

# Ashley Shade

## List of Publications by Year in descending order

Source: <https://exaly.com/author-pdf/5836274/publications.pdf>

Version: 2024-02-01

71  
papers

10,294  
citations

87723

38  
h-index

79541

73  
g-index

104  
all docs

104  
docs citations

104  
times ranked

13844  
citing authors

#	ARTICLE	IF	CITATIONS
1	A communal catalogue reveals Earth's multiscale microbial diversity. <i>Nature</i> , 2017, 551, 457-463.	13.7	1,942
2	Fundamentals of Microbial Community Resistance and Resilience. <i>Frontiers in Microbiology</i> , 2012, 3, 417.	1.5	1,131
3	Beyond the Venn diagram: the hunt for a core microbiome. <i>Environmental Microbiology</i> , 2012, 14, 4-12.	1.8	940
4	Conditionally Rare Taxa Disproportionately Contribute to Temporal Changes in Microbial Diversity. <i>MBio</i> , 2014, 5, e01371-14.	1.8	549
5	Diversity is the question, not the answer. <i>ISME Journal</i> , 2017, 11, 1-6.	4.4	358
6	Controls on soil microbial community stability under climate change. <i>Frontiers in Microbiology</i> , 2013, 4, 265.	1.5	353
7	Ecological patterns of seed microbiome diversity, transmission, and assembly. <i>Current Opinion in Microbiology</i> , 2017, 37, 15-22.	2.3	331
8	A meta-analysis of changes in bacterial and archaeal communities with time. <i>ISME Journal</i> , 2013, 7, 1493-1506.	4.4	322
9	Meeting Report: The Terabase Metagenomics Workshop and the Vision of an Earth Microbiome Project. <i>Standards in Genomic Sciences</i> , 2010, 3, 243-248.	1.5	228
10	Genome-wide selective sweeps and gene-specific sweeps in natural bacterial populations. <i>ISME Journal</i> , 2016, 10, 1589-1601.	4.4	226
11	Unexpected Diversity during Community Succession in the Apple Flower Microbiome. <i>MBio</i> , 2013, 4, .	1.8	210
12	Lake microbial communities are resilient after a whole-ecosystem disturbance. <i>ISME Journal</i> , 2012, 6, 2153-2167.	4.4	198
13	Assembly and seasonality of core phyllosphere microbiota on perennial biofuel crops. <i>Nature Communications</i> , 2019, 10, 4135.	5.8	182
14	Culturing captures members of the soil rare biosphere. <i>Environmental Microbiology</i> , 2012, 14, 2247-2252.	1.8	174
15	The under-ice microbiome of seasonally frozen lakes. <i>Limnology and Oceanography</i> , 2013, 58, 1998-2012.	1.6	173
16	The microbial ecology of flowers: an emerging frontier in phyllosphere research. <i>Botany</i> , 2014, 92, 253-266.	0.5	173
17	Interannual dynamics and phenology of bacterial communities in a eutrophic lake. <i>Limnology and Oceanography</i> , 2007, 52, 487-494.	1.6	167
18	Abundance-occupancy distributions to prioritize plant core microbiome membership. <i>Current Opinion in Microbiology</i> , 2019, 49, 50-58.	2.3	136

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19	Resistance, resilience and recovery: aquatic bacterial dynamics after water column disturbance. <i>Environmental Microbiology</i> , 2011, 13, 2752-2767.	1.8	127
20	Temporal patterns of rarity provide a more complete view of microbial diversity. <i>Trends in Microbiology</i> , 2015, 23, 335-340.	3.5	124
21	The influence of habitat heterogeneity on freshwater bacterial community composition and dynamics. <i>Environmental Microbiology</i> , 2008, 10, 1057-1067.	1.8	120
22	Macroecology to Unite All Life, Large and Small. <i>Trends in Ecology and Evolution</i> , 2018, 33, 731-744.	4.2	118
23	Divergent extremes but convergent recovery of bacterial and archaeal soil communities to an ongoing subterranean coal mine fire. <i>ISME Journal</i> , 2017, 11, 1447-1459.	4.4	108
24	Characterizing microbial communities through space and time. <i>Current Opinion in Biotechnology</i> , 2012, 23, 431-436.	3.3	98
25	Assembly of seed-associated microbial communities within and across successive plant generations. <i>Plant and Soil</i> , 2018, 422, 67-79.	1.8	91
26	Trait-based community assembly and succession of the infant gut microbiome. <i>Nature Communications</i> , 2019, 10, 512.	5.8	88
27	Bridging the gap between micro - and macro-scale perspectives on the role of microbial communities in global change ecology. <i>Plant and Soil</i> , 2006, 289, 59-70.	1.8	86
28	Bacterial Community Composition and Dynamics Spanning Five Years in Freshwater Bog Lakes. <i>MSphere</i> , 2017, 2, .	1.3	84
29	Effects of Short-Term Warming and Altered Precipitation on Soil Microbial Communities in Alpine Grassland of the Tibetan Plateau. <i>Frontiers in Microbiology</i> , 2016, 7, 1032.	1.5	81
30	Persistent microbiome members in the common bean rhizosphere: an integrated analysis of space, time, and plant genotype. <i>ISME Journal</i> , 2021, 15, 2708-2722.	4.4	76
31	Typhoons initiate predictable change in aquatic bacterial communities. <i>Limnology and Oceanography</i> , 2008, 53, 1319-1326.	1.6	73
32	Our microbial selves: what ecology can teach us. <i>EMBO Reports</i> , 2011, 12, 775-784.	2.0	71
33	Microbial metabolites in the marine carbon cycle. <i>Nature Microbiology</i> , 2022, 7, 508-523.	5.9	71
34	A global survey of arsenic-related genes in soil microbiomes. <i>BMC Biology</i> , 2019, 17, 45.	1.7	70
35	A Synthetic Community System for Probing Microbial Interactions Driven by Exometabolites. <i>MSystems</i> , 2017, 2, .	1.7	61
36	Comparison of Primer Sets for Use in Automated Ribosomal Intergenic Spacer Analysis of Aquatic Bacterial Communities: an Ecological Perspective. <i>Applied and Environmental Microbiology</i> , 2007, 73, 659-662.	1.4	56

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37	Trait-based patterns of microbial dynamics in dormancy potential and heterotrophic strategy: case studies of resource-based and post-pess succession. <i>ISME Journal</i> , 2018, 12, 2575-2581.	4.4	50
38	Differential bacterial dynamics promote emergent community robustness to lake mixing: an epilimnion to hypolimnion transplant experiment. <i>Environmental Microbiology</i> , 2010, 12, 455-466.	1.8	44
39	16S rRNA/rRNA Gene Ratios and Cell Activity Staining Reveal Consistent Patterns of Microbial Activity in Plant-Associated Soil. <i>MSystems</i> , 2019, 4, .	1.7	44
40	When, where and how does microbial community composition matter?. <i>Frontiers in Microbiology</i> , 2014, 5, 497.	1.5	43
41	Temporal dynamics of bacterial communities during seed development and maturation. <i>FEMS Microbiology Ecology</i> , 2020, 96, .	1.3	43
42	Ecological selection for small microbial genomes along a temperate-to-thermal soil gradient. <i>Nature Microbiology</i> , 2019, 4, 55-61.	5.9	42
43	Community structure explains antibiotic resistance gene dynamics over a temperature gradient in soil. <i>FEMS Microbiology Ecology</i> , 2018, 94, .	1.3	40
44	Lifestyles of rarity: understanding heterotrophic strategies to inform the ecology of the microbial rare biosphere. <i>Aquatic Microbial Ecology</i> , 2016, 78, 51-63.	0.9	39
45	Streptomycin Application Has No Detectable Effect on Bacterial Community Structure in Apple Orchard Soil. <i>Applied and Environmental Microbiology</i> , 2013, 79, 6617-6625.	1.4	38
46	Manipulating Wild and Tamed Phytobiomes: Challenges and Opportunities. <i>Phytobiomes Journal</i> , 2019, 3, 3-21.	1.4	38
47	Dormancy dynamics and dispersal contribute to soil microbiome resilience. <i>Philosophical Transactions of the Royal Society B: Biological Sciences</i> , 2020, 375, 20190255.	1.8	38
48	Toward a Generalizable Framework of Disturbance Ecology Through Crowdsourced Science. <i>Frontiers in Ecology and Evolution</i> , 2021, 9, .	1.1	34
49	Computing Workflows for Biologists: A Roadmap. <i>PLoS Biology</i> , 2015, 13, e1002303.	2.6	33
50	Can the black box be cracked? The augmentation of microbial ecology by high-resolution, automated sensing technologies. <i>ISME Journal</i> , 2009, 3, 881-888.	4.4	32
51	Seasonal and Episodic Lake Mixing Stimulate Differential Planktonic Bacterial Dynamics. <i>Microbial Ecology</i> , 2010, 59, 546-554.	1.4	31
52	Broadening Participation in Scientific Conferences during the Era of Social Distancing. <i>Trends in Microbiology</i> , 2020, 28, 949-952.	3.5	31
53	Seasonal Dynamics of Core Fungi in the Switchgrass Phyllosphere, and Co-Occurrence with Leaf Bacteria. <i>Phytobiomes Journal</i> , 2021, 5, 60-68.	1.4	29
54	Endophytic Microbiome Variation Among Single Plant Seeds. <i>Phytobiomes Journal</i> , 2022, 6, 45-55.	1.4	24

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55	NITROGEN CHANGES AND DOMAIN BACTERIA RIBOTYPE DIVERSITY IN SOILS OVERLYING THE CENTRALIA, PENNSYLVANIA UNDERGROUND COAL MINE FIRE. <i>Soil Science</i> , 2005, 170, 191-201.	0.9	22
56	Lineage-Specific Responses of Microbial Communities to Environmental Change. <i>Applied and Environmental Microbiology</i> , 2013, 79, 39-47.	1.4	20
57	RefSoil+: a Reference Database for Genes and Traits of Soil Plasmids. <i>MSystems</i> , 2019, 4, .	1.7	16
58	Taxonomically-linked growth phenotypes during arsenic stress among arsenic resistant bacteria isolated from soils overlying the Centralia coal seam fire. <i>PLoS ONE</i> , 2018, 13, e0191893.	1.1	16
59	Temporal Dynamics of South End Tidal Creek (Sapelo Island, Georgia) Bacterial Communities. <i>Applied and Environmental Microbiology</i> , 2009, 75, 1058-1064.	1.4	14
60	Abiotic Treatment to Common Bean Plants Results in an Altered Endophytic Seed Microbiome. <i>Microbiology Spectrum</i> , 2022, 10, e0021021.	1.2	12
61	Strategies for Building Computing Skills To Support Microbiome Analysis: a Five-Year Perspective from the EDAMAME Workshop. <i>MSystems</i> , 2019, 4, .	1.7	10
62	“Gradual Entrainment Lake Inverter” (GELI): A novel device for experimental lake mixing. <i>Limnology and Oceanography: Methods</i> , 2011, 9, 14-28.	1.0	9
63	Understanding Microbiome Stability in a Changing World. <i>MSystems</i> , 2018, 3, .	1.7	9
64	Resource Competition and Host Feedbacks Underlie Regime Shifts in Gut Microbiota. <i>American Naturalist</i> , 2021, 198, 1-12.	1.0	9
65	Biogeography and Diversity of Multi-Trophic Root Zone Microbiomes in Michigan Apple Orchards: Analysis of Rootstock, Scion, and Local Growing Region. <i>Phytobiomes Journal</i> , 2020, 4, 122-132.	1.4	8
66	Managing Plant Microbiomes for Sustainable Biofuel Production. <i>Phytobiomes Journal</i> , 2021, 5, 3-13.	1.4	8
67	Exometabolite Dynamics over Stationary Phase Reveal Strain-Specific Responses. <i>MSystems</i> , 2020, 5, .	1.7	7
68	Locally Adapted <i>Mimulus</i> Ecotypes Differentially Impact Rhizosphere Bacterial and Archaeal Communities in an Environment-Dependent Manner. <i>Phytobiomes Journal</i> , 2020, 4, 53-63.	1.4	6
69	A town on fire! Integrating 16S rRNA gene amplicon analyses into an undergraduate microbiology lecture class. <i>FEMS Microbiology Letters</i> , 2018, 365, .	0.7	5
70	Comparing gut resistome composition among patients with acute <i>Campylobacter</i> infections and healthy family members. <i>Scientific Reports</i> , 2021, 11, 22368.	1.6	2
71	Frontiers and Opportunities in Bioenergy Crop Microbiome Research Networks. <i>Phytobiomes Journal</i> , 2022, 6, 118-126.	1.4	1