Hurng-Yi Wang

List of Publications by Year in descending order

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		236925	189892
52	2,675	25	50
papers	2,675 citations	h-index	g-index
55	55	55	4638
all docs	docs citations	times ranked	citing authors

#	Article	IF	CITATIONS
1	Evolution under Spatially Heterogeneous Selection in Solid Tumors. Molecular Biology and Evolution, 2022, 39, .	8.9	5
2	The origin and past demography of murine astrovirus 1 in laboratory mice. Journal of General Virology, $2021,102,.$	2.9	1
3	Whole genome deep sequencing analysis of viral quasispecies diversity and evolution in HBeAg seroconverters. JHEP Reports, 2021, 3, 100254.	4.9	9
4	Gut microbiota in the innate immunity against hepatitis B virus $\hat{a} \in \text{``implication in age-dependent HBV clearance. Current Opinion in Virology, 2021, 49, 194-202.}$	5.4	7
5	A rapid insect species identification system using miniâ€barcode pyrosequencing. Pest Management Science, 2020, 76, 1222-1227.	3.4	4
6	Cover Image, Volume 76, Issue 4. Pest Management Science, 2020, 76, i.	3.4	0
7	Implication of single year seasonal sampling to genetic diversity fluctuation that coordinates with oceanographic dynamics in torpedo scads near Taiwan. Scientific Reports, 2020, 10, 16829.	3.3	1
8	The origin and underlying driving forces of the SARS-CoV-2 outbreak. Journal of Biomedical Science, 2020, 27, 73.	7.0	82
9	Clonality of Fluconazole-Nonsusceptible <i>Candida tropicalis</i> in Bloodstream Infections, Taiwan, 2011–2017. Emerging Infectious Diseases, 2019, 25, 1660-1667.	4.3	40
10	Lymphocyte Antigen 6 Complex, Locus C+ Monocytes and Kupffer Cells Orchestrate Liver Immune Responses Against Hepatitis B Virus in Mice. Hepatology, 2019, 69, 2364-2380.	7.3	36
11	Effects of indoor residual spraying and outdoor larval control on Anopheles coluzzii from São Tomé and PrÃncipe, two islands with pre-eliminated malaria. Malaria Journal, 2019, 18, 405.	2.3	8
12	Genetic Load and Potential Mutational Meltdown in Cancer Cell Populations. Molecular Biology and Evolution, 2019, 36, 541-552.	8.9	14
13	Is the evolution in tumors Darwinian or non-Darwinian?. National Science Review, 2018, 5, 15-17.	9.5	24
14	On the low reproducibility of cancer studies. National Science Review, 2018, 5, 619-624.	9.5	38
15	De novo assembly of highly polymorphic metagenomic data using in situ generated reference sequences and a novel BLAST-based assembly pipeline. BMC Bioinformatics, 2017, 18, 223.	2.6	14
16	Hydrodynamic HBV Transfection Mouse Model. Methods in Molecular Biology, 2017, 1540, 227-235.	0.9	7
17	The Ecology and Evolution of Cancer: The Ultra-Microevolutionary Process. Annual Review of Genetics, 2016, 50, 347-369.	7.6	86
18	PATHOLOGY AND MOLECULAR DETECTION OF RABIES VIRUS IN FERRET BADGERS ASSOCIATED WITH A RABIES OUTBREAK IN TAIWAN. Journal of Wildlife Diseases, 2016, 52, 57-69.	0.8	9

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19	Microâ€evolution of the hepatitis <scp>B</scp> virus genome in hepatitis <scp>B</scp> eâ€antigenâ€positive carriers: Comparison of genotypes <scp>B</scp> and <scp>C</scp> at various immune stages. Journal of Gastroenterology and Hepatology (Australia), 2015, 30, 172-177.		2
20	New Insights into the Evolutionary Rate of Hepatitis B Virus at Different Biological Scales. Journal of Virology, 2015, 89, 3512-3522.	3.4	33
21	Age-related immune clearance of hepatitis B virus infection requires the establishment of gut microbiota. Proceedings of the National Academy of Sciences of the United States of America, 2015, 112, 2175-2180.	7.1	175
22	Instability of Succinate Dehydrogenase in SDHD Polymorphism Connects Reactive Oxygen Species Production to Nuclear and Mitochondrial Genomic Mutations in Yeast. Antioxidants and Redox Signaling, 2015, 22, 587-602.	5 . 4	9
23	The CRISPR/Cas9 System Facilitates Clearance of the Intrahepatic HBV Templates In Vivo. Molecular Therapy - Nucleic Acids, 2014, 3, e186.	5.1	319
24	Molecular Characterization of Cryptically Circulating Rabies Virus from Ferret Badgers, Taiwan. Emerging Infectious Diseases, 2014, 20, 790-798.	4.3	49
25	Influence of gene flow on divergence dating – implications for the speciation history of <i>Takydromus</i> grass lizards. Molecular Ecology, 2014, 23, 4770-4784.	3.9	17
26	Multilocus approach to clarify species status and the divergence history of the Bemisia tabaci (Hemiptera: Aleyrodidae) species complex. Molecular Phylogenetics and Evolution, 2014, 76, 172-180.	2.7	23
27	Clinical features and phylogenetic analysis of Coxsackievirus A9 in Northern Taiwan in 2011. BMC Infectious Diseases, 2013, 13, 33.	2.9	19
28	Genetic and physiological data suggest demographic and adaptive responses in complex interactions between populations of figs (<i><scp>F</scp>icus pumila</i>) and their pollinating wasps (<i><scp>W</scp>iebesia pumilae</i>). Molecular Ecology, 2013, 22, 3814-3832.	3.9	7
29	Ribavirin Enhances the Action of Interferon- \hat{l} ± against Hepatitis C Virus by Promoting the p53 Activity through the ERK1/2 Pathway. PLoS ONE, 2012, 7, e43824.	2.5	13
30	Targeted Disruption in Mice of a Neural Stem Cell-Maintaining, KRAB-Zn Finger-Encoding Gene That Has Rapidly Evolved in the Human Lineage. PLoS ONE, 2012, 7, e47481.	2.5	11
31	Loopâ€mediated isothermal amplification for rapid identification of biotypes B and Q of the globally invasive pest <i>Bemisia tabaci⟨ i⟩, and studying population dynamics. Pest Management Science, 2012, 68, 1206-1213.</i>	3.4	26
32	Permanent Genetic Resources added to Molecular Ecology Resources Database 1 April 2012 – 31 May 2012. Molecular Ecology Resources, 2012, 12, 972-974.	4.8	18
33	Systematics and phylogeography of the Taiwanese endemic minnow Candidia barbatus (Pisces:) Tj ETQq1 1 0.78 Linnean Society, 2011, 161, 613-632.	4314 rgB1 2.3	「Overlock 1 19
34	Rapid growth of a hepatocellular carcinoma and the driving mutations revealed by cell-population genetic analysis of whole-genome data. Proceedings of the National Academy of Sciences of the United States of America, 2011, 108, 12042-12047.	7.1	117
35	Inferring nonneutral evolution from contrasting patterns of polymorphisms and divergences in different protein coding regions of enterovirus 71 circulating in Taiwan during 1998-2003. BMC Evolutionary Biology, 2010, 10, 294.	3.2	11
36	Molecular and Phylogenetic Analyses Suggest an Additional Hepatitis B Virus Genotype "l― PLoS ONE, 2010, 5, e9297.	2.5	123

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37	Distinct Hepatitis B Virus Dynamics in the Immunotolerant and Early Immunoclearance Phases. Journal of Virology, 2010, 84, 3454-3463.	3.4	81
38	Pre-S2 Deletions of Hepatitis B Virus and Hepatocellular Carcinoma in Children. Pediatric Research, 2010, 67, 90-94.	2.3	36
39	Identification of androgen response elements in the enhancer I of hepatitis B virus: A mechanism for sex disparity in chronic hepatitis B. Hepatology, 2009, 50, 1392-1402.	7.3	151
40	Predation Risks and Antiâ€Predation Parental Care Behavior: An Experimental Study in a Tropical Skink. Ethology, 2009, 115, 273-279.	1.1	23
41	Complex genetic interactions underlying expression differences between <i>Drosophila</i> races: Analysis of chromosome substitutions. Proceedings of the National Academy of Sciences of the United States of America, 2008, 105, 6362-6367.	7.1	29
42	Adaptive genic evolution in the Drosophila genomes. Proceedings of the National Academy of Sciences of the United States of America, 2007, 104, 2271-2276.	7.1	210
43	Rate of Evolution in Brain-Expressed Genes in Humans and Other Primates. PLoS Biology, 2006, 5, e13.	5.6	93
44	Characterization of severe acute respiratory syndrome coronavirus genomes in Taiwan: Molecular epidemiology and genome evolution. Proceedings of the National Academy of Sciences of the United States of America, 2004, 101, 2542-2547.	7.1	121
45	Origin of serum hepatitis B virus in acute exacerbation: Comparison with HBV in the liver and from other exacerbation. Hepatology, 2004, 40, 310-317.	7.3	22
46	Structural diversity and functional implications of the eukaryotic TDP gene family. Genomics, 2004, 83, 130-139.	2.9	261
47	Lack of Interferon Sensitivity-Determining Region in the Genome of Hepatitis B Virus Genotype Ba. Antiviral Therapy, 2004, 9, 895-903.	1.0	10
48	Rapidly Evolving Genes in Human. I. The Glycophorins and Their Possible Role in Evading Malaria Parasites. Molecular Biology and Evolution, 2003, 20, 1795-1804.	8.9	85
49	Secondary Structure of Mitochondrial 12S rRNA Among Fish and Its Phylogenetic Applications. Molecular Biology and Evolution, 2002, 19, 138-148.	8.9	42
50	Molecular Phylogeny of Gobioid Fishes (Perciformes: Gobioidei) Based on Mitochondrial 12S rRNA Sequences. Molecular Phylogenetics and Evolution, 2001, 20, 390-408.	2.7	50
51	Phylogeny, Taxonomy, and Biogeography of the Oriental Pitvipers of the Genus Trimeresurus (Reptilia:) Tj ETQq1	1 8:78431	14ggBT/Ove
52	Influence of glaciation on divergence patterns of the endemic minnow, Zacco pachycephalus, in Taiwan. Molecular Ecology, 1999, 8, 1879-1888.	3.9	46