

Michael A Purdy

List of Publications by Citations

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The third column is the impact factor (IF) of the journal, and the fourth column is the number of citations of the article.

28

papers

1,702

citations

14

h-index

29

g-index

29

ext. papers

1,891

ext. citations

6.1

avg, IF

4.49

L-index

#	Paper	IF	Citations
28	Consensus proposals for classification of the family Hepeviridae. <i>Journal of General Virology</i> , 2014 , 95, 2223-2232	4.9	468
27	Chronic Infection With Camelid Hepatitis E Virus in a Liver Transplant Recipient Who Regularly Consumes Camel Meat and Milk. <i>Gastroenterology</i> , 2016 , 150, 355-7.e3	13.3	306
26	Proposed reference sequences for hepatitis E virus subtypes. <i>Journal of General Virology</i> , 2016 , 97, 537-542	4.7	284
25	Genetic variability and the classification of hepatitis E virus. <i>Journal of Virology</i> , 2013 , 87, 4161-9	6.6	109
24	Evolutionary history and population dynamics of hepatitis E virus. <i>PLoS ONE</i> , 2010 , 5, e14376	3.7	86
23	The molecular epidemiology of hepatitis E virus infection. <i>Virus Research</i> , 2011 , 161, 31-9	6.4	73
22	Epidemic history and evolutionary dynamics of hepatitis B virus infection in two remote communities in rural Nigeria. <i>PLoS ONE</i> , 2010 , 5, e11615	3.7	71
21	Restricted enzooticity of hepatitis E virus genotypes 1 to 4 in the United States. <i>Journal of Clinical Microbiology</i> , 2011 , 49, 4164-72	9.7	63
20	The hepatitis E virus polyproline region is involved in viral adaptation. <i>PLoS ONE</i> , 2012 , 7, e35974	3.7	59
19	Hepatitis E virus genotype 4, Nanjing, China, 2001-2011. <i>Emerging Infectious Diseases</i> , 2013 , 19, 1528-30	10.2	40
18	Evolution of the hepatitis E virus polyproline region: order from disorder. <i>Journal of Virology</i> , 2012 , 86, 10186-93	6.6	22
17	Rabbit and human hepatitis E virus strains belong to a single serotype. <i>Virus Research</i> , 2013 , 176, 101-6	6.4	20
16	Recent population expansions of hepatitis B virus in the United States. <i>Journal of Virology</i> , 2014 , 88, 13971-80	6.6	18
15	Human and porcine hepatitis E viruses, southeastern Bolivia. <i>Emerging Infectious Diseases</i> , 2012 , 18, 339-40	10.2	17
14	Disparities in detection of antibodies against hepatitis E virus in US blood donor samples using commercial assays. <i>Transfusion</i> , 2018 , 58, 1254-1263	2.9	14
13	Hepatitis E virus isolated from rabbits is genetically heterogeneous but with very similar antigenicity to human HEV. <i>Journal of Medical Virology</i> , 2013 , 85, 627-35	19.7	14
12	Genetic host specificity of hepatitis E virus. <i>Infection, Genetics and Evolution</i> , 2014 , 24, 127-39	4.5	13

11	Supragenotypic groups of the hepatitis B virus genome. <i>Journal of General Virology</i> , 2008 , 89, 1179-1183	4.9	10
10	The effect of phylogenetic signal reduction on genotyping of hepatitis E viruses of the species <i>Orthohepevirus A</i> . <i>Archives of Virology</i> , 2017 , 162, 645-656	2.6	9
9	Distribution of hepatitis A antibodies in US blood donors. <i>Transfusion</i> , 2018 , 58, 2761-2765	2.9	4
8	Hepatitis E in Singapore: A Case-Series and Viral Phylodynamics Study. <i>American Journal of Tropical Medicine and Hygiene</i> , 2017 , 96, 922-928	3.2	2
7	Transcriptome analysis in rhesus macaques infected with hepatitis E virus genotype 1/3 infections and genotype 1 re-infection. <i>PLoS ONE</i> , 2020 , 15, e0237618	3.7	0
6	Evaluation of performance characteristics of hepatitis B e antigen serologic assays. <i>Journal of Clinical Virology</i> , 2018 , 109, 22-28	14.5	0
5	Permissive, in vitro replication of hepatitis B virus genotype E. <i>Journal of Virological Methods</i> , 2017 , 243, 20-24	2.6	
4	Transcriptome analysis in rhesus macaques infected with hepatitis E virus genotype 1/3 infections and genotype 1 re-infection 2020 , 15, e0237618		
3	Transcriptome analysis in rhesus macaques infected with hepatitis E virus genotype 1/3 infections and genotype 1 re-infection 2020 , 15, e0237618		
2	Transcriptome analysis in rhesus macaques infected with hepatitis E virus genotype 1/3 infections and genotype 1 re-infection 2020 , 15, e0237618		
1	Transcriptome analysis in rhesus macaques infected with hepatitis E virus genotype 1/3 infections and genotype 1 re-infection 2020 , 15, e0237618		