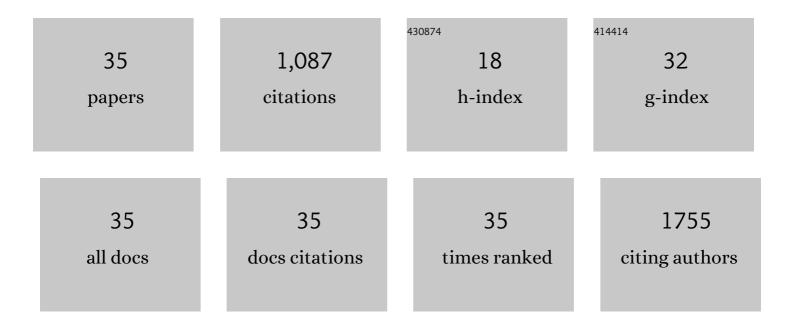
## Antonios Magoulas

List of Publications by Year in descending order

Source: https://exaly.com/author-pdf/11147234/publications.pdf Version: 2024-02-01



#	Article	IF	CITATIONS
1	Identifying Fishes through DNA Barcodes and Microarrays. PLoS ONE, 2010, 5, e12620.	2.5	145
2	Evaluation of candidate reference genes for QPCR during ontogenesis and of immune-relevant tissues of European seabass (Dicentrarchus labrax). Comparative Biochemistry and Physiology - B Biochemistry and Molecular Biology, 2009, 153, 340-347.	1.6	89
3	Quantitative Trait Loci Involved in Sex Determination and Body Growth in the Gilthead Sea Bream (Sparus aurata L.) through Targeted Genome Scan. PLoS ONE, 2011, 6, e16599.	2.5	70
4	A gene-based radiation hybrid map of the gilthead sea bream Sparus aurata refines and exploits conserved synteny with Tetraodon nigroviridis. BMC Genomics, 2007, 8, 44.	2.8	52
5	Profiling of infection specific mRNA transcripts of the European seabass Dicentrarchus labrax. BMC Genomics, 2009, 10, 157.	2.8	52
6	Kolumbo submarine volcano (Greece): An active window into the Aegean subduction system. Scientific Reports, 2016, 6, 28013.	3.3	52
7	Exploring a Nonmodel Teleost Genome Through RAD Sequencing—Linkage Mapping in Common Pandora, <i>Pagellus erythrinus</i> and Comparative Genomic Analysis. G3: Genes, Genomes, Genetics, 2016, 6, 509-519.	1.8	46
8	Mapping quantitative trait loci in European sea bass (Dicentrarchus labrax): The BASSMAP pilot study. Aquaculture, 2007, 272, S172-S182.	3.5	45
9	Origin, diversification, and historical biogeography of the genus Trachurus (Perciformes:) Tj ETQq1 1 0.784314 rg	BT /Overlc	$c_{42}$ 10 Tf 50
10	Evolution of a multifunctional gene: The warm temperature acclimation protein Wap65 in the European seabass Dicentrarchus labrax. Molecular Phylogenetics and Evolution, 2010, 55, 640-649.	2.7	40
11	Genetic population structure in the Chilean jack mackerel, Trachurus murphyi (Nichols) across the South-eastern Pacific Ocean. Fisheries Research, 2009, 100, 109-115.	1.7	39
12	Metagenomic investigation of the geologically unique <scp>H</scp> ellenic <scp>V</scp> olcanic <scp>A</scp> rc reveals a distinctive ecosystem with unexpected physiology. Environmental Microbiology, 2016, 18, 1122-1136.	3.8	37
13	Geochemistry of CO2-Rich Gases Venting From Submarine Volcanism: The Case of Kolumbo (Hellenic) Tj ETQq1 1	0,784314 1.8	rgBT /Overl
14	Population structure of anchovy Engraulis encrasicolus L. in the Mediterranean Sea inferred from multiple methods. Fisheries Research, 2008, 91, 187-195.	1.7	35
15	Comparative analysis and characterization of expressed sequence tags in gilthead sea bream (Sparus) Tj ETQq1 1	0 <sub>3</sub> 784314	rgBT /Overl
16	Second generation genetic linkage map for the gilthead sea bream Sparus aurata L. Marine Genomics, 2014, 18, 77-82.	1.1	27
17	Micro-CT for Biological and Biomedical Studies: A Comparison of Imaging Techniques. Journal of Imaging, 2021, 7, 172.	3.0	22
18	In silico mining and characterization of simple sequence repeats from gilthead sea bream (Sparus) Tj ETQq0 0 0 $r_{ m s}$	gBT /Overl 1.1	ock 10 Tf 50 21

multiplexing and cross-species assays. Marine Genomics, 2011, 4, 83-91.

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19	Microbial community differentiation between active and inactive sulfide chimneys of the Kolumbo submarine volcano, Hellenic Volcanic Arc. Extremophiles, 2018, 22, 13-27.	2.3	21
20	Os and 1s in marine molecular research: a regional HPC perspective. GigaScience, 2021, 10, .	6.4	19
21	Assessing the genetic landscape of a contact zone: the case of European hare in northeastern Greece. Genetica, 2013, 141, 23-40.	1.1	16
22	Full genome survey and dynamics of gene expression in the greater amberjack Seriola dumerili. GigaScience, 2017, 6, 1-13.	6.4	16
23	Functional and transcriptomic investigation of laccase activity in the presence of PCB29 identifies two novel enzymes and the multicopper oxidase repertoire of a marine-derived fungus. Science of the Total Environment, 2021, 775, 145818.	8.0	16
24	Simulating social-ecological systems: the Island Digital Ecosystem Avatars (IDEA) consortium. GigaScience, 2016, 5, 14.	6.4	15
25	Reduced genetic variation and strong genetic population structure in the freshwater killifish <i>Valencia letourneuxi</i> (Valenciidae) based on nuclear and mitochondrial markers. Biological Journal of the Linnean Society, 2014, 111, 334-349.	1.6	14
26	Application of Mitochondrial DNA in Stock Identification. , 2014, , 257-295.		14
27	Microbial strains isolated from CO2-venting Kolumbo submarine volcano show enhanced co-tolerance to acidity and antibiotics. Marine Environmental Research, 2019, 144, 102-110.	2.5	13
28	Functional Divergences of GAPDH Isoforms During Early Development in Two Perciform Fish Species. Marine Biotechnology, 2011, 13, 1115-1124.	2.4	11
29	Genetic diversity of Atlantic Bluefin tuna in the Mediterranean Sea: insights from genome-wide SNPs and microsatellites. Journal of Biological Research, 2017, 24, 3.	2.1	9
30	A de novo transcriptome assembly for the bath sponge Spongia officinalis, adjusting for microsymbionts. BMC Research Notes, 2019, 12, 813.	1.4	9
31	Genomic adaptation of Pseudomonas strains to acidity and antibiotics in hydrothermal vents at Kolumbo submarine volcano, Greece. Scientific Reports, 2021, 11, 1336.	3.3	9
32	Absence of spatial genetic structure in common dentex (Dentex dentex Linnaeus, 1758) in the Mediterranean Sea as evidenced by nuclear and mitochondrial molecular markers. PLoS ONE, 2018, 13, e0203866.	2.5	8
33	Genome Analysis of Lagocephalus sceleratus: Unraveling the Genomic Landscape of a Successful Invader. Frontiers in Genetics, 2021, 12, 790850.	2.3	6
34	Microbial Benthic Communities in the Aegean Sea. Handbook of Environmental Chemistry, 2020, , 1.	0.4	4
35	Characterization of polymorphic microsatellite markers in Valencia letourneuxi (Valenciidae) and cross-amplification in two other cyprinodontiform species. Conservation Genetics Resources, 2009, 1, 27-30.	0.8	3