

# Bettina MÃ¼ller

## List of Publications by Year in descending order

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15  
papers

419  
citations

1163117

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h-index

1281871

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g-index

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17  
docs citations

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times ranked

530  
citing authors

#	ARTICLE	IF	CITATIONS
1	Near Chromosome-Level Genome Assembly and Annotation of <i>Rhodotorula babjevae</i> Strains Reveals High Intraspecific Divergence. <i>Journal of Fungi</i> (Basel, Switzerland), 2022, 8, 323.	3.5	1
2	Metagenomics workflow for hybrid assembly, differential coverage binning, metatranscriptomics and pathway analysis (MUFFIN). <i>PLoS Computational Biology</i> , 2021, 17, e1008716.	3.2	18
3	Fecal Short-Chain Fatty Acid Ratios as Related to Gastrointestinal and Depressive Symptoms in Young Adults. <i>Psychosomatic Medicine</i> , 2021, 83, 693-699.	2.0	37
4	Chromosome-level genome assembly and transcriptome-based annotation of the oleaginous yeast <i>Rhodotorula toruloides</i> CBS 14. <i>Genomics</i> , 2021, 113, 4022-4027.	2.9	9
5	Impact of time and temperature on gut microbiota and SCFA composition in stool samples. <i>PLoS ONE</i> , 2020, 15, e0236944.	2.5	12
6	Abundance Tracking by Long-Read Nanopore Sequencing of Complex Microbial Communities in Samples from 20 Different Biogas/Wastewater Plants. <i>Applied Sciences</i> (Switzerland), 2020, 10, 7518.	2.5	15
7	Impact of time and temperature on gut microbiota and SCFA composition in stool samples. , 2020, 15, e0236944.		0
8	Impact of time and temperature on gut microbiota and SCFA composition in stool samples. , 2020, 15, e0236944.		0
9	Impact of time and temperature on gut microbiota and SCFA composition in stool samples. , 2020, 15, e0236944.		0
10	Impact of time and temperature on gut microbiota and SCFA composition in stool samples. , 2020, 15, e0236944.		0
11	Genome-Guided Analysis of <i>Clostridium ultunense</i> and Comparative Genomics Reveal Different Strategies for Acetate Oxidation and Energy Conservation in Syntrophic Acetate-Oxidising Bacteria. <i>Genes</i> , 2018, 9, 225.	2.4	27
12	Genome-Guided Analysis and Whole Transcriptome Profiling of the Mesophilic Syntrophic Acetate Oxidising Bacterium <i>Syntrophaceticus schinkii</i> . <i>PLoS ONE</i> , 2016, 11, e0166520.	2.5	53
13	Genome-Guided Analysis of Physiological Capacities of <i>Tepidanaerobacter acetatoxydans</i> Provides Insights into Environmental Adaptations and Syntrophic Acetate Oxidation. <i>PLoS ONE</i> , 2015, 10, e0121237.	2.5	28
14	Syntrophic acetate oxidation in industrial CSTR biogas digesters. <i>Journal of Biotechnology</i> , 2014, 171, 39-44.	3.8	92
15	First insights into the syntrophic acetate-oxidizing bacteria – a genetic study. <i>MicrobiologyOpen</i> , 2013, 2, 35-53.	3.0	126