Wen-Han Yu

List of Publications by Year
in descending order

14	4,155	687363	1058476
papers	**citations**	13	14
		h-index	**g-index**

14	14	14	7140
all docs	**docs citations**	**times ranked**	**citing authors**

WEN-HAN YU

#	Article	IF	Citations
1	The Human Oral Microbiome. Journal of Bacteriology, 2010, 192, 5002-5017.	2.2	2,536
2	The Human Oral Microbiome Database: a web accessible resource for investigating oral microbe taxonomic and genomic information. Database: the Journal of Biological Databases and Curation, 2010, 2010, baq013-baq013.	3.0	874
3	Antibody and TLR7 agonist delay viral rebound in SHIV-infected monkeys. Nature, 2018, 563, 360-364.	27.8	246
4	A Role for Fc Function in Therapeutic Monoclonal Antibody-Mediated Protection against Ebola Virus. Cell Host and Microbe, 2018, 24, 221-233.e5.	11.0	182
5	Pyramidobacter piscolens gen. nov., sp. nov., a member of the phylum 'Synergistetes' isolated from the human oral cavity. International Journal of Systematic and Evolutionary Microbiology, 2009, 59, 972-980.	1.7	108
6	Selective induction of antibody effector functional responses using MF59-adjuvanted vaccination. Journal of Clinical Investigation, 2019, 130, 662-672.	8.2	50
7	Comprehensive Transcriptome Analysis of the Periodontopathogenic Bacterium Porphyromonas gingivalis W83. Journal of Bacteriology, 2012, 194, 100-114.	2.2	31
8	Prediction of VRC01 neutralization sensitivity by HIV-1 gp160 sequence features. PLoS Computational Biology, 2019, 15, e1006952.	3.2	25
9	Predicting the broadly neutralizing antibody susceptibility of the HIV reservoir. JCI Insight, 2019, 4, .	5.0	25
10	Exploiting glycan topography for computational design of Env glycoprotein antigenicity. PLoS Computational Biology, 2018, 14, e1006093.	3.2	19
11	HIV Antibody Fc N-Linked Glycosylation Is Associated with Viral Rebound. Cell Reports, 2020, 33, 108502.	6.4	19
12	Strand-specific transcriptome profiling with directly labeled RNA on genomic tiling microarrays. BMC Molecular Biology, 2011, 12, 3.	3.0	15
13	A novel retrieval system for nearly complete microbial genomic fragments from soil samples. Journal of Microbiological Methods, 2008, 72, 197-205.	1.6	14
14	A hidden Markov support vector machine framework incorporating profile geometry learning for identifying microbial RNA in tiling array data. Bioinformatics, 2010, 26, 1423-1430.	4.1	11