Pei-Ying Hong

List of Publications by Year in descending order

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94269 128067 4,232 115 37 60 citations h-index g-index papers 119 119 119 6513 docs citations times ranked citing authors all docs

#	Article	IF	CITATIONS
1	Biofilm Formation Characteristics of Bacterial Isolates Retrieved from a Reverse Osmosis Membrane. Environmental Science & Env	4.6	219
2	Removal of bacterial contaminants and antibiotic resistance genes by conventional wastewater treatment processes in Saudi Arabia: Is the treated wastewater safe to reuse for agricultural irrigation?. Water Research, 2015, 73, 277-290.	5.3	180
3	Pyrosequencing Analysis of Bacterial Biofilm Communities in Water Meters of a Drinking Water Distribution System. Applied and Environmental Microbiology, 2010, 76, 5631-5635.	1.4	166
4	Phylogenetic analysis of the fecal microbial community in herbivorous land and marine iguanas of the Gal $ ilde{A}_i$ pagos Islands using 16S rRNA-based pyrosequencing. ISME Journal, 2011, 5, 1461-1470.	4.4	142
5	Xylan utilization in human gut commensal bacteria is orchestrated by unique modular organization of polysaccharide-degrading enzymes. Proceedings of the National Academy of Sciences of the United States of America, 2014, 111, E3708-17.	3.3	137
6	Quantitative effects of position and type of single mismatch on single base primer extension. Journal of Microbiological Methods, 2009, 77, 267-275.	0.7	127
7	Environmental and Public Health Implications of Water Reuse: Antibiotics, Antibiotic Resistant Bacteria, and Antibiotic Resistance Genes. Antibiotics, 2013, 2, 367-399.	1.5	100
8	Metagenomic analysis of DNA viruses in a wastewater treatment plant in tropical climate. Environmental Microbiology, 2012, 14, 441-452.	1.8	98
9	Removal of Antibiotic-Resistant Bacteria and Antibiotic Resistance Genes Affected by Varying Degrees of Fouling on Anaerobic Microfiltration Membranes. Environmental Science & Environmental Science	4.6	95
10	Monitoring airborne biotic contaminants in the indoor environment of pig and poultry confinement buildings. Environmental Microbiology, 2012, 14, 1420-1431.	1.8	94
11	Pyrosequencing-Based Analysis of the Mucosal Microbiota in Healthy Individuals Reveals Ubiquitous Bacterial Groups and Micro-Heterogeneity. PLoS ONE, 2011, 6, e25042.	1.1	91
12	Hydroxyl Functionalized Polytriazole- <i>co</i> -polyoxadiazole as Substrates for Forward Osmosis Membranes. ACS Applied Materials & Samp; Interfaces, 2015, 7, 3960-3973.	4.0	88
13	Comparative Analysis of Fecal Microbiota in Infants with and without Eczema. PLoS ONE, 2010, 5, e9964.	1.1	87
14	Reusing Treated Wastewater: Consideration of the Safety Aspects Associated with Antibiotic-Resistant Bacteria and Antibiotic Resistance Genes. Water (Switzerland), 2018, 10, 244.	1.2	83
15	Sustainable organic loading rate and energy recovery potential of mesophilic anaerobic membrane bioreactor for municipal wastewater treatment. Bioresource Technology, 2014, 166, 326-334.	4.8	78
16	Water Disinfection Byproducts Increase Natural Transformation Rates of Environmental DNA in <i>Acinetobacter baylyi</i> ADP1. Environmental Science & Environmental DNA in Control of Environmental DNA in Con	4.6	76
17	Design of anaerobic membrane bioreactors for the valorization of dilute organic carbon waste streams. Energy and Environmental Science, 2016, 9, 1102-1112.	15.6	74
18	Monitoring the Perturbation of Soil and Groundwater Microbial Communities Due to Pig Production Activities. Applied and Environmental Microbiology, 2013, 79, 2620-2629.	1.4	73

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19	Interfacial Polymerization of Zwitterionic Building Blocks for High-Flux Nanofiltration Membranes. Langmuir, 2019, 35, 1284-1293.	1.6	71
20	Does chlorination of seawater reverse osmosis membranes control biofouling?. Water Research, 2015, 78, 84-97.	5.3	67
21	Organic micropollutants in aerobic and anaerobic membrane bioreactors: Changes in microbial communities and gene expression. Bioresource Technology, 2016, 218, 882-891.	4.8	66
22	Estimating the minimum number of SARS-CoV-2 infected cases needed to detect viral RNA in wastewater: To what extent of the outbreak can surveillance of wastewater tell us?. Environmental Research, 2021, 195, 110748.	3.7	64
23	Ecological drift and local exposures drive enteric bacterial community differences within species of Galápagos iguanas. Molecular Ecology, 2012, 21, 1779-1788.	2.0	61
24	Silver-Enhanced Block Copolymer Membranes with Biocidal Activity. ACS Applied Materials & Samp; Interfaces, 2014, 6, 18497-18501.	4.0	58
25	CARRIAGE OF ANTIBIOTIC-RESISTANT ENTERIC BACTERIA VARIES AMONG SITES IN GALÃPAGOS REPTILES. Journal of Wildlife Diseases, 2012, 48, 56-67.	0.3	57
26	Characterization of biofoulants illustrates different membrane fouling mechanisms for aerobic and anaerobic membrane bioreactors. Separation and Purification Technology, 2016, 157, 192-202.	3.9	56
27	<scp><i>H</i></scp> <i>alomonas sulfidaeris</i> êdominated microbial community inhabits a 1.8 kmâ€deep subsurface <scp>C</scp> ambrian <scp>S</scp> andstone reservoir. Environmental Microbiology, 2014, 16, 1695-1708.	1.8	52
28	Molecular-based detection of potentially pathogenic bacteria in membrane bioreactor (MBR) systems treating municipal wastewater: a case study. Environmental Science and Pollution Research, 2017, 24, 5370-5380.	2.7	52
29	Two New Xylanases with Different Substrate Specificities from the Human Gut Bacterium Bacteroides intestinalis DSM 17393. Applied and Environmental Microbiology, 2014, 80, 2084-2093.	1.4	50
30	Drinking Water Microbiome Project: Is it Time?. Trends in Microbiology, 2019, 27, 670-677.	3.5	50
31	Differences in microbial communities and performance between suspended and attached growth anaerobic membrane bioreactors treating synthetic municipal wastewater. Environmental Science: Water Research and Technology, 2015, 1, 800-813.	1.2	48
32	Making Waves: Collaboration in the time of SARS-CoV-2 - rapid development of an international co-operation and wastewater surveillance database to support public health decision-making. Water Research, 2021, 199, 117167.	5.3	48
33	Increasing tetracycline concentrations on the performance and communities of mixed microalgae-bacteria photo-bioreactors. Algal Research, 2018, 29, 249-256.	2.4	46
34	Relative Abundance of <i> Bacteroides < /i > spp. in Stools and Wastewaters as Determined by Hierarchical Oligonucleotide Primer Extension. Applied and Environmental Microbiology, 2008, 74, 2882-2893.</i>	1.4	45
35	Isolation and Characterization of NDM-Positive Escherichia coli from Municipal Wastewater in Jeddah, Saudi Arabia. Antimicrobial Agents and Chemotherapy, 2016, 60, 5223-5231.	1.4	44
36	Performance and microbial community variations of anaerobic digesters under increasing tetracycline concentrations. Applied Microbiology and Biotechnology, 2017, 101, 5505-5517.	1.7	42

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37	Identification and characterization of core sludge and biofilm microbiota in anaerobic membrane bioreactors. Environment International, 2019, 133, 105165.	4.8	40
38	Evaluation of stool microbiota signatures in two cohorts of Asian (Singapore and Indonesia) newborns at risk of atopy. BMC Microbiology, 2011, 11, 193.	1.3	39
39	Inactivation and Gene Expression of a Virulent Wastewater <i>Escherichia coli</i> Strain and the Nonvirulent Commensal <i>Escherichia coli</i> DSM1103 Strain upon Solar Irradiation. Environmental Science & Echnology, 2017, 51, 3649-3659.	4.6	38
40	Hollow fiber membrane lumen modified by polyzwitterionic grafting. Journal of Membrane Science, 2017, 522, 1-11.	4.1	38
41	Effect of pre-acclimation of granular activated carbon on microbial electrolysis cell startup and performance. Bioelectrochemistry, 2017, 113, 20-25.	2.4	36
42	Acquisition of Extracellular DNA by <i>Acinetobacter baylyi</i> ADP1 in Response to Solar and UV-C _{254nm} Disinfection. Environmental Science & Environmental Science	4.6	35
43	Genomic characterization of NDM-1 and 5, and OXA-181 carbapenemases in uropathogenic Escherichia coli isolates from Riyadh, Saudi Arabia. PLoS ONE, 2018, 13, e0201613.	1.1	34
44	Anaerobic Membrane Bioreactor Effluent Reuse: A Review of Microbial Safety Concerns. Fermentation, 2017, 3, 39.	1.4	33
45	Removal of bacterial cells, antibiotic resistance genes and integrase genes by on-site hospital wastewater treatment plants: surveillance of treated hospital effluent quality. Environmental Science: Water Research and Technology, 2017, 3, 293-303.	1.2	31
46	Evaluation of two autoinducer-2 quantification methods for application in marine environments. Journal of Applied Microbiology, 2018, 124, 1469-1479.	1.4	31
47	Microbial diversity and biosignatures of amorphous silica deposits in orthoquartzite caves. Scientific Reports, 2018, 8, 17569.	1.6	30
48	Influence of biofilm thickness on the removal of thirteen different organic micropollutants via a Membrane Aerated Biofilm Reactor (MABR). Journal of Hazardous Materials, 2022, 432, 128698.	6.5	30
49	Assessing the Groundwater Quality at a Saudi Arabian Agricultural Site and the Occurrence of Opportunistic Pathogens on Irrigated Food Produce. International Journal of Environmental Research and Public Health, 2015, 12, 12391-12411.	1.2	28
50	Synthesis of highly porous poly(tert-butyl acrylate)-b-polysulfone-b-poly(tert-butyl acrylate) asymmetric membranes. Polymer Chemistry, 2016, 7, 3076-3089.	1.9	28
51	Rapid Size-Based Protein Discrimination inside Hybrid Isoporous Membranes. ACS Applied Materials & Lamp; Interfaces, 2019, 11, 8507-8516.	4.0	28
52	Quorum Sensing and the Use of Quorum Quenchers as Natural Biocides to Inhibit Sulfate-Reducing Bacteria. Antibiotics, 2016, 5, 39.	1.5	27
53	Membrane Bioreactor-Based Wastewater Treatment Plant in Saudi Arabia: Reduction of Viral Diversity, Load, and Infectious Capacity. Water (Switzerland), 2017, 9, 534.	1.2	27
54	Lignocellulose-derived thin stillage composition and efficient biological treatment with a high-rate hybrid anaerobic bioreactor system. Biotechnology for Biofuels, 2016, 9, 120.	6.2	25

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55	Chlorination or monochloramination: Balancing the regulated trihalomethane formation and microbial inactivation in marine aquaculture waters. Aquaculture, 2017, 480, 94-102.	1.7	25
56	Bacteriophages To Sensitize a Pathogenic New Delhi Metallo \hat{l}^2 -Lactamase-Positive <i>Escherichia coli</i> to Solar Disinfection. Environmental Science & Environmental S	4.6	25
57	Salinity-Mediated Increment in Sulfate Reduction, Biofilm Formation, and Quorum Sensing: A Potential Connection Between Quorum Sensing and Sulfate Reduction?. Frontiers in Microbiology, 2019, 10, 188.	1.5	25
58	Effect of Quorum Sensing on the Ability of <i>Desulfovibrio vulgaris</i> To Form Biofilms and To Biocorrode Carbon Steel in Saline Conditions. Applied and Environmental Microbiology, 2019, 86, .	1.4	25
59	Molecular-based approaches to characterize coastal microbial community and their potential relation to the trophic state of Red Sea. Scientific Reports, 2015, 5, 9001.	1.6	24
60	The Effect of the 2015 Earthquake on the Bacterial Community Compositions in Water in Nepal. Frontiers in Microbiology, 2017, 8, 2380.	1.5	24
61	Translational Molecular Ecology in practice: Linking DNA-based methods to actionable marine environmental management. Science of the Total Environment, 2020, 744, 140780.	3.9	24
62	Dynamics of microbial communities in an integrated ultrafiltration–reverse osmosis desalination pilot plant located at theÂArabian Gulf. Desalination and Water Treatment, 2016, 57, 16310-16323.	1.0	23
63	The use of UV/H2O2 to facilitate removal of emerging contaminants in anaerobic membrane bioreactor effluents. Environmental Research, 2021, 198, 110479.	3.7	23
64	A highâ€throughput and quantitative hierarchical oligonucleotide primer extension (HOPE)â€based approach to identify sources of faecal contamination in water bodies. Environmental Microbiology, 2009, 11, 1672-1681.	1.8	22
65	Phenotypic and Phylogenetic Identification of Coliform Bacteria Obtained Using 12 Coliform Methods Approved by the U.S. Environmental Protection Agency. Applied and Environmental Microbiology, 2015, 81, 6012-6023.	1.4	21
66	Antibiofilm effect enhanced by modification of 1,2,3-triazole and palladium nanoparticles on polysulfone membranes. Scientific Reports, 2016, 6, 24289.	1.6	21
67	Metagenomics-based evaluation of groundwater microbial profiles in response to treated wastewater discharge. Environmental Research, 2020, 180, 108835.	3.7	18
68	Fate and Persistence of a Pathogenic NDM-1-Positive Escherichia coli Strain in Anaerobic and Aerobic Sludge Microcosms. Applied and Environmental Microbiology, 2017, 83, .	1.4	17
69	Identification of methanogenesis and syntrophy as important microbial metabolic processes for optimal thermophilic anaerobic digestion of energy cane thin stillage. Bioresource Technology Reports, 2019, 7, 100254.	1.5	17
70	An Increase of Abundance and Transcriptional Activity for Acinetobacter junii Post Wastewater Treatment. Water (Switzerland), 2018, 10, 436.	1.2	16
71	Application of hierarchical oligonucleotide primer extension (HOPE) to assess relative abundances of ammonia- and nitrite-oxidizing bacteria. BMC Microbiology, 2017, 17, 85.	1.3	15
72	Metagenomic-Based Study of the Phylogenetic and Functional Gene Diversity in Galápagos Land and Marine Iguanas. Microbial Ecology, 2015, 69, 444-456.	1.4	14

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73	Understanding the antifouling mechanisms related to copper oxide and zinc oxide nanoparticles in anaerobic membrane bioreactors. Environmental Science: Nano, 2019, 6, 3467-3479.	2.2	14
74	Metagenomics as a Tool To Monitor Reclaimed-Water Quality. Applied and Environmental Microbiology, 2020, 86, .	1.4	14
75	Recent Update on UV Disinfection to Fulfill the Disinfection Credit Value for Enteric Viruses in Water. Environmental Science & Environmental Science	4.6	14
76	Discovering, Characterizing, and Applying Acyl Homoserine Lactone-Quenching Enzymes to Mitigate Microbe-Associated Problems Under Saline Conditions. Frontiers in Microbiology, 2019, 10, 823.	1.5	13
77	Water quality, seasonality, and trajectory of an aquaculture-wastewater plume in the Red Sea. Aquaculture Environment Interactions, 2018, 10, 61-77.	0.7	13
78	DNAzyme-based biosensor as a rapid and accurate verification tool to complement simultaneous enzyme-based media forE. colidetection. Environmental Science: Water Research and Technology, 2019, 5, 2260-2268.	1.2	12
79	Transition from unclassified Ktedonobacterales to Actinobacteria during amorphous silica precipitation in a quartzite cave environment. Scientific Reports, 2021, 11, 3921.	1.6	12
80	A type dependent effect of treated wastewater matrix on seed germination and food production. Science of the Total Environment, 2021, 769, 144573.	3.9	12
81	Attached-growth configuration outperforms continuously stirred tank anaerobic membrane bioreactors in alleviating membrane biofouling. Environmental Research, 2021, 199, 111272.	3.7	12
82	Immune-modulatory genomic properties differentiate gut microbiota of infants with and without eczema. PLoS ONE, 2017, 12, e0184955.	1.1	12
83	In situ growth of biocidal AgCl crystals in the top layer of asymmetric polytriazole membranes. RSC Advances, 2016, 6, 46696-46701.	1.7	11
84	Bacteriophage Infectivity Against Pseudomonas aeruginosa in Saline Conditions. Frontiers in Microbiology, 2018, 9, 875.	1.5	11
85	UV and bacteriophages as a chemical-free approach for cleaning membranes from anaerobic bioreactors. Proceedings of the National Academy of Sciences of the United States of America, 2021, 118, .	3.3	11
86	Sliding window neural network based sensing of bacteria in wastewater treatment plants. Journal of Process Control, 2022, 110, 35-44.	1.7	11
87	A Robust, Safe, and Scalable Magnetic Nanoparticle Workflow for RNA Extraction of Pathogens from Clinical and Wastewater Samples. Global Challenges, 2021, 5, 2000068.	1.8	10
88	Persistence of Bacteroides ovatus under simulated sunlight irradiation. BMC Microbiology, 2014, 14, 178.	1.3	9
89	An aerated and fluidized bed membrane bioreactor for effective wastewater treatment with low membrane fouling. Environmental Science: Water Research and Technology, 2016, 2, 994-1003.	1.2	9
90	Varying occurrence of extendedâ€spectrum betaâ€lactamase bacteria among three produce types. Journal of Food Safety, 2018, 38, e12373.	1.1	9

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91	Plankton community assessment in anthropogenic-impacted oligotrophic coastal regions. Environmental Science and Pollution Research, 2018, 25, 31017-31030.	2.7	9
92	Genome-Resolved Metagenomics and Antibiotic Resistance Genes Analysis in Reclaimed Water Distribution Systems. Water (Switzerland), 2020, 12, 3477.	1.2	7
93	Editorial perspective: Viruses in wastewater: Wading into the knowns and unknowns. Environmental Research, 2021, 196, 110255.	3.7	7
94	Impact of acclimation methods on microbial communities and performance of anaerobic fluidized bed membrane bioreactors. Environmental Science: Water Research and Technology, 2016, 2, 1041-1048.	1.2	6
95	Thin Film Polyamide Membranes with Photoresponsive Antibacterial Activity. ChemistrySelect, 2017, 2, 6612-6616.	0.7	5
96	Nanoparticles applied in membrane bioreactors: potential impact on reactor performance and microbial communities., 2020,, 207-236.		5
97	Calibration and validation for a real-time membrane bioreactor: A sliding window approach. Journal of Process Control, 2021, 98, 92-105.	1.7	5
98	Enteric virus in reclaimed water from treatment plants with different multi-barrier strategies: Trade-off assessment in treatment extent and risks. Science of the Total Environment, 2021, 776, 146039.	3.9	5
99	Digital E. coli Counter: A Microfluidics and Computer Vision-Based DNAzyme Method for the Isolation and Specific Detection of E. coli from Water Samples. Biosensors, 2022, 12, 34.	2.3	5
100	Hierarchical Oligonucleotide Primer Extension as a Time- and Cost-Effective Approach for Quantitative Determination of <i>Bifidobacterium</i> spp. in Infant Feces. Applied and Environmental Microbiology, 2009, 75, 2573-2576.	1.4	4
101	Flexible isoporous air filters for high-efficiency particle capture. Polymer, 2021, 213, 123278.	1.8	4
102	Inactivation and Loss of Infectivity of Enterovirus 70 by Solar Irradiation. Water (Switzerland), 2019, 11, 64.	1.2	3
103	Temperature Responses of Heterotrophic Bacteria in Co-culture With a Red Sea Synechococcus Strain. Frontiers in Microbiology, 2021, 12, 612732.	1.5	3
104	Antibiotic-Resistant Bacteria and Resistance Genes in the Water–Food Nexus of the Agricultural Environment. , 2015, , 325-346.		2
105	Potential Dissemination of ARB and ARGs into Soil Through the Use of Treated Wastewater for Agricultural Irrigation: Is It a True Cause for Concern?. Soil Biology, 2017, , 105-139.	0.6	2
106	Nonlinear Model Predictive Control Design for BSM-MBR: Benchmark of Membrane Bioreactor. IFAC-PapersOnLine, 2020, 53, 16524-16530.	0.5	2
107	Bacterial cell numbers and community structures of seawater biofilms depend on the attachment substratum., 0, 97, 41-71.		2
108	Characterizing the Chemical Contaminants Diversity and Toxic Potential of Untreated Wastewater From a Drug Rehabilitation Hospital: Understanding Impact on Downstream Environment. Frontiers in Environmental Science, 2022, 10, .	1.5	2

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109	Editorial: Microbial Safety in Water Resources. Frontiers in Microbiology, 2018, 9, 3064.	1.5	1
110	A DNA-mimic contact-active functional group for antifouling ultrafiltration membranes. Chemosphere, 2019, 216, 669-676.	4.2	1
111	Elucidating the Role of Virulence Traits in the Survival of Pathogenic E. coli PI-7 Following Disinfection. Frontiers in Bioengineering and Biotechnology, 2020, 8, 614186.	2.0	0
112	Mitigating Antimicrobial Resistance Risks When Using Reclaimed Municipal Wastewater for Agriculture. Handbook of Environmental Chemistry, 2020, , 245-265.	0.2	0
113	Terrestrial Vertebrate Animal Metagenomics: Herbivorous Reptiles. , 2014, , 1-11.		0
114	Terrestrial Vertebrate Animal Metagenomics: Herbivorous Reptiles. , 2015, , 693-703.		0
115	Understanding microbial assembly on seawater reverse osmosis membranes to facilitate evaluation of seawater pretreatment options., 0, 170, 1-10.		0