Jason P Mulvenna

List of Publications by Year in descending order

Source: https://exaly.com/author-pdf/8866759/publications.pdf

Version: 2024-02-01

92 papers 5,185 citations

94433 37 h-index 91884 69 g-index

97 all docs 97
docs citations

97 times ranked 5424 citing authors

#	Article	IF	Citations
1	The tumorigenic liver fluke Opisthorchis viverrini – multiple pathways to cancer. Trends in Parasitology, 2012, 28, 395-407.	3.3	376
2	Opisthorchiasis and Opisthorchis-associated cholangiocarcinoma in Thailand and Laos. Acta Tropica, 2011, 120, S158-S168.	2.0	262
3	Helminth Immunomodulation in Autoimmune Disease. Frontiers in Immunology, 2017, 8, 453.	4.8	182
4	Discovery, Structure and Biological Activities of the Cyclotides. Current Protein and Peptide Science, 2004, 5, 297-315.	1.4	167
5	Proteomics Analysis of the Excretory/Secretory Component of the Blood-feeding Stage of the Hookworm, Ancylostoma caninum. Molecular and Cellular Proteomics, 2009, 8, 109-121.	3.8	167
6	Genome of the human hookworm Necator americanus. Nature Genetics, 2014, 46, 261-269.	21,4	166
7	A Granulin-Like Growth Factor Secreted by the Carcinogenic Liver Fluke, Opisthorchis viverrini, Promotes Proliferation of Host Cells. PLoS Pathogens, 2009, 5, e1000611.	4.7	162
8	Generalized urticaria induced by the Na-ASP-2 hookworm vaccine: Implications for the development of vaccines against helminths. Journal of Allergy and Clinical Immunology, 2012, 130, 169-176.e6.	2.9	151
9	Extracellular vesicles secreted by Schistosoma mansoni contain protein vaccine candidates. International Journal for Parasitology, 2016, 46, 1-5.	3.1	147
10	Carcinogenic Liver Fluke Secretes Extracellular Vesicles That Promote Cholangiocytes to Adopt a Tumorigenic Phenotype. Journal of Infectious Diseases, 2015, 212, 1636-1645.	4.0	141
11	CyBase: a database of cyclic protein sequence and structure. Nucleic Acids Research, 2006, 34, D192-D194.	14.5	137
12	The secreted and surface proteomes of the adult stage of the carcinogenic human liver fluke <i>Opisthorchis viverrini</i> . Proteomics, 2010, 10, 1063-1078.	2.2	135
13	Exposed proteins of the Schistosoma japonicum tegument. International Journal for Parasitology, 2010, 40, 543-554.	3.1	130
14	Infection with the carcinogenic liver fluke <i>Opisthorchis viverrini</i> modifies intestinal and biliary microbiome. FASEB Journal, 2013, 27, 4572-4584.	0.5	116
15	A quantitative proteomic analysis of the tegumental proteins from Schistosoma mansoni schistosomula reveals novel potential therapeutic targets. International Journal for Parasitology, 2015, 45, 505-516.	3.1	103
16	Transcriptome and venom proteome of the box jellyfish Chironex fleckeri. BMC Genomics, 2015, 16, 407.	2.8	103
17	Elevated Plasma IL-6 Associates with Increased Risk of Advanced Fibrosis and Cholangiocarcinoma in Individuals Infected by Opisthorchis viverrini. PLoS Neglected Tropical Diseases, 2012, 6, e1654.	3.0	96
18	Gene discovery for the carcinogenic human liver fluke, Opisthorchis viverrini. BMC Genomics, 2007, 8, 189.	2.8	90

#	Article	IF	CITATIONS
19	Proteomic characterisation of Echinococcus granulosus hydatid cyst fluid from sheep, cattle and humans. Journal of Proteomics, 2011, 74, 1560-1572.	2.4	88
20	Secreted Proteomes of Different Developmental Stages of the Gastrointestinal Nematode Nippostrongylus brasiliensis. Molecular and Cellular Proteomics, 2014, 13, 2736-2751.	3.8	88
21	An enzymatically inactivated hemoglobinase from <i>Necator americanus</i> induces neutralizing antibodies against multiple hookworm species and protects dogs against heterologous hookworm infection. FASEB Journal, 2009, 23, 3007-3019.	0.5	83
22	Semienzymatic Cyclization of Disulfide-rich Peptides Using Sortase A. Journal of Biological Chemistry, 2014, 289, 6627-6638.	3.4	83
23	Processing of a 22 kDa Precursor Protein to Produce the Circular Protein Tricyclon A. Structure, 2005, 13, 691-701.	3.3	78
24	Carcinogenic Parasite Secretes Growth Factor That Accelerates Wound Healing and Potentially Promotes Neoplasia. PLoS Pathogens, 2015, 11, e1005209.	4.7	78
25	Cloning and Characterisation of Schistosoma japonicum Insulin Receptors. PLoS ONE, 2010, 5, e9868.	2.5	76
26	Chironex fleckeri (Box Jellyfish) Venom Proteins. Journal of Biological Chemistry, 2014, 289, 4798-4812.	3.4	72
27	Discovery of Cyclotide-Like Protein Sequences in Graminaceous Crop Plants: Ancestral Precursors of Circular Proteins?. Plant Cell, 2006, 18, 2134-2144.	6.6	70
28	Tissue Specific Profiling of Females of Schistosoma japonicum by Integrated Laser Microdissection Microscopy and Microarray Analysis. PLoS Neglected Tropical Diseases, 2009, 3, e469.	3.0	70
29	Transcriptional Changes in Schistosoma mansoni during Early Schistosomula Development and in the Presence of Erythrocytes. PLoS Neglected Tropical Diseases, 2010, 4, e600.	3.0	70
30	Tentacle Transcriptome and Venom Proteome of the Pacific Sea Nettle, Chrysaora fuscescens (Cnidaria: Scyphozoa). Toxins, 2016, 8, 102.	3.4	70
31	Infection with the carcinogenic human liver fluke, Opisthorchis viverrini. Molecular BioSystems, 2011, 7, 1367.	2.9	60
32	Venom Proteome of the Box Jellyfish Chironex fleckeri. PLoS ONE, 2012, 7, e47866.	2.5	57
33	Neutralizing Antibodies to the Hookworm Hemoglobinase <i>Na</i> à€APRâ€1: Implications for a Multivalent Vaccine against Hookworm Infection and Schistosomiasis. Journal of Infectious Diseases, 2010, 201, 1561-1569.	4.0	53
34	Identification of a Novel, EBV-Based Antibody Risk Stratification Signature for Early Detection of Nasopharyngeal Carcinoma in Taiwan. Clinical Cancer Research, 2018, 24, 1305-1314.	7.0	52
35	A family of cathepsin B cysteine proteases expressed in the gut of the human hookworm, Necator americanus. Molecular and Biochemical Parasitology, 2008, 160, 90-99.	1.1	50
36	Evolution of resistance to chytridiomycosis is associated with a robust early immune response. Molecular Ecology, 2018, 27, 919-934.	3.9	50

#	Article	IF	Citations
37	Vaccinomics for the Major Blood Feeding Helminths of Humans. OMICS A Journal of Integrative Biology, 2011, 15, 567-577.	2.0	48
38	A modified FASP protocol for high-throughput preparation of protein samples for mass spectrometry. PLoS ONE, 2017, 12, e0175967.	2.5	44
39	Up-regulation of annexin A2 in cholangiocarcinoma caused by Opisthorchis viverrini and its implication as a prognostic marker. International Journal for Parasitology, 2010, 40, 1203-1212.	3.1	37
40	Distinct miRNA signatures associate with subtypes of cholangiocarcinoma from infection with the tumourigenic liver fluke Opisthorchis viverrini. Journal of Hepatology, 2014, 61, 850-858.	3.7	37
41	A Deep Exploration of the Transcriptome and "Excretory/Secretory―Proteome of Adult Fascioloides magna. Molecular and Cellular Proteomics, 2012, 11, 1340-1353.	3.8	35
42	Integrated Transcriptomic-Proteomic Analysis Using a Proteogenomic Workflow Refines Rat Genome Annotation. Molecular and Cellular Proteomics, 2016, 15, 329-339.	3.8	35
43	Expression, refolding and purification of Ov-GRN-1, a granulin-like growth factor from the carcinogenic liver fluke, that causes proliferation of mammalian host cells. Protein Expression and Purification, 2011, 79, 263-270.	1.3	34
44	Solution Structure, Membrane Interactions, and Protein Binding Partners of the Tetraspanin Sm-TSP-2, a Vaccine Antigen from the Human Blood Fluke Schistosoma mansoni. Journal of Biological Chemistry, 2014, 289, 7151-7163.	3.4	33
45	Unique molecular profile of exosomes derived from primary human proximal tubular epithelial cells under diseased conditions. Journal of Extracellular Vesicles, 2017, 6, 1314073.	12.2	33
46	Discovery, Structural Determination, and Putative Processing of the Precursor Protein That Produces the Cyclic Trypsin Inhibitor Sunflower Trypsin Inhibitor 1. Journal of Biological Chemistry, 2005, 280, 32245-32253.	3.4	32
47	Methods and matrices: approaches to identifying miRNAs for Nasopharyngeal carcinoma. Journal of Translational Medicine, 2014, 12, 3.	4.4	32
48	A microRNA profile associated with Opisthorchis viverrini-induced cholangiocarcinoma in tissue and plasma. BMC Cancer, 2015, 15, 309.	2.6	32
49	Ov-APR-1, an aspartic protease from the carcinogenic liver fluke, Opisthorchis viverrini: Functional expression, immunolocalization and subsite specificity. International Journal of Biochemistry and Cell Biology, 2009, 41, 1148-1156.	2.8	30
50	Qualitative and quantitative proteomic analyses of Schistosoma japonicum eggs and egg-derived secretory-excretory proteins. Parasites and Vectors, 2019, 12, 173.	2,5	29
51	Genomic-Bioinformatic Analysis of Transcripts Enriched in the Third-Stage Larva of the Parasitic Nematode Ascaris suum. PLoS Neglected Tropical Diseases, 2008, 2, e246.	3.0	27
52	Prognostic significance of peroxiredoxin 1 and ezrin-radixin-moesin–binding phosphoprotein 50 in cholangiocarcinoma. Human Pathology, 2012, 43, 1719-1730.	2.0	27
53	Exploiting Helminth–Host Interactomes through Big Data. Trends in Parasitology, 2017, 33, 875-888.	3.3	27
54	Anthelminthic activity of the cyclotides (kalata B1 and B2) against schistosome parasites. Biopolymers, 2013, 100, 461-470.	2.4	26

#	Article	IF	CITATIONS
55	Microproteinuria during Opisthorchis viverrini Infection: A Biomarker for Advanced Renal and Hepatobiliary Pathologies from Chronic Opisthorchiasis. PLoS Neglected Tropical Diseases, 2013, 7, e2228.	3.0	25
56	Circumventing qPCR inhibition to amplify miRNAs in plasma. Biomarker Research, 2014, 2, 13.	6.8	25
57	Comprehensive analysis of the secreted proteome of adult Necator americanusÂhookworms. PLoS Neglected Tropical Diseases, 2020, 14, e0008237.	3.0	25
58	Rapid short term and gradual permanent cardiotoxic effects of vertebrate toxins from Chironex fleckeri (Australian box jellyfish) venom. Toxicon, 2014, 80, 17-26.	1.6	24
59	Urine proteomics study reveals potential biomarkers for the differential diagnosis of cholangiocarcinoma and periductal fibrosis. PLoS ONE, 2019, 14, e0221024.	2.5	21
60	Tetraspanin-2 localisation in high pressure frozen and freeze-substituted Schistosoma mansoni adult males reveals its distribution in membranes of tegumentary vesicles. International Journal for Parasitology, 2013, 43, 785-793.	3.1	18
61	Discovery and Qualification of Serum Protein Biomarker Candidates for Cholangiocarcinoma Diagnosis. Journal of Proteome Research, 2019, 18, 3305-3316.	3.7	18
62	Proteomic profile of Bithynia siamensis goniomphalos snails upon infection with the carcinogenic liver fluke Opisthorchis viverrini. Journal of Proteomics, 2015, 113, 281-291.	2.4	17
63	Kunitz type protease inhibitor EgKI-1 from the canine tapeworm Echinococcus granulosus as a promising therapeutic against breast cancer. PLoS ONE, 2018, 13, e0200433.	2.5	17
64	Proteomic and genomic analyses suggest the association of apolipoprotein C1 with abdominal aortic aneurysm. Proteomics - Clinical Applications, 2014, 8, 762-772.	1.6	16
65	A comparative proteomic analysis of bile for biomarkers of cholangiocarcinoma. Tumor Biology, 2017, 39, 101042831770576.	1.8	16
66	Coming out of the Shell: Building the Molecular Infrastructure for Research on Parasite-Harbouring Snails. PLoS Neglected Tropical Diseases, 2013, 7, e2284.	3.0	15
67	Proteomic Identification of Plasma Protein Tyrosine Phosphatase Alpha and Fibronectin Associated with Liver Fluke, Opisthorchis viverrini, Infection. PLoS ONE, 2012, 7, e45460.	2.5	15
68	Discovery and structures of the cyclotides: novel macrocyclic peptides from plants. International Journal of Peptide Research and Therapeutics, 2001, 8, 119-128.	0.1	14
69	Insights into the Membrane Interactions of the Saposin-Like Proteins Na-SLP-1 and Ac-SLP-1 from Human and Dog Hookworm. PLoS ONE, 2011, 6, e25369.	2.5	14
70	Profiling miRNAs in nasopharyngeal carcinoma FFPE tissue by microarray and Next Generation Sequencing. Genomics Data, 2014, 2, 285-289.	1.3	13
71	Patterns of Interindividual Variability in the Antibody Repertoire Targeting Proteins Across the Epstein-Barr Virus Proteome. Journal of Infectious Diseases, 2018, 217, 1923-1931.	4.0	13
72	Mollusk microbiota shift during Angiostrongylus cantonensis infection in the freshwater snail Biomphalaria glabrata and the terrestrial slug Phillocaulis soleiformis. Parasitology Research, 2020, 119, 2495-2503.	1.6	12

#	Article	IF	Citations
73	Levels of 8-OxodG Predict Hepatobiliary Pathology in Opisthorchis viverrini Endemic Settings in Thailand. PLoS Neglected Tropical Diseases, 2015, 9, e0003949.	3.0	12
74	Differential Protein Expression in the Hemolymph of Bithynia siamensis goniomphalos Infected with Opisthorchis viverrini. PLoS Neglected Tropical Diseases, 2016, 10, e0005104.	3.0	12
75	Exploring transcriptional conservation between Ancylostoma caninum and Haemonchus contortus by oligonucleotide microarray and bioinformatic analyses. Molecular and Cellular Probes, 2009, 23, 1-9.	2.1	11
76	A primary human T-cell spectral library to facilitate large scale quantitative T-cell proteomics. Scientific Data, 2020, 7, 412.	5.3	11
77	Clinicopathological Significance of Osteopontin in Cholangiocarcinoma Cases. Asian Pacific Journal of Cancer Prevention, 2016, 17, 201-205.	1.2	11
78	Molecular and phylogenetic characterization of cytochromes c from Haemonchus contortus and Trichostrongylus vitrinus (Nematoda: Trichostrongylida). Gene, 2008, 424, 121-129.	2.2	10
79	Lysosome-associated membrane glycoprotein (LAMP) $\hat{a}\in$ preliminary study on a hidden antigen target for vaccination against schistosomiasis. Scientific Reports, 2015, 5, 15069.	3.3	10
80	Discovery and structures of the cyclotides: novel macrocyclic peptides from plants. International Journal of Peptide Research and Therapeutics, 2001, 8, 119-128.	0.1	9
81	Differential Protein Expression Marks the Transition From Infection With Opisthorchis viverrini to Cholangiocarcinoma. Molecular and Cellular Proteomics, 2017, 16, 911-923.	3.8	9
82	Survival, gene and metabolite responses of Litoria verreauxii alpina frogs to fungal disease chytridiomycosis. Scientific Data, 2018, 5, 180033.	5.3	9
83	Discovering proteins for chemoprevention and chemotherapy by curcumin in liver fluke infection-induced bile duct cancer. PLoS ONE, 2018, 13, e0207405.	2.5	9
84	Changes in protein expression after treatment with Ancylostoma caninum excretory/secretory products in a mouse model of colitis. Scientific Reports, 2017, 7, 41883.	3.3	8
85	Banking on the future: Biobanking for "omics―approaches to biomarker discovery for Opisthorchis-induced cholangiocarcinoma in Thailand. Parasitology International, 2012, 61, 173-177.	1.3	7
86	ERK and mTORC1 Inhibitors Enhance the Anti-Cancer Capacity of the Octpep-1 Venom-Derived Peptide in Melanoma BRAF(V600E) Mutations. Toxins, 2021, 13, 146.	3.4	7
87	Bioclojure: a functional library for the manipulation of biological sequences. Bioinformatics, 2014, 30, 2537-2539.	4.1	6
88	Data set from the proteomic analysis of Bithynia siamensis goniomphalos snails upon infection with the carcinogenic liver fluke Opisthorchis viverrini. Data in Brief, 2015, 2, 16-20.	1.0	6
89	Proteomic identification of the contents of small extracellular vesicles from in vivo Plasmodium yoelii infection. International Journal for Parasitology, 2022, 52, 35-45.	3.1	6
90	The miRNAome of Opisthorchis viverrini induced intrahepatic cholangiocarcinoma. Genomics Data, 2014, 2, 274-279.	1.3	5

#	Article	IF	CITATIONS
91	Synthetic hookworm-derived peptides are potent modulators of primary human immune cell function that protect against experimental colitis inÂvivo. Journal of Biological Chemistry, 2021, 297, 100834.	3.4	5
92	Characterization and binding affinities of SmLANP: A new Schistosoma mansoni member of the ANP32 family of regulatory proteins. Molecular and Biochemical Parasitology, 2009, 165, 95-102.	1.1	1