

Jackie L Collier

List of Publications by Year in descending order

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papers

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304743

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docs citations

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4352
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#	ARTICLE	IF	CITATIONS
1	The Marine Microbial Eukaryote Transcriptome Sequencing Project (MMETSP): Illuminating the Functional Diversity of Eukaryotic Life in the Oceans through Transcriptome Sequencing. <i>PLoS Biology</i> , 2014, 12, e1001889.	5.6	885
2	Chlorosis induced by nutrient deprivation in <i>Synechococcus</i> sp. strain PCC 7942: not all bleaching is the same. <i>Journal of Bacteriology</i> , 1992, 174, 4718-4726.	2.2	280
3	Niche of harmful alga <i>Aureococcus anophagefferens</i> revealed through ecogenomics. <i>Proceedings of the National Academy of Sciences of the United States of America</i> , 2011, 108, 4352-4357.	7.1	256
4	Evolution of the Phycobiliproteins. <i>Journal of Molecular Biology</i> , 1995, 248, 79-96.	4.2	238
5	Expression of <i>nifH</i> Genes in Natural Microbial Assemblages in Lake George, New York, Detected by Reverse Transcriptase PCR. <i>Applied and Environmental Microbiology</i> , 2000, 66, 3119-3124.	3.1	235
6	Role of urea in microbial metabolism in aquatic systems: a biochemical and molecular review. <i>Aquatic Microbial Ecology</i> , 2010, 59, 67-88.	1.8	233
7	FLOW CYTOMETRY AND THE SINGLE CELL IN PHYCOLOGY. <i>Journal of Phycology</i> , 2000, 36, 628-644.	2.3	119
8	The marine cyanobacterium <i>Synechococcus</i> sp. WH7805 requires urease (urea amidohydrolase, EC 3.5.1.5) to utilize urea as a nitrogen source: molecular-genetic and biochemical analysis of the enzyme. <i>Microbiology (United Kingdom)</i> , 1999, 145, 447-459.	1.8	118
9	The Responses of Cyanobacteria to Environmental Conditions: Light and Nutrients. , 1994, , 641-675.		98
10	Genetic tool development in marine protists: emerging model organisms for experimental cell biology. <i>Nature Methods</i> , 2020, 17, 481-494.	19.0	97
11	Environmental effects on the light-harvesting complex of cyanobacteria. <i>Journal of Bacteriology</i> , 1993, 175, 575-582.	2.2	75
12	Changes in the cyanobacterial photosynthetic apparatus during acclimation to macronutrient deprivation. <i>Photosynthesis Research</i> , 1994, 42, 173-183.	2.9	71
13	Ocean urea fertilization for carbon credits poses high ecological risks. <i>Marine Pollution Bulletin</i> , 2008, 56, 1049-1056.	5.0	58
14	REVERSE TRANSCRIPTION PCR AMPLIFICATION OF CYANOBACTERIAL SYMBIONT 16S RRNA SEQUENCES FROM SINGLE NON-PHOTOSYNTHETIC EUKARYOTIC MARINE PLANKTONIC HOST CELLS1. <i>Journal of Phycology</i> , 2006, 42, 243-250.	2.3	50
15	Reconstruction and analysis of the genome-scale metabolic model of <i>schizochytrium limacinum</i> SR21 for docosahexaenoic acid production. <i>BMC Genomics</i> , 2015, 16, 799.	2.8	50
16	A calcium-binding, asparagine-linked oligosaccharide is involved in skeleton formation in the sea urchin embryo.. <i>Journal of Cell Biology</i> , 1989, 109, 1289-1299.	5.2	43
17	Novel uncultivated labyrinthulomycetes revealed by 18S rDNA sequences from seawater and sediment samples. <i>Aquatic Microbial Ecology</i> , 2010, 58, 215-228.	1.8	42
18	Phycocerythrin-containing picoplankton in the Southern California Bight. <i>Deep-Sea Research Part II: Topical Studies in Oceanography</i> , 2003, 50, 2405-2422.	1.4	35

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19	Diversity of urea-degrading microorganisms in open ocean and estuarine planktonic communities. <i>Environmental Microbiology</i> , 2009, 11, 3118-3131.	3.8	35
20	Quantitative Real-Time PCR Assay for QPX (Thraustochytriidae), a Parasite of the Hard Clam (<i> Mercenaria mercenaria</i>). <i>Overlook</i> 10, 50-70.	3.1	33
21	Alteration of plankton communities and biogeochemical cycles by harmful <i>Cochlodinium polykrikoides</i> (Dinophyceae) blooms. <i>Harmful Algae</i> , 2014, 33, 41-54.	4.8	31
22	DIFFERENCES IN GROWTH AND PHYSIOLOGY OF MARINE <i>SYNECHOCOCCUS</i> (CYANOBACTERIA) ON NITRATE VERSUS AMMONIUM ARE NOT DETERMINED SOLELY BY NITROGEN SOURCE REDOX STATE. <i>Journal of Phycology</i> , 2012, 48, 106-116.	2.3	27
23	Picoplankton contribution to biogenic silica stocks and production rates in the Sargasso Sea. <i>Global Biogeochemical Cycles</i> , 2017, 31, 762-774.	4.9	27
24	Patterns and regulation of silicon accumulation in <i>Synechococcus</i> spp.. <i>Journal of Phycology</i> , 2017, 53, 746-761.	2.3	26
25	Silicon content of individual cells of <i>Synechococcus</i> from the North Atlantic Ocean. <i>Marine Chemistry</i> , 2016, 187, 16-24.	2.3	24
26	Molecular genetic variation within and among isolates of QPX (Thraustochytridae), a parasite of the hard clam <i>Mercenaria mercenaria</i> . <i>Diseases of Aquatic Organisms</i> , 2007, 77, 159-168.	1.0	23
27	Effects of temperature on hard clam (<i>Mercenaria mercenaria</i>) immunity and QPX (Quahog Parasite). <i>Overlook</i> 10, 314-321.	3.2	22
28	UREASE GENE SEQUENCES FROM ALGAE AND HETEROTROPHIC BACTERIA IN AXENIC AND NONAXENIC PHYTOPLANKTON CULTURES. <i>Journal of Phycology</i> , 2009, 45, 625-634.	2.3	20
29	Strength in numbers: Collaborative science for new experimental model systems. <i>PLoS Biology</i> , 2018, 16, e2006333.	5.6	15
30	The chemical form of silicon in marine <i>Synechococcus</i> . <i>Marine Chemistry</i> , 2018, 206, 44-51.	2.3	14
31	A New PCR-Based Method Shows That Blue Crabs (<i>Callinectes sapidus</i> (Rathbun)) Consume Winter Flounder (<i>Pseudopleuronectes americanus</i> (Walbaum)). <i>PLoS ONE</i> , 2014, 9, e85101.	2.5	13
32	Accidental ecosystem restoration? Assessing the estuary-wide impacts of a new ocean inlet created by Hurricane Sandy. <i>Estuarine, Coastal and Shelf Science</i> , 2019, 221, 132-146.	2.1	11
33	Possible impacts of zoosporic parasites in diseases of commercially important marine mollusc species: part II. <i>Labyrinthulomycota</i> . <i>Botanica Marina</i> , 2017, 60, .	1.2	10
34	Swimming, gliding, and rolling toward the mainstream: cell biology of marine protists. <i>Molecular Biology of the Cell</i> , 2019, 30, 1245-1248.	2.1	10
35	Erection of a New Genus and Species for the Pathogen of Hard Clams ~Quahog Parasite Unknown™ (QPX): <i>Mucochytrium quahogii</i> gen. nov., sp. nov.. <i>Protist</i> , 2021, 172, 125793.	1.5	9
36	Disruption of a gene encoding a novel thioredoxin-like protein alters the cyanobacterial photosynthetic apparatus. <i>Journal of Bacteriology</i> , 1995, 177, 3269-3276.	2.2	8

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37	Microbial Communities in Partially and Fully Treated Effluent of Three Nitrogen-Removing Biofilters. <i>Journal of Sustainable Water in the Built Environment</i> , 2020, 6, 04020010.	1.6	7
38	Keeping up with advances in qPCR pathogen detection: an example for QPX disease in hard clams. <i>Diseases of Aquatic Organisms</i> , 2022, 148, 127-144.	1.0	7
39	FLOW CYTOMETRY AND THE SINGLE COMPOUND IN PLANKTON ECOLOGY. <i>Journal of Phycology</i> , 2004, 40, 805-807.	2.3	6
40	THE STRUCTURE OF PHYCOBILISOMES IN MUTANTS OF <i>Synechococcus</i> sp. STRAIN PCC 7942 DEVOID OF SPECIFIC LINKER POLYPEPTIDES. <i>Photochemistry and Photobiology</i> , 1995, 61, 298-302.	2.5	4
41	Seasonality of QPX disease in the Raritan Bay (NY) wild hard clam (<i>Mercenaria mercenaria</i>) population. <i>Aquaculture Research</i> , 2017, 48, 1269-1278.	1.8	4
42	Transcriptomic Responses of Four Pelagophytes to Nutrient (N, P) and Light Stress. <i>Frontiers in Marine Science</i> , 2021, 8, .	2.5	3
43	Evaluation of different materials used for sealing of implant abutment access channel and the peri-implant sulcus microbiota: A 6-month, randomized controlled trial. <i>Clinical Oral Implants Research</i> , 2021, 32, 941-950.	4.5	3
44	Possible impacts of zoosporic parasites in diseases of commercially important marine mollusc species: part I. Perkinsozoa. <i>Botanica Marina</i> , 2017, 60, .	1.2	1
45	Nitrogen transformations and microbial characterization in passive nitrogen removing biofilters (NRBs) for onsite wastewater treatment. <i>Proceedings of the Water Environment Federation</i> , 2017, 2017, 898-906.	0.0	1