

Wen-Han Yu

List of Publications by Year in descending order

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Version: 2024-02-01

14
papers

4,155
citations

686830

13
h-index

1058022

14
g-index

14
all docs

14
docs citations

14
times ranked

7140
citing authors

#	ARTICLE	IF	CITATIONS
1	The Human Oral Microbiome. <i>Journal of Bacteriology</i> , 2010, 192, 5002-5017.	1.0	2,536
2	The Human Oral Microbiome Database: a web accessible resource for investigating oral microbe taxonomic and genomic information. <i>Database: the Journal of Biological Databases and Curation</i> , 2010, 2010, baq013-baq013.	1.4	874
3	Antibody and TLR7 agonist delay viral rebound in SHIV-infected monkeys. <i>Nature</i> , 2018, 563, 360-364.	13.7	246
4	A Role for Fc Function in Therapeutic Monoclonal Antibody-Mediated Protection against Ebola Virus. <i>Cell Host and Microbe</i> , 2018, 24, 221-233.e5.	5.1	182
5	<i>Pyramidobacter piscolens</i> gen. nov., sp. nov., a member of the phylum 'Synergistetes' isolated from the human oral cavity. <i>International Journal of Systematic and Evolutionary Microbiology</i> , 2009, 59, 972-980.	0.8	108
6	Selective induction of antibody effector functional responses using MF59-adjuvanted vaccination. <i>Journal of Clinical Investigation</i> , 2019, 130, 662-672.	3.9	50
7	Comprehensive Transcriptome Analysis of the Periodontopathogenic Bacterium <i>Porphyromonas gingivalis</i> W83. <i>Journal of Bacteriology</i> , 2012, 194, 100-114.	1.0	31
8	Prediction of VRC01 neutralization sensitivity by HIV-1 gp160 sequence features. <i>PLoS Computational Biology</i> , 2019, 15, e1006952.	1.5	25
9	Predicting the broadly neutralizing antibody susceptibility of the HIV reservoir. <i>JCI Insight</i> , 2019, 4, .	2.3	25
10	Exploiting glycan topography for computational design of Env glycoprotein antigenicity. <i>PLoS Computational Biology</i> , 2018, 14, e1006093.	1.5	19
11	HIV Antibody Fc N-Linked Glycosylation Is Associated with Viral Rebound. <i>Cell Reports</i> , 2020, 33, 108502.	2.9	19
12	Strand-specific transcriptome profiling with directly labeled RNA on genomic tiling microarrays. <i>BMC Molecular Biology</i> , 2011, 12, 3.	3.0	15
13	A novel retrieval system for nearly complete microbial genomic fragments from soil samples. <i>Journal of Microbiological Methods</i> , 2008, 72, 197-205.	0.7	14
14	A hidden Markov support vector machine framework incorporating profile geometry learning for identifying microbial RNA in tiling array data. <i>Bioinformatics</i> , 2010, 26, 1423-1430.	1.8	11