Bernd W Brandt

List of Publications by Year in descending order

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Version: 2024-02-01

60 papers

1,842 citations

304743 22 h-index 289244 40 g-index

61 all docs

61 docs citations

61 times ranked

3057 citing authors

#	Article	IF	CITATIONS
1	The microbiome of dental and periâ€implant subgingival plaque during periâ€implant mucositis therapy: A randomized clinical trial. Journal of Clinical Periodontology, 2022, 49, 28-38.	4.9	18
2	Comparability of microbiota of swabbed and spit saliva. European Journal of Oral Sciences, 2022, 130, e12858.	1.5	5
3	Reply. Arthritis and Rheumatology, 2022, 74, 1297-1298.	5.6	O
4	The Evaluation of the Effects of Two Probiotic Strains on the Oral Ecosystem: A Randomized Clinical Trial. Frontiers in Oral Health, 2022, 3, 825017.	3.0	1
5	Long-Term Analysis of Resilience of the Oral Microbiome in Allogeneic Stem Cell Transplant Recipients. Microorganisms, 2022, 10, 734.	3.6	8
6	Adaptive changes of sediment microbial communities associated with cleanup of oil spills in Nigerian mangrove forests. Marine Pollution Bulletin, 2022, 176, 113406.	5.0	10
7	Manipulation of Saliva-Derived Microcosm Biofilms To Resemble Dysbiotic Subgingival Microbiota. Applied and Environmental Microbiology, 2021, 87, .	3.1	6
8	Optimizing the quality of clinical studies on oral microbiome: A practical guide for planning, performing, and reporting. Periodontology 2000, 2021, 85, 210-236.	13.4	51
9	Influence of short- and long-term exposure on the biodegradation capacity of activated sludge microbial communities in ready biodegradability tests. Environmental Science: Water Research and Technology, 2021, 7, 107-121.	2.4	3
10	Microbiome analysis of feline odontoclastic resorptive lesion (FORL) and feline oral health. Journal of Medical Microbiology, 2021, 70, .	1.8	3
11	Effects of DNA preservation solution and DNA extraction methods on microbial community profiling of soil. Folia Microbiologica, 2021, 66, 597-606.	2.3	14
12	High biodiversity in a benzene-degrading nitrate-reducing culture is sustained by a few primary consumers. Communications Biology, 2021, 4, 530.	4.4	11
13	The microbiological load and microbiome of the Dutch dental unit; â€~please, hold your breath'. Water Research, 2021, 200, 117205.	11.3	9
14	Submucosal microbiome of periâ€implant sites: A crossâ€sectional study. Journal of Clinical Periodontology, 2021, 48, 1228-1239.	4.9	21
15	16S rDNA sequencing and metadata of Dutch dental unit water. Data in Brief, 2021, 37, 107221.	1.0	2
16	Saliva-derived microcosm biofilms grown on different oral surfaces in vitro. Npj Biofilms and Microbiomes, 2021, 7, 74.	6.4	8
17	Differences in the Oral Microbiome in Patients With Early Rheumatoid Arthritis and Individuals at Risk of Rheumatoid Arthritis Compared to Healthy Individuals. Arthritis and Rheumatology, 2021, 73, 1986-1993.	5.6	33
18	Reprocessing 16S rRNA Gene Amplicon Sequencing Studies: (Meta)Data Issues, Robustness, and Reproducibility. Frontiers in Cellular and Infection Microbiology, 2021, 11, 720637.	3.9	4

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19	Comparison of Red-Complex Bacteria Between Saliva and Subgingival Plaque of Periodontitis Patients: A Systematic Review and Meta-Analysis. Frontiers in Cellular and Infection Microbiology, 2021, 11, 727732.	3.9	28
20	Correlating Biodegradation Kinetics of 2,4-Dichlorophenoxyacetic Acid (2,4-D) and 2,4,5-Trichlorophenoxyacetic Acid (2,4,5-T) to the Dynamics of Microbial Communities Originating From Soil in Vietnam Contaminated With Herbicides. Frontiers in Sustainable Cities, 2021, 3, .	2.4	8
21	Home sampling is a feasible method for oral microbiota analysis for infants and mothers. Journal of Dentistry, 2020, 100, 103428.	4.1	4
22	An in-vitro dynamic flow model for translational research into dental unit water system biofilms. Journal of Microbiological Methods, 2020, 171, 105879.	1.6	9
23	Short-Chain <i>N</i> -Acylhomoserine Lactone Quorum-Sensing Molecules Promote Periodontal Pathogens in <i>In Vitro</i> Oral Biofilms. Applied and Environmental Microbiology, 2020, 86, .	3.1	26
24	Impact of the early-life skin microbiota on the development of canine atopic dermatitis in a high-risk breed birth cohort. Scientific Reports, 2020, 10, 1044.	3.3	11
25	Dysbiosis of the Oral Ecosystem in Severe Congenital Neutropenia Patients. Proteomics - Clinical Applications, 2020, 14, e1900058.	1.6	7
26	Dental aerosols: microbial composition and spatial distribution. Journal of Oral Microbiology, 2020, 12, 1762040.	2.7	72
27	Interkingdom interactions on the denture surface: Implications for oral hygiene. Biofilm, 2019, 1, 100002 .	3.8	15
28	Tumor microbiome: Pancreatic cancer and duodenal fluids contain multitudes, …but do they contradict themselves?. Critical Reviews in Oncology/Hematology, 2019, 144, 102824.	4.4	6
29	Microbial changes in relation to oral mucositis in autologous hematopoietic stem cell transplantation recipients. Scientific Reports, 2019, 9, 16929.	3.3	32
30	Physico-chemical and biological aspects of a serially connected lab-scale constructed wetland-stabilization tank-GAC slow sand filtration system during removal of selected PPCPs. Chemical Engineering Journal, 2019, 369, 1109-1118.	12.7	29
31	Regrowth of Microcosm Biofilms on Titanium Surfaces After Various Antimicrobial Treatments. Frontiers in Microbiology, 2019, 10, 2693.	3.5	14
32	Long-term impact of oral surgery with or without amoxicillin on the oral microbiome-A prospective cohort study. Scientific Reports, 2019, 9, 18761.	3.3	12
33	Diversity of SpaP in genetic and salivary agglutinin mediated adherence among Streptococcus mutans strains. Scientific Reports, 2019, 9, 19943.	3.3	12
34	The efficacy of whole human genome capture on ancient dental calculus and dentin. American Journal of Physical Anthropology, 2019, 168, 496-509.	2.1	24
35	Microcosm biofilms cultured from different oral niches in periodontitis patients. Journal of Oral Microbiology, 2019, 11, 1551596.	2.7	38
36	Microbiomes associated with bovine periodontitis and oral health. Veterinary Microbiology, 2018, 218, 1-6.	1.9	33

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37	Limited added value of fungal ITS amplicon sequencing in the study of bovine abortion. Heliyon, 2018, 4, e00915.	3.2	3
38	Genetic Loci Associated With Fluoride Resistance in Streptococcus mutans. Frontiers in Microbiology, 2018, 9, 3093.	3.5	16
39	Subgingival microbiome of rheumatoid arthritis patients in relation to their disease status and periodontal health. PLoS ONE, 2018, 13, e0202278.	2.5	50
40	Red fluorescence of dental plaque in children â€"A cross-sectional study. Journal of Dentistry, 2017, 58, 40-47.	4.1	17
41	<i>Candida albicans</i> alters the bacterial microbiome of early <i>in vitro</i> oral biofilms. Journal of Oral Microbiology, 2017, 9, 1270613.	2.7	57
42	Fungal mitochondrial oxygen consumption induces the growth of strict anaerobic bacteria. Fungal Genetics and Biology, 2017, 109, 1-6.	2.1	32
43	Fluoride resistance in <i>Streptococcus mutans</i> : a mini review. Journal of Oral Microbiology, 2017, 9, 1344509.	2.7	99
44	The mycobiome of root canal infections is correlated to the bacteriome. Clinical Oral Investigations, 2017, 21, 1871-1881.	3.0	55
45	Changes in the oral ecosystem induced by the use of 8% arginine toothpaste. Archives of Oral Biology, 2017, 73, 79-87.	1.8	39
46	The Fitness Cost of Fluoride Resistance for Different Streptococcus mutans Strains in Biofilms. Frontiers in Microbiology, 2017, 8, 1630.	3.5	16
47	A novel compound to maintain a healthy oral plaque ecology <i>in vitro</i> . Journal of Oral Microbiology, 2016, 8, 32513.	2.7	19
48	Microbial profiles at baseline and not the use of antibiotics determine the clinical outcome of the treatment of chronic periodontitis. Scientific Reports, 2016, 6, 20205.	3.3	51
49	A Single Nucleotide Change in the Promoter <i>mutp</i> Enhances Fluoride Resistance of Streptococcus mutans. Antimicrobial Agents and Chemotherapy, 2016, 60, 7509-7512.	3.2	21
50	Integrating Candida albicans metabolism with biofilm heterogeneity by transcriptome mapping. Scientific Reports, 2016, 6, 35436.	3.3	39
51	Nitrate and the Origin of Saliva Influence Composition and Short Chain Fatty Acid Production of Oral Microcosms. Microbial Ecology, 2016, 72, 479-492.	2.8	58
52	metaModules identifies key functional subnetworks in microbiome-related disease. Bioinformatics, 2016, 32, 1678-1685.	4.1	21
53	Intrinsic challenges in ancient microbiome reconstruction using 16S rRNA gene amplification. Scientific Reports, 2015, 5, 16498.	3.3	153
54	Proteins and peptides in parotid saliva of irradiated patients compared to that of healthy controls using SELDI-TOF-MS. BMC Research Notes, 2015, 8, 639.	1.4	12

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55	The Oral Microbiome of Denture Wearers Is Influenced by Levels of Natural Dentition. PLoS ONE, 2015, 10, e0137717.	2.5	82
56	NGS-eval: NGS Error analysis and novel sequence VAriant detection tool. Nucleic Acids Research, 2015, 43, W301-W305.	14.5	16
57	Same Exposure but Two Radically Different Responses to Antibiotics: Resilience of the Salivary Microbiome versus Long-Term Microbial Shifts in Feces. MBio, 2015, 6, e01693-15.	4.1	333
58	TreeSeq, a Fast and Intuitive Tool for Analysis of Whole Genome and Metagenomic Sequence Data. PLoS ONE, 2015, 10, e0123851.	2.5	3
59	Unraveling the outcome of 16S rDNA-based taxonomy analysis through mock data and simulations. Bioinformatics, 2014, 30, 1530-1538.	4.1	29
60	TaxMan: a server to trim rRNA reference databases and inspect taxonomic coverage. Nucleic Acids Research, 2012, 40, W82-W87.	14.5	33