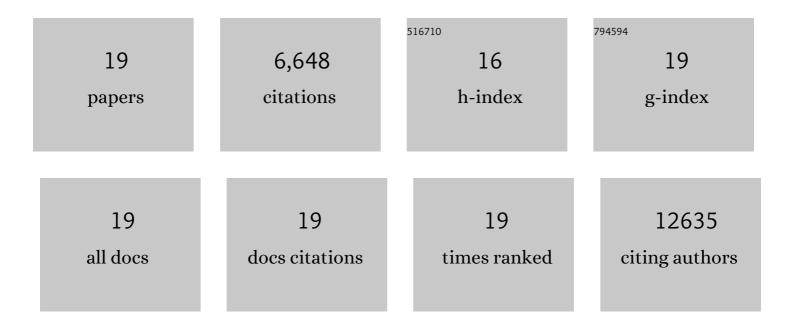
Kelvin Li

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 |