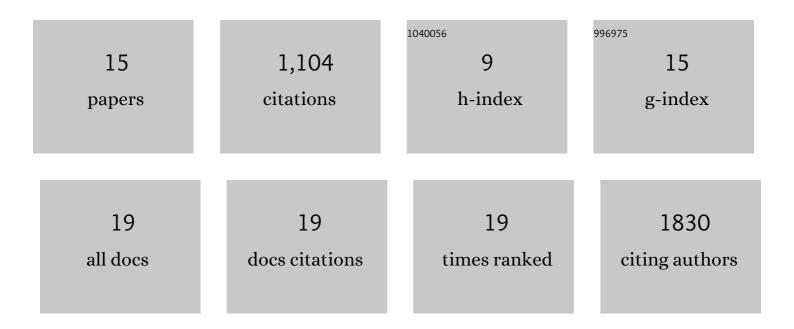
## Jacquelyn S Meisel

List of Publications by Year in descending order

Source: https://exaly.com/author-pdf/948247/publications.pdf Version: 2024-02-01



IACOLIELYN S MEISEL

#	Article	IF	CITATIONS
1	Binnacle: Using Scaffolds to Improve the Contiguity and Quality of Metagenomic Bins. Frontiers in Microbiology, 2021, 12, 638561.	3.5	2
2	Three transmission events of Vibrio cholerae O1 into Lusaka, Zambia. BMC Infectious Diseases, 2021, 21, 570.	2.9	5
3	Human macrophage response to microbial supernatants from diabetic foot ulcers. Wound Repair and Regeneration, 2019, 27, 598-608.	3.0	9
4	Strain- and Species-Level Variation in the Microbiome of Diabetic Wounds Is Associated with Clinical Outcomes and Therapeutic Efficacy. Cell Host and Microbe, 2019, 25, 641-655.e5.	11.0	192
5	Current State of and Future Opportunities for Prediction in Microbiome Research: Report from the Mid-Atlantic Microbiome Meet-up in Baltimore on 9 January 2019. MSystems, 2019, 4, .	3.8	6
6	Embracing Ambiguity in the Taxonomic Classification of Microbiome Sequencing Data. Frontiers in Genetics, 2019, 10, 1022.	2.3	6
7	HmmUFOtu: An HMM and phylogenetic placement based ultra-fast taxonomic assignment and OTU picking tool for microbiome amplicon sequencing studies. Genome Biology, 2018, 19, 82.	8.8	32
8	Antiseptic Agents Elicit Short-Term, Personalized, and Body Site–Specific Shifts in Resident Skin Bacterial Communities. Journal of Investigative Dermatology, 2018, 138, 2234-2243.	0.7	52
9	Commensal microbiota modulate gene expression in the skin. Microbiome, 2018, 6, 20.	11.1	147
10	The Human Microbiome. , 2017, , 63-77.		4
11	Topical Antimicrobial Treatments Can Elicit Shifts to Resident Skin Bacterial Communities and Reduce Colonization by Staphylococcus aureus Competitors. Antimicrobial Agents and Chemotherapy, 2017, 61,	3.2	48
12	Cutaneous Leishmaniasis Induces a Transmissible Dysbiotic Skin Microbiota that Promotes Skin Inflammation. Cell Host and Microbe, 2017, 22, 13-24.e4.	11.0	82
13	Evolutionary and functional implications of hypervariable loci within the skin virome. PeerJ, 2017, 5, e2959.	2.0	28
14	Skin Microbiome Surveys Are Strongly Influenced by Experimental Design. Journal of Investigative Dermatology, 2016, 136, 947-956.	0.7	249
15	The Human Skin Double-Stranded DNA Virome: Topographical and Temporal Diversity, Genetic Enrichment, and Dynamic Associations with the Host Microbiome. MBio, 2015, 6, e01578-15.	4.1	232