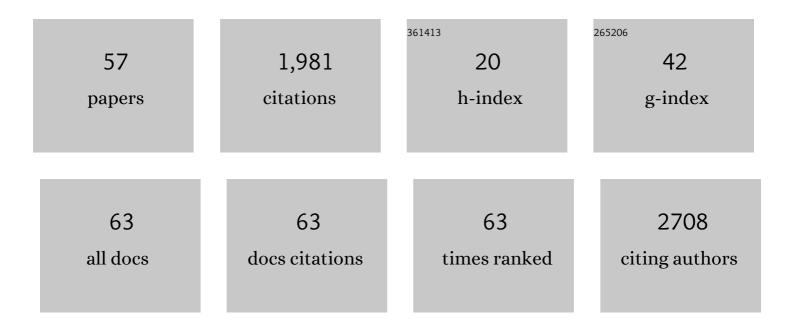
## James W Wynne

List of Publications by Year in descending order

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#	Article	IF	CITATIONS
1	Comparative Analysis of Bat Genomes Provides Insight into the Evolution of Flight and Immunity. Science, 2013, 339, 456-460.	12.6	522
2	Contraction of the type I IFN locus and unusual constitutive expression of <i>IFN-α</i> in bats. Proceedings of the National Academy of Sciences of the United States of America, 2016, 113, 2696-2701.	7.1	272
3	A microbial sea of possibilities: current knowledge and prospects for an improved understanding of the fish microbiome. Reviews in Aquaculture, 2020, 12, 1101-1134.	9.0	117
4	Transcriptome Analyses of Amoebic Gill Disease-affected Atlantic Salmon (Salmo salar) Tissues Reveal Localized Host Gene Suppression. Marine Biotechnology, 2008, 10, 388-403.	2.4	83
5	Bats and Viruses: Friend or Foe?. PLoS Pathogens, 2013, 9, e1003651.	4.7	65
6	Major histocompatibility polymorphism associated with resistance towards amoebic gill disease in Atlantic salmon (Salmo salar L.). Fish and Shellfish Immunology, 2007, 22, 707-717.	3.6	60
7	Resequencing the <i>Mycobacterium avium</i> subsp. <i>paratuberculosis</i> K10 Genome: Improved Annotation and Revised Genome Sequence. Journal of Bacteriology, 2010, 192, 6319-6320.	2.2	56
8	Evolution and comparative analysis of the bat MHC-I region. Scientific Reports, 2016, 6, 21256.	3.3	56
9	Exploring the Zoonotic Potential of Mycobacterium avium Subspecies paratuberculosis through Comparative Genomics. PLoS ONE, 2011, 6, e22171.	2.5	55
10	A diversity of amoebae colonise the gills of farmed Atlantic salmon (Salmo salar) with amoebic gill disease (AGD). European Journal of Protistology, 2019, 67, 27-45.	1.5	49
11	Genetic variation of resistance to amoebic gill disease in Atlantic salmon (Salmo salar) assessed in a challenge system. Aquaculture, 2007, 272, S94-S99.	3.5	46
12	Resistance to amoebic gill disease (AGD) is characterised by the transcriptional dysregulation of immune and cell cycle pathways. Developmental and Comparative Immunology, 2008, 32, 1539-1560.	2.3	46
13	Gene expression analysis of whole blood RNA from pigs infected with low and high pathogenic African swine fever viruses. Scientific Reports, 2017, 7, 10115.	3.3	45
14	Proteomics informed by transcriptomics reveals Hendra virus sensitizes bat cells to TRAIL-mediated apoptosis. Genome Biology, 2014, 15, 532.	8.8	42
15	Characterization of the Antigen Processing Machinery and Endogenous Peptide Presentation of a Bat MHC Class I Molecule. Journal of Immunology, 2016, 196, 4468-4476.	0.8	30
16	Microbial biomass, marine invertebrate meals and feed restriction influence the biological and gut microbiota response of shrimp Penaeus monodon. Aquaculture, 2020, 520, 734679.	3.5	30
17	Proteomics informed by transcriptomics reveals Hendra virus sensitizes bat cells to TRAIL mediated apoptosis. Genome Biology, 2014, 15, 532.	9.6	30
18	Comparative Transcriptomics Highlights the Role of the Activator Protein 1 Transcription Factor in the Host Response to Ebolavirus. Journal of Virology, 2017, 91, .	3.4	27

#	Article	IF	CITATIONS
	Purification and Characterisation of Immunoglobulins from the Australian Black Flying Fox (Pteropus) Tj ETQq1 1		<u> </u>
19	e52930.	2.5	26
20	Exposure Risk for Infection and Lack of Human-to-Human Transmission of <i>Mycobacterium ulcerans</i> Disease, Australia. Emerging Infectious Diseases, 2017, 23, 837-840.	4.3	24
21	Antibiotic-induced alterations and repopulation dynamics of yellowtail kingfish microbiota. Animal Microbiome, 2020, 2, 26.	3.8	23
22	Bacteriomic Profiling of Branchial Lesions Induced by Neoparamoeba perurans Challenge Reveals Commensal Dysbiosis and an Association with Tenacibaculum dicentrarchi in AGD-Affected Atlantic Salmon (Salmo salar L.). Microorganisms, 2020, 8, 1189.	3.6	22
23	Investigating Both Mucosal Immunity and Microbiota in Response to Gut Enteritis in Yellowtail Kingfish. Microorganisms, 2020, 8, 1267.	3.6	22
24	Host-Parasite Interaction of Atlantic salmon (Salmo salar) and the Ectoparasite Neoparamoeba perurans in Amoebic Gill Disease. Frontiers in Immunology, 2021, 12, 672700.	4.8	22
25	The interaction between temperature and dose on the efficacy and biochemical response of Atlantic salmon to hydrogen peroxide treatment for amoebic gill disease. Journal of Fish Diseases, 2020, 43, 39-48.	1.9	17
26	Production and proteomic characterisation of purified protein derivative from Mycobacterium avium subsp. paratuberculosis. Proteome Science, 2012, 10, 22.	1.7	15
27	The Effect of Antimicrobial Treatment upon the Gill Bacteriome of Atlantic Salmon (Salmo salar L.) and Progression of Amoebic Gill Disease (AGD) In Vivo. Microorganisms, 2021, 9, 987.	3.6	11
28	Transcriptome Response of Atlantic Salmon (Salmo salar) to a New Piscine Orthomyxovirus. Pathogens, 2020, 9, 807.	2.8	10
29	Gill Mucus and Gill Mucin O-glycosylation in Healthy and Amebic Gill Disease-Affected Atlantic Salmon. Microorganisms, 2020, 8, 1871.	3.6	10
30	Effect of a prophylactic treatment with chloramine-T on gill histology and microbiome of Atlantic salmon (Salmo salar) under commercial conditions. Aquaculture, 2022, 546, 737319.	3.5	10
31	Prevalence of six amoeba species colonising the gills of farmed Atlantic salmon with amoebic gill disease (AGD) using qPCR. Aquaculture Environment Interactions, 2019, 11, 405-415.	1.8	10
32	Pilchard orthomyxovirus (POMV). I. Characterisation of an emerging virus isolated from pilchards Sardinops sagax and Atlantic salmon Salmo salar. Diseases of Aquatic Organisms, 2020, 139, 35-50.	1.0	10
33	Mouse fibroblast L929 cells are less permissive to infection by Nelson Bay orthoreovirus compared to other mammalian cell lines. Journal of General Virology, 2015, 96, 1787-1794.	2.9	9
34	Comparative transcriptome analysis of pilchard orthomyxovirus (POMV) and infectious salmon anaemia virus (ISAV). Fish and Shellfish Immunology, 2020, 105, 415-426.	3.6	8
35	Dead or alive: microbial viability treatment reveals both active and inactive bacterial constituents in the fish gut microbiota. Journal of Applied Microbiology, 2021, 131, 2528-2538.	3.1	8
36	Hydrogen peroxide treatment of Atlantic salmon temporarily decreases oxygen consumption but has negligible effects on hypoxia tolerance and aerobic performance. Aquaculture, 2021, 540, 736676.	3.5	8

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#	Article	IF	CITATIONS
37	Comparison of bacterial diversity and distribution on the gills of Atlantic salmon ( <i>Salmo salar</i> ) Tj ETQq1 1 (	).784314 3.1	rgBT /Overl
38	Seawater transmission and infection dynamics of pilchard orthomyxovirus (POMV) in Atlantic salmon ( <i>Salmo salar</i> ). Journal of Fish Diseases, 2021, 44, 73-88.	1.9	7
39	Allelic and haplotypic diversity at the major histocompatibility class II within domesticated Australian Atlantic salmon (Salmo salar L.). Journal of Fish Biology, 2007, 70, 45-59.	1.6	6
40	Proteomic analysis of Pteropus alecto kidney cells in response to the viral mimic, Poly I:C. Proteome Science, 2015, 13, 25.	1.7	6
41	Proteomics informed by transcriptomics for characterising differential cellular susceptibility to Nelson Bay orthoreovirus infection. BMC Genomics, 2017, 18, 615.	2.8	6
42	A High Throughput Viability Screening Method for the Marine Ectoparasite Neoparamoeba perurans. Protist, 2020, 171, 125773.	1.5	6
43	Sensory Rewiring in an Echolocator: Genome-Wide Modification of Retinogenic and Auditory Genes in the Bat <i>Myotis davidii</i> . G3: Genes, Genomes, Genetics, 2014, 4, 1825-1835.	1.8	5
44	Immersion challenge of naÃ⁻ve Atlantic salmon with cultured <i>Nolandella</i> sp. and <i>Pseudoparamoeba</i> sp. did not increase the severity of <i>Neoparamoeba perurans</i> â€induced amoebic gill disease (AGD). Journal of Fish Diseases, 2021, 44, 149-160.	1.9	5
45	Evaluation of sodium percarbonate as a bath treatment for amoebic gill disease in Atlantic salmon. Aquaculture Research, 2021, 52, 117-129.	1.8	5
46	Low incidence of recurrent Buruli ulcers in treated Australian patients living in an endemic region. PLoS Neglected Tropical Diseases, 2018, 12, e0006724.	3.0	4
47	Novacqâ"¢ improves survival of Penaeus vannamei when challenged with pathogenic Vibrio parahaemolyticus causing acute hepatopancreatic necrosis disease. Aquaculture, 2021, 545, 737235.	3.5	4
48	Profiling Branchial Bacteria of Atlantic Salmon (Salmo salar L.) Following Exposure to Antimicrobial Agents. Frontiers in Animal Science, 2021, 2, .	1.9	4
49	Searching for the sweet spot of amoebic gill disease of farmed Atlantic salmon: the potential role of glycan-lectin interactions in the adhesion of Neoparamoeba perurans. International Journal for Parasitology, 2021, 51, 545-557.	3.1	3
50	The ability of Neoparamoeba perurans to bind to and digest nonâ€fishâ€derived mucin: Insights into the amoeba's mechanism of action to overcome gill mucus production. Journal of Fish Diseases, 2021, 44, 1355-1367.	1.9	3
51	Characterization of a major histocompatibility class II <i>A </i> gene ( <i>Clhaâ€DAA</i> ) with an embedded microsatellite marker in Atlantic herring ( <i>Clupea harengus </i> L.). Journal of Fish Biology, 2008, 73, 367-381.	1.6	2
52	SNP genotyping of animal and human derived isolates of Mycobacterium avium subsp. paratuberculosis. Veterinary Microbiology, 2014, 172, 479-485.	1.9	2
53	Evaluation of the Infectious Potential of Neoparamoeba perurans Following Freshwater Bathing Treatments. Microorganisms, 2021, 9, 967.	3.6	2
54	Will Australia's common carp (Cyprinus carpio) populations develop resistance to Cyprinid herpesvirus 3 (CyHV-3) if released as a biocontrol agent? Identification of pathways and knowledge gaps. Biological Control, 2021, 157, 104571.	3.0	2

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55	Competing endogenous RNA-networks reveal key regulatory microRNAs involved in the response of Atlantic salmon to a novel orthomyxovirus. Developmental and Comparative Immunology, 2022, 132, 104396.	2.3	2
56	First detection of a novel â€~unknown host' flavivirus in a Malaysian rodent. Access Microbiology, 2021, 3, 000223.	0.5	1
57	Development of a new real-time PCR for the detection of pilchard orthomyxovirus (POMV) in apparently healthy fish. Aquaculture, 2022, 547, 737404.	3.5	0