Paola Venier

List of Publications by Year in descending order

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#	Article	IF	CITATIONS
1	A glimpse on metazoan ZNFX1 helicases, ancient players of antiviral innate immunity. Fish and Shellfish Immunology, 2022, 121, 456-466.	1.6	6
2	Paired miRNA and RNA sequencing provides a first insight into molecular defense mechanisms of Scapharca broughtonii during ostreid herpesvirus-1 infection. Fish and Shellfish Immunology, 2022, , .	1.6	1
3	ADAR-Editing during Ostreid Herpesvirus 1 Infection in <i>Crassostrea gigas</i> : Facts and Limitations. MSphere, 2022, , e0001122.	1.3	4
4	Digging into bivalve miRNAomes: between conservation and innovation. Philosophical Transactions of the Royal Society B: Biological Sciences, 2021, 376, 20200165.	1.8	16
5	NMR Metabolite Profiles of the Bivalve Mollusc Mytilus galloprovincialis Before and After Immune Stimulation With Vibrio splendidus. Frontiers in Molecular Biosciences, 2021, 8, 686770.	1.6	11
6	Massive gene presence-absence variation shapes an open pan-genome in the Mediterranean mussel. Genome Biology, 2020, 21, 275.	3.8	105
7	Parallel analysis of miRNAs and mRNAs suggests distinct regulatory networks in Crassostrea gigas infected by Ostreid herpesvirus 1. BMC Genomics, 2020, 21, 620.	1.2	12
8	Functional Insights From the Evolutionary Diversification of Big Defensins. Frontiers in Immunology, 2020, 11, 758.	2.2	35
9	Expansion and loss events characterized the occurrence of MIF-like genes in bivalves. Fish and Shellfish Immunology, 2019, 93, 39-49.	1.6	4
10	An Evolutionary Perspective of Dopachrome Tautomerase Enzymes in Metazoans. Genes, 2019, 10, 495.	1.0	6
11	A-to-I editing of Malacoherpesviridae RNAs supports the antiviral role of ADAR1 in mollusks. BMC Evolutionary Biology, 2019, 19, 149.	3.2	20
12	A Needle in A Haystack: Tracing Bivalve-Associated Viruses in High-Throughput Transcriptomic Data. Viruses, 2019, 11, 205.	1.5	28
13	Dual Analysis of Virus-Host Interactions: The Case of Ostreid herpesvirus 1 and the Cupped Oyster Crassostrea gigas. Evolutionary Bioinformatics, 2019, 15, 117693431983130.	0.6	4
14	Biotechnologies from Marine Bivalves. , 2019, , 95-112.		0
15	Induced expression of cathelicidins in trout (<scp><i>Oncorhynchus mykiss</i></scp>) challenged with four different bacterial pathogens. Journal of Peptide Science, 2018, 24, e3089.	0.8	16
16	Immunity in Molluscs: Recognition and Effector Mechanisms, with a Focus on Bivalvia. , 2018, , 225-341.		43
17	Identification of a newly described OsHV-1 Âμvar from the North Adriatic Sea (Italy). Journal of General Virology, 2018, 99, 693-703.	1.3	41
18	Diversity and evolution of TIR-domain-containing proteins in bivalves and Metazoa: New insights from comparative genomics. Developmental and Comparative Immunology, 2017, 70, 145-164.	1.0	43

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19	Structural Insights into the Mode of Action of the Peptide Antibiotic Copsin. Biochemistry, 2017, 56, 4992-5001.	1.2	8
20	Oyster RNA-seq Data Support the Development of Malacoherpesviridae Genomics. Frontiers in Microbiology, 2017, 8, 1515.	1.5	20
21	Myticalins: A Novel Multigenic Family of Linear, Cationic Antimicrobial Peptides from Marine Mussels (Mytilus spp.). Marine Drugs, 2017, 15, 261.	2.2	54
22	Immune-Related Signaling in Mussel and Bivalves. , 2016, , 93-105.		3
23	Cathelicidins. , 2016, , 225-237.		4
24	Colloidal Surface Active Maghemite Nanoparticles for Biologically Safe Cr ^{VI} Remediation: from Coreâ€ s hell Nanostructures to Pilot Plant Development. Chemistry - A European Journal, 2016, 22, 14219-14226.	1.7	16
25	Chaperone patterns in vernal keratoconjunctivitis are distinctive of cell and Hsp type and are modified by inflammatory stimuli. Allergy: European Journal of Allergy and Clinical Immunology, 2016, 71, 403-411.	2.7	8
26	Serum amyloid A in marine bivalves: An acute phase and innate immunity protein. Developmental and Comparative Immunology, 2016, 59, 136-144.	1.0	7
27	The miRNA biogenesis in marine bivalves. PeerJ, 2016, 4, e1763.	0.9	32
28	IL-17 signaling components in bivalves: Comparative sequence analysis and involvement in the immune responses. Developmental and Comparative Immunology, 2015, 52, 255-268.	1.0	41
29	An updated molecular basis for mussel immunity. Fish and Shellfish Immunology, 2015, 46, 17-38.	1.6	135
30	Dual analysis of host and pathogen transcriptomes in ostreid herpesvirus 1â€positive <scp><i>C</i></scp> <i>rassostrea gigas</i> . Environmental Microbiology, 2015, 17, 4200-4212.	1.8	75
31	Identification and Characterization of a Novel Family of Cysteine-Rich Peptides (MgCRP-I) from <i>Mytilus galloprovincialis</i> . Genome Biology and Evolution, 2015, 7, 2203-2219.	1.1	16
32	Structural and Antimicrobial Features of Peptides Related to Myticin C, a Special Defense Molecule from the Mediterranean Mussel <i>Mytilus galloprovincialis</i> . Journal of Agricultural and Food Chemistry, 2015, 63, 9251-9259.	2.4	28
33	Aquatic ecology of the oyster pathogens <scp><i>V</i></scp> <i>ibrio splendidus</i> and <scp><i>V</i></scp> <i>ibrio aestuarianus</i> . Environmental Microbiology, 2015, 17, 1065-1080.	1.8	72
34	The genome of the Pacific oyster Crassostrea gigas brings new insights on the massive expansion of the C1q gene family in Bivalvia. Developmental and Comparative Immunology, 2015, 49, 59-71.	1.0	95
35	Atmospheric-Pressure Cold Plasma Induces Transcriptional Changes in Ex Vivo Human Corneas. PLoS ONE, 2015, 10, e0133173.	1.1	21
36	Analysis of synonymous codon usage patterns in sixty-four different bivalve species. PeerJ, 2015, 3, e1520.	0.9	11

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37	Target Capture and Massive Sequencing of Genes Transcribed inMytilus galloprovincialis. BioMed Research International, 2014, 2014, 1-9.	0.9	9
38	RNA sequencing and de novo assembly of the digestive gland transcriptome in Mytilus galloprovincialis fed with toxinogenic and non-toxic strains of Alexandrium minutum. BMC Research Notes, 2014, 7, 722.	0.6	47
39	Toll signal transduction pathway in bivalves: Complete cds of intermediate elements and related gene transcription levels in hemocytes of immune stimulated Mytilus galloprovincialis. Developmental and Comparative Immunology, 2014, 45, 300-312.	1.0	61
40	Mortality occurrence and pathogen detection in Crassostrea gigas and Mytilus galloprovincialis close-growing in shallow waters (Goro lagoon, Italy). Fish and Shellfish Immunology, 2014, 41, 37-44.	1.6	79
41	Characterization of the salmonid cathelicidins and of their biological activities. Fish and Shellfish Immunology, 2013, 34, 1651.	1.6	0
42	DNA Damage and Transcriptional Changes in the Gills of Mytilus galloprovincialis Exposed to Nanomolar Doses of Combined Metal Salts (Cd, Cu, Hg). PLoS ONE, 2013, 8, e54602.	1.1	68
43	Forcing host-pathogen interactions in C. gigas and M. galloprovincialis transplanted into the goro lagoon (North Adriatic sea, Italy). Fish and Shellfish Immunology, 2013, 34, 1682.	1.6	Ο
44	First evidence of a Toll signaling pathway involved in innate immune response in Lophotrochozoa. Fish and Shellfish Immunology, 2013, 34, 1742.	1.6	0
45	Toll-like receptors and MyD88 adaptors in Mytilus: Complete cds and gene expression levels. Developmental and Comparative Immunology, 2013, 40, 158-166.	1.0	110
46	Bivalve Omics: State of the Art and Potential Applications for the Biomonitoring of Harmful Marine Compounds. Marine Drugs, 2013, 11, 4370-4389.	2.2	67
47	Big defensins and mytimacins, new AMP families of the Mediterranean mussel Mytilus galloprovincialis. Developmental and Comparative Immunology, 2012, 36, 390-399.	1.0	106
48	International roundâ€robin study on the Ames fluctuation test. Environmental and Molecular Mutagenesis, 2012, 53, 185-197.	0.9	68
49	Disinfection of Ocular Cells and Tissues by Atmospheric-Pressure Cold Plasma. PLoS ONE, 2012, 7, e33245.	1.1	97
50	Massively Parallel Amplicon Sequencing Reveals Isotype-Specific Variability of Antimicrobial Peptide Transcripts in Mytilus galloprovincialis. PLoS ONE, 2011, 6, e26680.	1.1	28
51	Diversity of Coding Sequences and Gene Structures of the Antifungal Peptide Mytimycin (MytM) from the Mediterranean Mussel, Mytilus galloprovincialis. Marine Biotechnology, 2011, 13, 857-867.	1.1	31
52	The C1q domain containing proteins of the Mediterranean mussel Mytilus galloprovincialis: A widespread and diverse family of immune-related molecules. Developmental and Comparative Immunology, 2011, 35, 635-643.	1.0	142
53	Insights into the innate immunity of the Mediterranean mussel Mytilus galloprovincialis. BMC Genomics, 2011, 12, 69.	1.2	159
54	Expression and diversity of AMP and other immune-related molecules in M. galloprovincialis. Comparative Biochemistry and Physiology Part A, Molecular & Integrative Physiology, 2010, 157, S6.	0.8	0

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55	Mediterranean Mussel Gene Expression Profile Induced by Okadaic Acid Exposure. Environmental Science & Technology, 2010, 44, 8276-8283.	4.6	45
56	Expression of Mytilus immune genes in response to experimental challenges varied according to the site of collection. Fish and Shellfish Immunology, 2010, 28, 640-648.	1.6	31
57	MytiBase: a knowledgebase of mussel (M. galloprovincialis) transcribed sequences. BMC Genomics, 2009, 10, 72.	1.2	102
58	Study of Diseases and the Immune System of Bivalves Using Molecular Biology and Genomics. Reviews in Fisheries Science, 2008, 16, 133-156.	2.1	95
59	High sequence variability of myticin transcripts in hemocytes of immune-stimulated mussels suggests ancient host–pathogen interactions. Developmental and Comparative Immunology, 2008, 32, 213-226.	1.0	83
60	Gene transcription profiling in pollutant exposed mussels (Mytilus spp.) using a new low-density oligonucleotide microarray. Gene, 2006, 376, 24-36.	1.0	102
61	Development of mussel mRNA profiling: Can gene expression trends reveal coastal water pollution?. Mutation Research - Fundamental and Molecular Mechanisms of Mutagenesis, 2006, 602, 121-134.	0.4	113
62	Characterization of mussel gill cells in vivo and in vitro. Cell and Tissue Research, 2005, 321, 131-140.	1.5	89
63	Evidence of genetic damage in grass gobies and mussels from the Venice lagoon. Environment International, 2005, 31, 1053-1064.	4.8	33
64	Towards a catalogue of genes transcribed in multiple tissues of Mytilus galloprovincialis. Gene, 2003, 314, 29-40.	1.0	65
65	Characterization of Coastal Sites by Applying Genetic and Genotoxicity Markers inMytilus GalloprovincialisandTapes Philippinarum. Chemistry and Ecology, 2003, 19, 113-128.	0.6	9
66	Evaluation of the random amplified polymorphic DNA (RAPD) assay for the detection of DNA damage and mutations. Mutation Research - Genetic Toxicology and Environmental Mutagenesis, 2002, 521, 151-163.	0.9	148
67	Susceptibility to genetic damage and cell types in Mediterranean mussels. Marine Environmental Research, 2002, 54, 487-491.	1.1	31
68	DNA adducts in mussels and fish exposed to bulky genotoxic compounds. Marine Environmental Research, 2002, 54, 481-486.	1.1	14
69	DNA ADDUCT DETECTION IN MUSSELS EXPOSED TO BULKY AROMATIC COMPOUNDS IN LABORATORY AND FIELD CONDITIONS. , 2001, , 65-83.		3
70	Enzymatic biomarker measurement and study of DNA adduct formation in benzo[a]pyrene-contaminated mussels, Mytilus galloprovincialis. Aquatic Toxicology, 2000, 49, 269-287.	1.9	188
71	Relationship Between Kinetics of Benzo[a]pyrene Bioaccumulation and DNA Binding in the Mussel Mytilus galloprovincialis. Bulletin of Environmental Contamination and Toxicology, 1999, 62, 455-462.	1.3	21
72	Tissue dose, DNA adducts, oxidative DNA damage and CYP1A-immunopositive proteins in mussels exposed to waterborne benzo[a]pyrene. Mutation Research - Fundamental and Molecular Mechanisms of Mutagenesis, 1998, 399, 17-30.	0.4	129

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73	Detection of micronuclei in gill cells and haemocytes of mussels exposed to benzo[a]pyrene. Mutation Research - Genetic Toxicology and Environmental Mutagenesis, 1997, 390, 33-44.	0.9	151
74	Metabolic activation of benzo[a]pyrene in two fetal mouse hepatocyte lines: induction of DNA adducts and micronuclei. Mutation Research - Genetic Toxicology Testing and Biomonitoring of Environmental Or Occupational Exposure, 1996, 367, 135-141.	1.2	11
75	Formation of DNA adducts in the gill tissue of Mytilus galloprovincialis treated with benzo[a]pyrene. Aquatic Toxicology, 1996, 34, 119-133.	1.9	65
76	DNA Adducts inMytilus GalloprovincialisandZosterisessor OphiocephalusCollected from PAC-Polluted and Reference Sites of the Venice Lagoon. Polycyclic Aromatic Compounds, 1996, 11, 67-73.	1.4	9
77	Mutagenic activity of chlorinated surface waters and humic acid solutions. International Journal of Environmental Health Research, 1994, 4, 17-31.	1.3	4
78	Alcohol metabolism in the upper digestive tract. European Journal of Cancer Prevention, 1992, 1, 25-32.	0.6	3
79	Evaluation of the mutagenicity of water organic extracts by the Salmonella/microsome assay. Mutation Research - Environmental Mutagenesis and Related Subjects Including Methodology, 1992, 271, 182.	0.4	0
80	Induction of gene mutations insalmonellaandDrosophilaby soluble Cr(Vi) compounds: Synergistic effects of nitrilotriacetic acid. Toxicological and Environmental Chemistry, 1989, 22, 27-38.	0.6	9
81	Release of mutagens from finished leather. Mutation Research-Fundamental and Molecular Mechanisms of Mutagenesis, 1989, 226, 229-233.	1.2	1
82	Biological monitoring of human exposure to coal tar. International Archives of Occupational and Environmental Health, 1989, 61, 363-368.	1.1	61
83	Induction of SOS response in Escherichia coli strain PQ37 by 16 chemical compounds and human urine extracts. Mutagenesis, 1989, 4, 51-57.	1.0	19
84	Validation of SOS chromotest with chemical compounds and biological samples. Mutation Research - Environmental Mutagenesis and Related Subjects Including Methodology, 1988, 203, 228.	0.4	0
85	Relationship between total urinary PAH, 1-OH pyrene levels and mutagenicity of urinary extracts from psoriatic patients exposed to therapeutical coal tar-based ointment. Mutation Research - Environmental Mutagenesis and Related Subjects Including Methodology, 1988, 203, 246.	0.4	0
86	Mutagenic activity of carbon black dyes used in the leather industry. Mutagenesis, 1987, 2, 19-22.	1.0	7
87	Interactions of chromium with nitrilotriacetic acid (NTA) in the induction of genetic effects in bacteriaâ€. Toxicological and Environmental Chemistry, 1987, 14, 201-218.	0.6	15
88	Mutagenic activity and polycyclic aromatic hydrocarbon levels in urine of humans exposed to therapeutical coal tar. Carcinogenesis, 1986, 7, 819-823.	1.3	34
89	Increased mutagenicity of chromium compounds by nitrilotriacetic acid. Environmental Mutagenesis, 1985, 7, 185-200.	1.4	31
90	Interaction of nitrilotriacetic acid with heavy metals in the induction of sister chromatid exchanges in cultured mammalian cells. Environmental Mutagenesis, 1985, 7, 381-390.	1.4	39

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91	Mutagenic activity and polycydic aromatic hydrocarbon levels in urine of workers exposed to coal tar pitch volatiles in an anode plant. Carcinogenesis, 1985, 6, 749-752.	1.3	35
92	Genetic effects of chromium tannins. Carcinogenesis, 1985, 6, 1327-1335.	1.3	17
93	Effects of nitrilotriacetic acid on the induction of gene mutations and sister-chromatid exchanges by insoluble chromium compounds. Mutation Research - Genetic Toxicology Testing and Biomonitoring of Environmental Or Occupational Exposure, 1985, 156, 219-228.	1.2	35
94	Mutagenic Activity of Nitriloacetic Acid. Archives of Toxicology Supplement, 1984, , 407-407.	0.7	0
95	Mutagenesis test on urine of workers exposed to polycyclic aromatic hydrocarbons in a anode plant. Medicina Del Lavoro, 1984, 75, 275-81.	0.3	4
96	Cytotoxic, mutagenic and clastogenic activity of industrial chromium compounds. Mutation Research - Environmental Mutagenesis and Related Subjects Including Methodology, 1983, 113, 316-317.	0.4	1
97	Genetic effects of chromium compounds. Mutation Research - Genetic Toxicology Testing and Biomonitoring of Environmental Or Occupational Exposure, 1983, 117, 279-300.	1.2	136
98	Cytotoxic, mutagenic and clastogenic effects of industrial chromium compounds. Carcinogenesis, 1982, 3, 1331-1338.	1.3	47