Sylvie Lapegue

List of Publications by Year in descending order

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#	Article	IF	CITATIONS
1	Mortality of marine mussels <i>Mytilus edulis</i> and <i>M.Âgalloprovincialis</i> : systematic literature review of risk factors and recommendations for future research. Reviews in Aquaculture, 2021, 13, 504-536.	4.6	18
2	Genetic Characterization of Cupped Oyster Resources in Europe Using Informative Single Nucleotide Polymorphism (SNP) Panels. Genes, 2020, 11, 451.	1.0	4
3	Analysis of Genome-Wide Differentiation between Native and Introduced Populations of the Cupped Oysters Crassostrea gigas and Crassostrea angulata. Genome Biology and Evolution, 2018, 10, 2518-2534.	1.1	52
4	Contribution of in Vivo Experimental Challenges to Understanding Flat Oyster Ostrea edulis Resistance to Bonamia ostreae. Frontiers in Cellular and Infection Microbiology, 2017, 7, 433.	1.8	12
5	Rapid expansion of the invasive oyster Crassostrea gigas at its northern distribution limit in Europe: Naturally dispersed or introduced?. PLoS ONE, 2017, 12, e0177481.	1.1	29
6	Can survival of European flat oysters following experimental infection with Bonamia ostreae be predicted using QTLs?. Aquaculture, 2015, 448, 521-530.	1.7	17
7	Additive transcriptomic variation associated with reproductive traits suggest local adaptation in a recently settled population of the Pacific oyster, Crassostrea gigas. BMC Genomics, 2015, 16, 808.	1.2	15
8	Invasion genetics of the Pacific oyster Crassostrea gigas in the British Isles inferred from microsatellite and mitochondrial markers. Biological Invasions, 2015, 17, 2581-2595.	1.2	38
9	Genotyping of a microsatellite locus to differentiate clinical Ostreid herpesvirus 1 specimens. Veterinary Research, 2014, 45, 3.	1.1	31
10	A High Load of Non-neutral Amino-Acid Polymorphisms Explains High Protein Diversity Despite Moderate Effective Population Size in a Marine Bivalve With Sweepstakes Reproduction. G3: Genes, Genomes, Genetics, 2013, 3, 333-341.	0.8	31
11	Multiplex PCR sets of novel microsatellite loci for the great scallop <i>Pecten maximus</i> and their application in parentage assignment. Aquatic Living Resources, 2013, 26, 207-213.	0.5	19
12	Population genomics shed light on the demographic and adaptive histories of European invasion in the Pacific oyster, <i>Crassostrea gigas</i> . Evolutionary Applications, 2013, 6, 1064-1078.	1.5	51
13	Gonad volume assessment in the oyster Crassostrea gigas: Comparison between a histological method and a magnetic resonance imaging (MRI) method. Aquaculture, 2012, 370-371, 84-89.	1.7	5
14	Complete mitochondrial DNA sequence of the European flat oyster Ostrea edulis confirms Ostreidae classification. BMC Research Notes, 2011, 4, 400.	0.6	32
15	Strategies for the retention of high genetic variability in European flat oyster (Ostrea edulis) restoration programmes. Conservation Genetics, 2010, 11, 1899-1910.	0.8	63
16	Genomic Approaches in Aquaculture and Fisheries. , 2010, , 213-286.		5
17	Development of four EST-SSR multiplex PCRs in the Pacific oyster (Crassostrea gigas) and their validation in parentage assignment. Aquaculture, 2010, 310, 234-239.	1.7	28
18	Generation and analysis of a 29,745 unique Expressed Sequence Tags from the Pacific oyster (Crassostrea gigas) assembled into a publicly accessible database: the GigasDatabase. BMC Genomics, 2009, 10, 341.	1.2	127

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19	Detection of phenoloxidase activity in early stages of the Pacific oyster Crassostrea gigas (Thunberg). Developmental and Comparative Immunology, 2009, 33, 653-659.	1.0	32
20	Ostreid herpes virus 1 infection in families of the Pacific oyster, Crassostrea gigas, during a summer mortality outbreak: Differences in viral DNA detection and quantification using real-time PCR. Virus Research, 2009, 142, 181-187.	1.1	106
21	Characterization of 27 microsatellite loci in the European flat oyster <i>Ostrea edulis</i> . Molecular Ecology Resources, 2009, 9, 960-963.	2.2	14
22	Increasing genomic information in bivalves through new EST collections in four species: Development of new genetic markers for environmental studies and genome evolution. Gene, 2008, 408, 27-36.	1.0	132
23	<i>Bonamia ostreae</i> -induced mortalities in one-year old European flat oysters <i>Ostrea edulis</i> : experimental infection by cohabitation challenge. Aquatic Living Resources, 2008, 21, 423-439.	0.5	37
24	Impact of Diuron on Aneuploidy and Hemocyte Parameters in Pacific Oyster, Crassostrea gigas. Archives of Environmental Contamination and Toxicology, 2007, 52, 58-63.	2.1	48
25	Effects of cadmium on aneuploidy and hemocyte parameters in the Pacific oyster, Crassostrea gigas. Aquatic Toxicology, 2006, 78, 149-156.	1.9	38
26	Phylogeographic study of the dwarf oyster, Ostreola stentina, from Morocco, Portugal and Tunisia: evidence of a geographic disjunction with the closely related taxa, Ostrea aupouria and Ostreola equestris. Marine Biology, 2006, 150, 103-110.	0.7	24
27	GENETIC DIVERSITY OF THE EUROPEAN OYSTER (OSTREA EDULIS L.) IN NOVA SCOTIA: COMPARISON WITH OTHER PARTS OF CANADA, MAINE AND EUROPE AND IMPLICATIONS FOR BROODSTOCK MANAGEMENT. Journal of Shellfish Research, 2006, 25, 543-551.	0.3	12
28	A Complementary Method for Production of Tetraploid Crassostrea gigas Using Crosses Between Diploids and Tetraploids with Cytochalasin B Treatments. Marine Biotechnology, 2005, 7, 318-330.	1.1	31
29	Endonuclease banding reveals that atrazine-induced aneuploidy resembles spontaneous chromosome loss in Crassostrea gigas. Genome, 2005, 48, 177-180.	0.9	14
30	Chromosome loss in bi-parental progenies of tetraploid Pacific oyster Crassostrea gigas. Aquaculture, 2005, 247, 97-105.	1.7	39
31	Impact of atrazine on aneuploidy in pacific oysters, <i>Crassostrea gigas</i> . Environmental Toxicology and Chemistry, 2003, 22, 219-223.	2.2	24
32	Mitochondrial and nuclear DNA sequence variation of presumed Crassostrea gigas and Crassostrea angulata specimens: a new oyster species in Hong Kong?. Aquaculture, 2003, 228, 15-25.	1.7	67
33	IMPACT OF ATRAZINE ON ANEUPLOIDY IN PACIFIC OYSTERS, CRASSOSTREA GIGAS. Environmental Toxicology and Chemistry, 2003, 22, 219.	2.2	2
34	Genetic parallelism between European flat oyster populations at the edge of their natural range. Evolutionary Applications, 0, , .	1.5	2