

Gianluca Corno

List of Publications by Year in descending order

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Version: 2024-02-01

73
papers

3,010
citations

147566

31
h-index

174990

52
g-index

76
all docs

76
docs citations

76
times ranked

3508
citing authors

#	ARTICLE	IF	CITATIONS
1	Co-occurrence of integrase 1, antibiotic and heavy metal resistance genes in municipal wastewater treatment plants. <i>Water Research</i> , 2016, 94, 208-214.	5.3	397
2	Microplastics increase impact of treated wastewater on freshwater microbial community. <i>Environmental Pollution</i> , 2018, 234, 495-502.	3.7	195
3	Direct and Indirect Effects of Protist Predation on Population Size Structure of a Bacterial Strain with High Phenotypic Plasticity. <i>Applied and Environmental Microbiology</i> , 2006, 72, 78-86.	1.4	147
4	Fitness and Recovery of Bacterial Communities and Antibiotic Resistance Genes in Urban Wastewaters Exposed to Classical Disinfection Treatments. <i>Environmental Science & Technology</i> , 2016, 50, 10153-10161.	4.6	110
5	Constitutive presence of antibiotic resistance genes within the bacterial community of a large subalpine lake. <i>Molecular Ecology</i> , 2015, 24, 3888-3900.	2.0	108
6	Rainfall increases the abundance of antibiotic resistance genes within a riverine microbial community. <i>Environmental Pollution</i> , 2017, 226, 473-478.	3.7	103
7	Co-selection of antibiotic and heavy metal resistance in freshwater bacteria. <i>Journal of Limnology</i> , 2016, 75, .	0.3	98
8	Effluents of wastewater treatment plants promote the rapid stabilization of the antibiotic resistome in receiving freshwater bodies. <i>Water Research</i> , 2019, 158, 72-81.	5.3	82
9	Assessing the Influence of Vegan, Vegetarian and Omnivore Oriented Westernized Dietary Styles on Human Gut Microbiota: A Cross Sectional Study. <i>Frontiers in Microbiology</i> , 2018, 9, 317.	1.5	78
10	Contribution of microplastic particles to the spread of resistances and pathogenic bacteria in treated wastewaters. <i>Water Research</i> , 2021, 201, 117368.	5.3	67
11	Phylogenetic diversity of nonmarine picocyanobacteria. <i>FEMS Microbiology Ecology</i> , 2013, 85, 293-301.	1.3	66
12	The role of metal contamination in shaping microbial communities in heavily polluted marine sediments. <i>Environmental Pollution</i> , 2020, 265, 114823.	3.7	65
13	Structural and functional patterns of bacterial communities in response to protist predation along an experimental productivity gradient. <i>Environmental Microbiology</i> , 2008, 10, 2857-2871.	1.8	63
14	Are microplastic particles a hotspot for the spread and the persistence of antibiotic resistance in aquatic systems?. <i>Environmental Pollution</i> , 2021, 279, 116896.	3.7	60
15	Antibiotics promote aggregation within aquatic bacterial communities. <i>Frontiers in Microbiology</i> , 2014, 5, 297.	1.5	59
16	A global multinational survey of cefotaxime-resistant coliforms in urban wastewater treatment plants. <i>Environment International</i> , 2020, 144, 106035.	4.8	55
17	Persistence of antibiotic resistance genes in large subalpine lakes: the role of anthropogenic pollution and ecological interactions. <i>Hydrobiologia</i> , 2018, 824, 93-108.	1.0	52
18	Coaggregation in a microbial predator-prey system affects competition and trophic transfer efficiency. <i>Ecology</i> , 2013, 94, 870-881.	1.5	50

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19	Resistance to Biocides in <i>Listeria monocytogenes</i> Collected in Meat-Processing Environments. <i>Frontiers in Microbiology</i> , 2016, 7, 1627.	1.5	48
20	Impact of industrial wastewater on the dynamics of antibiotic resistance genes in a full-scale urban wastewater treatment plant. <i>Science of the Total Environment</i> , 2019, 646, 1204-1210.	3.9	47
21	Disinfection of urban wastewater by a new photo-Fenton like process using Cu-iminodisuccinic acid complex as catalyst at neutral pH. <i>Water Research</i> , 2018, 146, 206-215.	5.3	46
22	Diverse distribution of Toxin-Antitoxin II systems in <i>Salmonella enterica</i> serovars. <i>Scientific Reports</i> , 2016, 6, 28759.	1.6	44
23	<i>Daphnia</i> as a refuge for an antibiotic resistance gene in an experimental freshwater community. <i>Science of the Total Environment</i> , 2016, 571, 77-81.	3.9	43
24	A microbial perspective on biological invasions in aquatic ecosystems. <i>Hydrobiologia</i> , 2015, 746, 13-22.	1.0	40
25	Antibiotic disturbance affects aquatic microbial community composition and food web interactions but not community resilience. <i>Molecular Ecology</i> , 2019, 28, 1170-1182.	2.0	39
26	The mesopelagic anoxic Black Sea as an unexpected habitat for <i>Synechococcus</i> challenges our understanding of global "deep red fluorescence". <i>ISME Journal</i> , 2019, 13, 1676-1687.	4.4	39
27	Effects of nutrient availability and <i>Ochromonas</i> sp. predation on size and composition of a simplified aquatic bacterial community. <i>FEMS Microbiology Ecology</i> , 2006, 58, 354-363.	1.3	36
28	<i>Bacteria</i> , <i>Archaea</i> , and <i>Crenarchaeota</i> in the Epilimnion and Hypolimnion of a Deep Holo-Oligomictic Lake. <i>Applied and Environmental Microbiology</i> , 2009, 75, 7298-7300.	1.4	35
29	Grazing-induced <i>Synechococcus</i> microcolony formation: experimental insights from two freshwater phylotypes. <i>FEMS Microbiology Ecology</i> , 2016, 92, fiw154.	1.3	34
30	Defence strategies and antibiotic resistance gene abundance in enterococci under stress by exposure to low doses of peracetic acid. <i>Chemosphere</i> , 2017, 185, 480-488.	4.2	34
31	Combination of flow cytometry and molecular analysis to monitor the effect of UVC/H ₂ O ₂ vs UVC/H ₂ O ₂ /Cu-IDS processes on pathogens and antibiotic resistant genes in secondary wastewater effluents. <i>Water Research</i> , 2020, 184, 116194.	5.3	34
32	Long-term trends of epilimnetic and hypolimnetic bacteria and organic carbon in a deep holo-oligomictic lake. <i>Hydrobiologia</i> , 2010, 644, 279-287.	1.0	33
33	High-quality treated wastewater causes remarkable changes in natural microbial communities and <i>int1</i> gene abundance. <i>Water Research</i> , 2019, 167, 114895.	5.3	33
34	Ecology and Distribution of Thaumarchaea in the Deep Hypolimnion of Lake Maggiore. <i>Archaea</i> , 2015, 1-11.	2.3	32
35	Effects of predation pressure on bacterial abundance, diversity, and size-structure distribution in an oligotrophic system. <i>Journal of Limnology</i> , 2008, 67, 107.	0.3	30
36	Bacterial diversity and morphology in deep ultraoligotrophic Andean lakes: The role of UVR on vertical distribution. <i>Limnology and Oceanography</i> , 2009, 54, 1098-1112.	1.6	27

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37	Picocyanobacterial community structure and space-time dynamics in the subalpine Lake Maggiore (N.) Tj ETQq1 1 0.784314 rgBT /Overl	0.3	26
38	Tracing particulate matter and associated microorganisms in freshwaters. <i>Hydrobiologia</i> , 2017, 800, 145-154.	1.0	26
39	Erect macroalgae influence epilithic bacterial assemblages and reduce coral recruitment. <i>Marine Ecology - Progress Series</i> , 2018, 597, 65-77.	0.9	25
40	Picocyanobacterial assemblages in ultraoligotrophic Andean lakes reveal high regional microdiversity. <i>Journal of Plankton Research</i> , 2010, 32, 357-366.	0.8	24
41	Impact of disinfection processes on bacterial community in urban wastewater: Should we rethink microbial assessment methods?. <i>Journal of Environmental Chemical Engineering</i> , 2020, 8, 104393.	3.3	24
42	Every Coin Has a Back Side: Invasion by <i>Limnohabitans planktonicus</i> Promotes the Maintenance of Species Diversity in Bacterial Communities. <i>PLoS ONE</i> , 2012, 7, e51576.	1.1	23
43	Interspecific interactions drive chitin and cellulose degradation by aquatic microorganisms. <i>Aquatic Microbial Ecology</i> , 2015, 76, 27-37.	0.9	23
44	Bioplastic accumulates antibiotic and metal resistance genes in coastal marine sediments. <i>Environmental Pollution</i> , 2021, 291, 118161.	3.7	20
45	Photosynthetic characteristics and diversity of freshwater <i>Synechococcus</i> at two depths during different mixing conditions in a deep oligotrophic lake. <i>Journal of Limnology</i> , 2007, 66, 81.	0.3	19
46	Spatial distribution of antibiotic and heavy metal resistance genes in the Black Sea. <i>Marine Pollution Bulletin</i> , 2020, 160, 111635.	2.3	19
47	Elimination from wastewater of antibiotics reserved for hospital settings, with a Fenton process based on zero-valent iron. <i>Chemosphere</i> , 2021, 283, 131170.	4.2	19
48	Assessing antimicrobial resistance gene load in vegan, vegetarian and omnivore human gut microbiota. <i>International Journal of Antimicrobial Agents</i> , 2018, 52, 702-705.	1.1	18
49	Every fifth published metagenome is not available to science. <i>PLoS Biology</i> , 2020, 18, e3000698.	2.6	18
50	PET particles raise microbiological concerns for human health while tyre wear microplastic particles potentially affect ecosystem services in waters. <i>Journal of Hazardous Materials</i> , 2022, 429, 128397.	6.5	18
51	Evaluation and quantification of antimicrobial residues and antimicrobial resistance genes in two Italian swine farms. <i>Environmental Pollution</i> , 2019, 255, 113183.	3.7	17
52	Different substrates within a lake harbour connected but specialised microbial communities. <i>Hydrobiologia</i> , 2020, 847, 1689-1704.	1.0	17
53	Ultraviolet Radiation Induces Filamentation in Bacterial Assemblages from North Andean Patagonian Lakes. <i>Photochemistry and Photobiology</i> , 2010, 86, 871-881.	1.3	16
54	Mechanisms regulating CO ₂ and CH ₄ dynamics in the Azorean volcanic lakes (SãŁo Miguel Island,) Tj ETQq0 0 0 rgBT /Overl	0.3	13

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55	Genomic Comparison and Spatial Distribution of Different <i>Synechococcus</i> Phylotypes in the Black Sea. <i>Frontiers in Microbiology</i> , 2020, 11, 1979.	1.5	13
56	Bacterial grazing by mixotrophic flagellates and <i>Daphnia longispina</i> : a comparison in a fishless alpine lake. <i>Aquatic Microbial Ecology</i> , 2006, 42, 127-137.	0.9	13
57	The microbiome associated with two <i>Synechococcus</i> ribotypes at different levels of ecological interaction. <i>Journal of Phycology</i> , 2017, 53, 1151-1158.	1.0	10
58	Seasonality of the antibiotic resistance gene blaCTX-M in temperate Lake Maggiore. <i>Hydrobiologia</i> , 2019, 843, 143-153.	1.0	10
59	Microplastic retention in small and medium municipal wastewater treatment plants and the role of the disinfection. <i>Environmental Science and Pollution Research</i> , 2022, 29, 10535-10546.	2.7	9
60	Dynamics of bacteria and mixotrophic flagellates in an Alpine lake in relation to <i>Daphnia</i> population development. <i>Journal of Limnology</i> , 2002, 61, 177.	0.3	8
61	The vertical distribution of tetA and int11 in a deep lake is rather due to sedimentation than to resuspension. <i>FEMS Microbiology Ecology</i> , 2020, 96, .	1.3	8
62	Transparent exopolymer particles (TEP) are driven by chlorophyll <i>a</i> and mainly confined to the euphotic zone in a deep subalpine lake. <i>Inland Waters</i> , 2017, 7, 118-127.	1.1	7
63	Tetracycline modifies competitive interactions in experimental microcosms containing bacteria isolated from freshwater. <i>FEMS Microbiology Ecology</i> , 2014, 90, 168-174.	1.3	6
64	Contribution of plasmidome, metal resistome and integrases to the persistence of the antibiotic resistome in aquatic environments. <i>Environmental Pollution</i> , 2022, 297, 118774.	3.7	6
65	The mixotrophic flagellates as key organisms from DOC to <i>Daphnia</i> in an oligotrophic alpine lake. <i>Verhandlungen Der Internationalen Vereinigung Fur Theoretische Und Angewandte Limnologie International Association of Theoretical and Applied Limnology</i> , 2002, 28, 392-395.	0.1	5
66	Tossed "good luck" coins as vectors for anthropogenic pollution into aquatic environment. <i>Environmental Pollution</i> , 2020, 259, 113800.	3.7	4
67	The ZVI-Fenton process affects the total load of human pathogenic bacteria in wastewater samples. <i>Journal of Water Process Engineering</i> , 2022, 47, 102668.	2.6	4
68	Are grazer-induced adaptations of bacterial abundance and morphology time-dependent?. <i>Journal of Limnology</i> , 2006, 65, 35.	0.3	3
69	Deconvolution models for a better understanding of natural microbial communities enumerated by flow cytometry. <i>Cytometry Part A: the Journal of the International Society for Analytical Cytology</i> , 2018, 93, 180-181.	1.1	3
70	Zooplankton as a Transitional Host for <i>Escherichia coli</i> in Freshwater. <i>Applied and Environmental Microbiology</i> , 2022, 88, e0252221.	1.4	2
71	Knowledge Gaps and Research Needs in Bacterial Co-Resistance in the Environment. , 2019, , 39-59.		1
72	Lanzarote and Chinijo Islands: An Anchialine UNESCO Global Geopark. <i>Volcanic Tourist Destinations</i> , 2019, , 109-121.	0.2	1

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73	Metagenome Analysis Reveals a Response of the Antibiotic Resistome to Mars-like Extraterrestrial Conditions. <i>Astrobiology</i> , 0, , .	1.5	1