

# Michael Schmid

## List of Publications by Year in descending order

Source: <https://exaly.com/author-pdf/6441165/publications.pdf>

Version: 2024-02-01

14  
papers

694  
citations

840776  
11  
h-index

1125743  
13  
g-index

18  
all docs

18  
docs citations

18  
times ranked

932  
citing authors

#	ARTICLE	IF	CITATIONS
1	Evaluation of Oxford Nanopore MinION RNA-Seq Performance for Human Primary Cells. International Journal of Molecular Sciences, 2021, 22, 6317.	4.1	8
2	Phospho-RNA sequencing with circAID-p-seq. Nucleic Acids Research, 2021, , .	14.5	0
3	Global transcriptome analysis reveals relevant effects at environmental concentrations of cypermethrin in honey bees ( <i>Apis mellifera</i> ). Environmental Pollution, 2020, 259, 113715.	7.5	15
4	The neonicotinoid thiacloprid causes transcriptional alteration of genes associated with mitochondria at environmental concentrations in honey bees. Environmental Pollution, 2020, 266, 115297.	7.5	24
5	<i>Pseudomonas orientalis</i> F9 Pyoverdine, Safracin, and Phenazine Mutants Remain Effective Antagonists against <i>Erwinia amylovora</i> in Apple Flowers. Applied and Environmental Microbiology, 2020, 86, .	3.1	18
6	Long-read based de novo assembly of low-complexity metagenome samples results in finished genomes and reveals insights into strain diversity and an active phage system. BMC Microbiology, 2019, 19, 143.	3.3	104
7	Pushing the limits of de novo genome assembly for complex prokaryotic genomes harboring very long, near identical repeats. Nucleic Acids Research, 2018, 46, 8953-8965.	14.5	104
8	Comparative Genomics of Completely Sequenced <i>Lactobacillus helveticus</i> Genomes Provides Insights into Strain-Specific Genes and Resolves Metagenomics Data Down to the Strain Level. Frontiers in Microbiology, 2018, 9, 63.	3.5	73
9	<i>Pseudomonas orientalis</i> F9: A Potent Antagonist against Phytopathogens with Phytotoxic Effect in the Apple Flower. Frontiers in Microbiology, 2018, 9, 145.	3.5	34
10	Biofilm Formation Potential of Heat-Resistant <i>Escherichia coli</i> Dairy Isolates and the Complete Genome of Multidrug-Resistant, Heat-Resistant Strain FAM21845. Applied and Environmental Microbiology, 2017, 83, .	3.1	29
11	An integrative strategy to identify the entire protein coding potential of prokaryotic genomes by proteogenomics. Genome Research, 2017, 27, 2083-2095.	5.5	112
12	Competition assays and physiological experiments of soil and phyllosphere yeasts identify <i>Candida subhashii</i> as a novel antagonist of filamentous fungi. BMC Microbiology, 2017, 17, 4.	3.3	77
13	Complete genome sequence of <i>Pseudomonas citronellolis</i> P3B5, a candidate for microbial phyllo-remediation of hydrocarbon-contaminated sites. Standards in Genomic Sciences, 2016, 11, 75.	1.5	49
14	Short communication: Heat-resistant <i>Escherichia coli</i> as potential persistent reservoir of extended-spectrum $\beta$ -lactamases and Shiga toxin-encoding phages in dairy. Journal of Dairy Science, 2016, 99, 8622-8632.	3.4	30