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List of Publications by Year
in descending order

86 papers	5,380 citations	87843 38 h-index	95218 68 g-index
96 all docs	96 docs citations	96 times ranked	3113 citing authors

#	Article	IF	Citations
1	Harnessing genomics to fast-track genetic improvement in aquaculture. Nature Reviews Genetics, 2020, 21, 389-409.	7.7	286
2	Major Quantitative Trait Loci Affect Resistance to Infectious Pancreatic Necrosis in Atlantic Salmon (Salmo salar). Genetics, 2008, 178, 1109-1115.	1.2	262
3	Development and validation of a high density SNP genotyping array for Atlantic salmon (Salmo salar). BMC Genomics, 2014, 15, 90.	1.2	219
4	Applications of genotyping by sequencing in aquaculture breeding and genetics. Reviews in Aquaculture, 2018, 10, 670-682.	4.6	217
5	Genome wide association and genomic prediction for growth traits in juvenile farmed Atlantic salmon using a high density SNP array. BMC Genomics, 2015, 16, 969.	1.2	211
6	Genomic prediction of host resistance to sea lice in farmed Atlantic salmon populations. Genetics Selection Evolution, 2016, 48, 47.	1.2	203
7	Linkage maps of the Atlantic salmon (Salmo salar) genome derived from RAD sequencing. BMC Genomics, 2014, 15, 166.	1.2	151
8	Genome-Wide Association and Genomic Selection for Resistance to Amoebic Gill Disease in Atlantic Salmon. G3: Genes, Genomes, Genetics, 2018, 8, 1195-1203.	0.8	142
9	The susceptibility of Atlantic salmon fry to freshwater infectious pancreatic necrosis is largely explained by a major QTL. Heredity, 2010, 105, 318-327.	1.2	139
10	Genomic Prediction of Resistance to Pasteurellosis in Gilthead Sea Bream (Sparus aurata) Using 2b-RAD Sequencing. G3: Genes, Genomes, Genetics, 2016, 6, 3693-3700.	0.8	129
11	Potential of Genome Editing to Improve Aquaculture Breeding and Production. Trends in Genetics, 2019, 35, 672-684.	2.9	125
12	Characterisation of QTL-linked and genome-wide restriction site-associated DNA (RAD) markers in farmed Atlantic salmon. BMC Genomics, 2012, 13, 244.	1.2	120
13	Genetics and genomics of disease resistance in salmonid species. Frontiers in Genetics, 2014, 5, 415.	1.1	120
14	Potential of genotyping-by-sequencing for genomic selection in livestock populations. Genetics Selection Evolution, 2015, 47, 12.	1.2	107
15	Genomic Selection for Growth Traits in Pacific Oyster (Crassostrea gigas): Potential of Low-Density Marker Panels for Breeding Value Prediction. Frontiers in Genetics, 2018, 9, 391.	1.1	105
16	Future directions in breeding for disease resistance in aquaculture species. Revista Brasileira De Zootecnia, 2017, 46, 545-551.	0.3	104
17	Functional Annotation of All Salmonid Genomes (FAASG): an international initiative supporting future salmonid research, conservation and aquaculture. BMC Genomics, 2017, 18, 484.	1.2	99
18	Development of a Medium Density Combined-Species SNP Array for Pacific and European Oysters (Crassostrea gigas and Ostrea edulis). G3: Genes, Genomes, Genetics, 2017, 7, 2209-2218.	0.8	97

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19	Genotype Imputation To Improve the Cost-Efficiency of Genomic Selection in Farmed Atlantic Salmon. G3: Genes, Genomes, Genetics, 2017, 7, 1377-1383.	0.8	93
20	A chromosome-level genome assembly for the Pacific oyster Crassostrea gigas. GigaScience, 2021, 10, .	3.3	88
21	Genome-wide association and genomic prediction of resistance to viral nervous necrosis in European sea bass (Dicentrarchus labrax) using RAD sequencing. Genetics Selection Evolution, 2018, 50, 30.	1.2	87
22	Accuracy of Genomic Evaluations of Juvenile Growth Rate in Common Carp (Cyprinus carpio) Using Genotyping by Sequencing. Frontiers in Genetics, 2018, 9, 82.	1.1	85
23	Genomics in aquaculture to better understand species biology and accelerate genetic progress. Frontiers in Genetics, 2015, 6, 128.	1.1	82
24	Amelanocortin-4 receptor(MC4R) polymorphism is associated with performance traits in divergently selected large white pig populations. Animal Genetics, 2004, 35, 386-390.	0.6	79
25	Gene expression comparison of resistant and susceptible Atlantic salmon fry challenged with Infectious Pancreatic Necrosis virus reveals a marked contrast in immune response. BMC Genomics, 2016, 17, 279.	1.2	78
26	The genetic architecture of growth and fillet traits in farmed Atlantic salmon (Salmo salar). BMC Genetics, 2015, 16, 51.	2.7	77
27	Mapping and validation of a major QTL affecting resistance to pancreas disease (salmonid alphavirus) in Atlantic salmon (Salmo salar). Heredity, 2015, 115, 405-414.	1.2	77
28	Atlantic salmon (Salmo salar L.) genetics in the 21st century: taking leaps forward in aquaculture and biological understanding. Animal Genetics, 2019, 50, 3-14.	0.6	66
29	A Genome-Wide Association Study for Host Resistance to Ostreid Herpesvirus in Pacific Oysters (Crassostrea gigas). G3: Genes, Genomes, Genetics, 2018, 8, 1273-1280.	0.8	63
30	Optimizing Low-Cost Genotyping and Imputation Strategies for Genomic Selection in Atlantic Salmon. G3: Genes, Genomes, Genetics, 2020, 10, 581-590.	0.8	61
31	Genomic Prediction Using Low Density Marker Panels in Aquaculture: Performance Across Species, Traits, and Genotyping Platforms. Frontiers in Genetics, 2020, 11, 124.	1.1	61
32	Discovery and Functional Annotation of Quantitative Trait Loci Affecting Resistance to Sea Lice in Atlantic Salmon. Frontiers in Genetics, 2019, 10, 56.	1.1	59
33	Accuracy of genotype imputation and genomic predictions in a two-generation farmed Atlantic salmon population using high-density and low-density SNP panels. Aquaculture, 2018, 491, 147-154.	1.7	56
34	Gene Expression Response to Sea Lice in Atlantic Salmon Skin: RNA Sequencing Comparison Between Resistant and Susceptible Animals. Frontiers in Genetics, 2018, 9, 287.	1.1	50
35	Optimizing Genomic Prediction of Host Resistance to Koi Herpesvirus Disease in Carp. Frontiers in Genetics, 2019, 10, 543.	1.1	48
36	Genetic differences in host infectivity affect disease spread and survival in epidemics. Scientific Reports, 2019, 9, 4924.	1.6	48

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37	Single nucleotide polymorphisms in the insulinâ€like growth factor 1 (<scp>IGF</scp>1) gene are associated with growthâ€related traits in farmed Atlantic salmon. Animal Genetics, 2014, 45, 709-715.	0.6	46
38	QTL affecting morphometric traits and stress response in the gilthead seabream (Sparus aurata). Aquaculture, 2011, 319, 58-66.	1.7	42
39	Potential of genomic selection for improvement of resistance to ostreid herpesvirus in Pacific oyster (Crassostrea gigas). Animal Genetics, 2020, 51, 249-257.	0.6	41
40	Construction and Annotation of a High Density SNP Linkage Map of the Atlantic Salmon (Salmo) Tj ETQq0 0 0 rgBT /Overlock 10 Tf 5	0.8	40
41	Highâ€resolution mapping of the recombination landscape of the phytopathogen Fusarium graminearum suggests twoâ€speed genome evolution. Molecular Plant Pathology, 2018, 19, 341-354.	2.0	40
42	Detailed insights into panâ€European population structure and inbreeding in wild and hatchery Pacific oysters (Crassostrea gigas) revealed by genomeâ€wide SNP data. Evolutionary Applications, 2019, 12, 519-534.	1.5	39
43	Efficient CRISPR/Cas9 genome editing in a salmonid fish cell line using a lentivirus delivery system. BMC Biotechnology, 2020, 20, 35.	1.7	39
44	Genomics Toolbox for Farmed Fish. Reviews in Fisheries Science, 2008, 16, 3-15.	2.1	38
45	Mapping and Sequencing of a Significant Quantitative Trait Locus Affecting Resistance to Koi Herpesvirus in Common Carp. G3: Genes, Genomes, Genetics, 2018, 8, 3507-3513.	0.8	38
46	Current status and potential of genomic selection to improve selective breeding in the main aquaculture species of International Council for the Exploration of the Sea (ICES) member countries. Aquaculture Reports, 2021, 20, 100700.	0.7	37
47	Verification of SNPs Associated with Growth Traits in Two Populations of Farmed Atlantic Salmon. International Journal of Molecular Sciences, 2016, 17, 5.	1.8	36
48	Genetic parameters for resistance to Tilapia Lake Virus (TiLV) in Nile tilapia (Oreochromis niloticus). Aquaculture, 2020, 522, 735126.	1.7	36
49	Segregation of infectious pancreatic necrosis resistance QTL in the early life cycle of Atlantic Salmon (Salmo salar). Animal Genetics, 2010, 41, 531-536.	0.6	34
50	Detection of QTL affecting harvest traits in a commercial Atlantic salmon population. Animal Genetics, 2009, 40, 753-755.	0.6	32
51	A QTL affecting daily feed intake maps to Chromosome 2 in pigs. Mammalian Genome, 2005, 16, 464-470.	1.0	31
52	Balancing selection at a premature stop mutation in the myostatin gene underlies a recessive leg weakness syndrome in pigs. PLoS Genetics, 2019, 15, e1007759.	1.5	31
53	Population Structure and Genetic Diversity of Nile Tilapia (Oreochromis niloticus) Strains Cultured in Tanzania. Frontiers in Genetics, 2019, 10, 1269.	1.1	31
54	Development and Validation of an Open Access SNP Array for Nile Tilapia (Oreochromis) Tj ETQq0 0 0 rgBT /Overlock 10 Tf 50 62 Td	0.8	31

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55	Genetic improvement technologies to support the sustainable growth of UK aquaculture. Reviews in Aquaculture, 2021, 13, 1958-1985.	4.6	31
56	Development and testing of a combined species SNP array for the European seabass (Dicentrarchus) Tj ETQq0 0 0 rgBT /Overlock 10 Tf 5	E.3	31
57	Exploring the utility of cross-laboratory RAD-sequencing datasets for phylogenetic analysis. BMC Research Notes, 2015, 8, 299.	0.6	29
58	A major quantitative trait locus affecting resistance to Tilapia lake virus in farmed Nile tilapia (Oreochromis niloticus). Heredity, 2021, 127, 334-343.	1.2	29
59	Detection and Confirmation of a Major QTL Affecting Resistance to Infectious Pancreatic Necrosis (IPN) in Atlantic Salmon (Salmo Salar). Developments in Biologicals, 2008, 132, 199-204.	0.4	29
60	Sequencing and Characterisation of an Extensive Atlantic Salmon (Salmo salar L.) MicroRNA Repertoire. PLoS ONE, 2013, 8, e70136.	1.1	29
61	miRNAs Predicted to Regulate Host Anti-viral Gene Pathways in IPNV-Challenged Atlantic Salmon Fry Are Affected by Viral Load, and Associated With the Major IPN Resistance QTL Genotypes in Late Infection. Frontiers in Immunology, 2020, 11, 2113.	2.2	28
62	Optimizing hatchery practices for genetic improvement of marine bivalves. Reviews in Aquaculture, 2021, 13, 2289-2304.	4.6	28
63	Surrogate broodstock to enhance biotechnology research and applications in aquaculture. Biotechnology Advances, 2021, 49, 107756.	6.0	28
64	A SNP in the 5â€² flanking region of the myostatin-1b gene is associated with harvest traits in Atlantic salmon (Salmo salar). BMC Genetics, 2013, 14, 112.	2.7	27
65	Maternal inheritance of deltamethrin resistance in the salmon louse Lepeophtheirus salmonis (KrÃ¸yer) is associated with unique mtDNA haplotypes. PLoS ONE, 2017, 12, e0180625.	1.1	27
66	SNP markers for the genetic characterization of Mexican shrimp broodstocks. Genomics, 2018, 110, 423-429.	1.3	26
67	Characterising the mechanisms underlying genetic resistance to amoebic gill disease in Atlantic salmon using RNA sequencing. BMC Genomics, 2020, 21, 271.	1.2	23
68	The nedd-8 activating enzyme gene underlies genetic resistance to infectious pancreatic necrosis virus in Atlantic salmon. Genomics, 2021, 113, 3842-3850.	1.3	22
69	A Polymorphism in the 5â€²-Untranslated Region of the Porcine Cholecystokinin Type A Receptor Gene Affects Feed Intake and Growth. Genetics, 2006, 174, 1555-1563.	1.2	21
70	Potential of genomic technologies to improve disease resistance in molluscan aquaculture. Philosophical Transactions of the Royal Society B: Biological Sciences, 2021, 376, 20200168.	1.8	18
71	Quantitative trait loci and genes associated with salmonid alphavirus load in Atlantic salmon: implications for pancreas disease resistance and tolerance. Scientific Reports, 2020, 10, 10393.	1.6	17
72	Characterizing the genetic structure of introduced Nile tilapia (Oreochromis niloticus) strains in Tanzania using double digest RAD sequencing. Aquaculture International, 2020, 28, 477-492.	1.1	16

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73	Characterization of OAR1 and OAR18 QTL associated with muscle depth in British commercial terminal sire sheep. Animal Genetics, 2011, 42, 172-180.	0.6	15
74	Efficient Genome Editing in Multiple Salmonid Cell Lines Using Ribonucleoprotein Complexes. Marine Biotechnology, 2020, 22, 717-724.	1.1	15
75	Changed Patterns of Genomic Variation Following Recent Domestication: Selection Sweeps in Farmed Atlantic Salmon. Frontiers in Genetics, 2020, 11, 264.	1.1	15
76	Investigating mechanisms underlying genetic resistance to Salmon Rickettsial Syndrome in Atlantic salmon using RNA sequencing. BMC Genomics, 2021, 22, 156.	1.2	15
77	Assessing the genetic diversity of farmed and wild Rufiji tilapia (Oreochromis urolepis) Tj ETQq1 1 0.784314 rgBT /Overlock 10 Tf 50	0.8	12
78	Developments in marine invertebrate primary culture reveal novel cell morphologies in the model bivalve Crassostrea gigas. PeerJ, 2020, 8, e9180.	0.9	12
79	The impact of genetic relationship between training and validation populations on genomic prediction accuracy in Atlantic salmon. Aquaculture Reports, 2022, 23, 101033.	0.7	12
80	Exploring genetic resistance to infectious salmon anaemia virus in Atlantic salmon by genome-wide association and RNA sequencing. BMC Genomics, 2021, 22, 345.	1.2	11
81	The cholecystokinin type A receptor g.179A>G polymorphism affects feeding rate. Animal Genetics, 2008, 39, 187-188.	0.6	10
82	Assessment of genetic diversity and population structure in cultured Australian Pacific oysters. Animal Genetics, 2019, 50, 686-694.	0.6	9
83	Novel insights into the genetic relationship between growth and disease resistance in an aquaculture strain of Coho salmon (Oncorhynchus kisutch). Aquaculture, 2019, 511, 734207.	1.7	9
84	Transcriptome Profiling of Pacu (Piaractus mesopotamicus) Challenged With Pathogenic Aeromonas hydrophila: Inference on Immune Gene Response. Frontiers in Genetics, 2020, 11, 604.	1.1	8
85	The role of energy reserves in common carp performance inferred from phenotypic and genetic parameters. Aquaculture, 2021, 541, 736799.	1.7	5
86	Genetic relationship between koi herpesvirus disease resistance and production traits inferred from sibling performance in Amur mirror carp. Aquaculture, 2020, 520, 734986.	1.7	4