Michael Snow

List of Publications by Year in descending order

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#	Article	IF	CITATIONS
1	Analysis of the nucleoprotein gene identifies distinct lineages of viral haemorrhagic septicaemia virus within the European marine environment. Virus Research, 1999, 63, 35-44.	2.2	119
2	Selection for upper thermal tolerance in rainbow trout (<i>Oncorhynchus mykiss</i> Walbaum). Journal of Experimental Biology, 2015, 218, 803-812.	1.7	110
3	A low-pathogenic variant of infectious salmon anemia virus (ISAV-HPRO) is highly prevalent and causes a non-clinical transient infection in farmed Atlantic salmon (Salmo salar L.) in the Faroe Islands. Journal of General Virology, 2011, 92, 909-918.	2.9	82
4	Characterization of the Infectious Salmon Anemia Virus Fusion Protein. Journal of Virology, 2005, 79, 12544-12553.	3.4	78
5	Characterization of the Infectious Salmon Anemia Virus Genomic Segment That Encodes the Putative Hemagglutinin. Journal of Virology, 2001, 75, 5352-5356.	3.4	76
6	Identification of an interferon antagonist protein encoded by segment 7 of infectious salmon anaemia virus. Virus Research, 2006, 115, 176-184.	2.2	68
7	Three species of Mytilus and their hybrids identified in a Scottish Loch: natives, relicts and invaders?. Journal of Experimental Marine Biology and Ecology, 2008, 367, 100-110.	1.5	54
8	Development and application of real-time PCR for specific detection of Lepeophtheirus salmonis and Caligus elongatus larvae in Scottish plankton samples. Diseases of Aquatic Organisms, 2006, 73, 141-150.	1.0	51
9	Surveillance for infectious salmon anaemia virus HPRO in marine Atlantic salmon farms across Scotland. Diseases of Aquatic Organisms, 2009, 87, 161-169.	1.0	45
10	Development of a sensitive and controlled real-time RT-PCR assay for viral haemorrhagic septicaemia virus (VHSV) in marine salmonid aquaculture. Diseases of Aquatic Organisms, 2008, 80, 137-144.	1.0	38
11	Characterisation of the putative nucleoprotein gene of infectious salmon anaemia virus (ISAV). Virus Research, 2001, 74, 111-118.	2.2	31
12	Identification and characterisation of the genomic segment 7 of the infectious salmon anaemia virus genome. Virus Research, 2002, 84, 161-170.	2.2	27
13	Survey of mussel (<i>Mytilus)</i> species at Scottish shellfish farms. Aquaculture Research, 2009, 40, 1715-1722.	1.8	24
14	The contribution of molecular epidemiology to the understanding and control of viral diseases of salmonid aquaculture. Veterinary Research, 2011, 42, 56.	3.0	24
15	Realâ€time <scp>PCR</scp> detection of <i>Didemnum perlucidum</i> (Monniot, 1983) and <i>Didemnum vexillum</i> (Kott, 2002) in an applied routine marine biosecurity context. Molecular Ecology Resources, 2017, 17, 443-453.	4.8	22
16	An unusually high upper thermal acclimation potential for rainbow trout. , 2022, 10, coab101.		22
17	A strand specific real-time RT-PCR method for the targeted detection of the three species (vRNA, cRNA) Tj ETQq1 2013, 187, 65-71.	1 0.7843 2.1	14 rgBT /Ov 16
18	Investigating the cryptogenic status of the sea squirt Didemnum perlucidum (Tunicata, Ascidiacea) in Australia based on a molecular study of its global distribution. Aquatic Invasions, 2016, 11, 239-245.	1.6	16

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19	Gametogenic asynchrony of mussels Mytilus in a mixed-species area: Implications for management. Aquaculture, 2009, 295, 175-182.	3.5	14
20	Development of sensitive and specific molecular tools for the efficient detection and discrimination of potentially invasive mussel species of the genus Perna. Management of Biological Invasions, 2013, 4, 155-165.	1.2	10
21	The convict cichlid Amatitlania nigrofasciata (Cichlidae): first record of this non-native species in Western Australian waterbodies. Records of the Western Australian Museum, 2013, 28, 7.	0.8	9
22	Molecular identification of the precise geographic origins of an invasive shrimp species in a globally significant Australian biodiversity hotspot. Biological Invasions, 2017, 19, 463-468.	2.4	8
23	Performance of Mussels, <i>Mytilus edulis</i> , <i>Mytilus trossulus,</i> and Their Hybrids in Cultivation at Three Scottish Lochs. Journal of the World Aquaculture Society, 2011, 42, 111-121.	2.4	7
24	Genotype-specific Taqman® assays for the detection and rapid characterisation of European strains of viral haemorrhagic septicaemia virus. Journal of Virological Methods, 2013, 187, 209-214.	2.1	7
25	Isolation and characterization of 13 polymorphic microsatellite loci for the smooth Cherax cainii and hairy marron C. tenuimanus (Decapoda: Parastacidae). Conservation Genetics Resources, 2014, 6, 337-339.	0.8	4
26	Introgression Threatens the Survival of the Critically Endangered Freshwater Crayfish Cherax tenuimanus (Decapoda: Parastacidae) in the Wild. PLoS ONE, 2015, 10, e0121075.	2.5	2