

Samad Amini-Bavil-Olyaei

List of Publications by Year in descending order

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39
papers

1,443
citations

304743

22
h-index

315739

38
g-index

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all docs

39
docs citations

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times ranked

2059
citing authors

#	ARTICLE	IF	CITATIONS
1	The Antiviral Effector IFITM3 Disrupts Intracellular Cholesterol Homeostasis to Block Viral Entry. <i>Cell Host and Microbe</i> , 2013, 13, 452-464.	11.0	290
2	Molecular identification of hepatitis B virus genotypes/subgenotypes: Revised classification hurdles and updated resolutions. <i>World Journal of Gastroenterology</i> , 2014, 20, 7152.	3.3	154
3	The rtA194T polymerase mutation impacts viral replication and susceptibility to tenofovir in hepatitis B e antigen-positive and hepatitis B e antigen-negative hepatitis B virus strains. <i>Hepatology</i> , 2009, 49, 1158-1165.	7.3	118
4	Negative regulation of NF- κ B activity by brain-specific TRIPartite Motif protein 9. <i>Nature Communications</i> , 2014, 5, 4820.	12.8	62
5	Are hepatitis B virus "subgenotypes" defined accurately?. <i>Journal of Clinical Virology</i> , 2010, 47, 356-360.	3.1	57
6	Genotype characterization and phylogenetic analysis of hepatitis B virus isolates from Iranian patients. <i>Journal of Medical Virology</i> , 2005, 75, 227-234.	5.0	55
7	Acute hepatitis B virus infection by genotype F despite successful vaccination in an immune-competent German patient. <i>Journal of Clinical Virology</i> , 2007, 38, 353-357.	3.1	55
8	Complete genomic sequence and phylogenetic relatedness of hepatitis B virus isolates from Iran. <i>Journal of Medical Virology</i> , 2005, 76, 318-326.	5.0	47
9	Novel hepatitis B virus subgenotype A6 in African-Belgian patients. <i>Journal of Clinical Virology</i> , 2010, 47, 93-96.	3.1	46
10	Development of a multiplex nested consensus PCR for detection and identification of major human herpesviruses in CNS infections. <i>Journal of Clinical Virology</i> , 2005, 32, 318-324.	3.1	44
11	Differential Impact of Immune Escape Mutations G145R and P120T on the Replication of Lamivudine-Resistant Hepatitis B Virus e Antigen-Positive and -Negative Strains. <i>Journal of Virology</i> , 2010, 84, 1026-1033.	3.4	40
12	Molecular evolutionary analysis and mutational pattern of full-length genomes of hepatitis B virus isolated from Belgian patients with different clinical manifestations. <i>Journal of Medical Virology</i> , 2010, 82, 379-389.	5.0	35
13	Phylogenetic analysis of hepatitis B virus full-length genomes reveals evidence for a large nosocomial outbreak in Belgium. <i>Journal of Clinical Virology</i> , 2009, 46, 61-68.	3.1	33
14	Clinical, virologic and phylogenetic features of hepatitis B infection in Iranian patients. <i>World Journal of Gastroenterology</i> , 2008, 14, 5448.	3.3	32
15	Hepatitis B virus genotypes in southwest Iran: Molecular, serological and clinical outcomes. <i>World Journal of Gastroenterology</i> , 2008, 14, 1510.	3.3	31
16	Hepatitis B virus genotyping, core promoter, and precore/core mutations among Afghan patients infected with hepatitis B: A preliminary report. <i>Journal of Medical Virology</i> , 2006, 78, 358-364.	5.0	29
17	Symptomatic Hepatitis B Virus (HBV) Reactivation despite Reduced Viral Fitness Is Associated with HBV Test and Immune Escape Mutations in an HIV-Coinfected Patient. <i>Journal of Infectious Diseases</i> , 2008, 198, 1620-1624.	4.0	28
18	Evolutionary analysis of HBV "S" antigen genetic diversity in Iranian blood donors: A nationwide study. <i>Journal of Medical Virology</i> , 2014, 86, 144-155.	5.0	28

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19	Mass migration to Europe: an opportunity for elimination of hepatitis B virus?. <i>The Lancet Gastroenterology and Hepatology</i> , 2019, 4, 315-323.	8.1	25
20	Hepatitis B virus (HBV) genotype and YMDD motif mutation profile among patients infected with HBV and untreated with lamivudine. <i>International Journal of Infectious Diseases</i> , 2008, 12, 83-87.	3.3	23
21	Prevalence, viral replication efficiency and antiviral drug susceptibility of rtQ215 polymerase mutations within the hepatitis B virus genome. <i>Journal of Hepatology</i> , 2009, 51, 647-654.	3.7	23
22	Molecular characterization of hepatitis B virus strains circulating in Belgian patients co-infected with HIV and HBV: Overt and occult infection. <i>Journal of Medical Virology</i> , 2011, 83, 1876-1884.	5.0	23
23	Molecular analysis of an HBsAg-negative hepatitis B virus mutant selected in a tenofovir-treated HIV-hepatitis B virus co-infected patient. <i>Aids</i> , 2009, 23, 268-272.	2.2	22
24	Characterization of hepatitis B virus genome variability in Iranian patients with chronic infection, a nationwide study. <i>Journal of Medical Virology</i> , 2012, 84, 414-423.	5.0	21
25	An aberrant high prevalence of hepatitis B infection among Afghans residing in one of the Bushehr refugee camps (Dalaki camp) in the southwest of Iran. <i>International Journal of Infectious Diseases</i> , 2008, 12, 101-102.	3.3	20
26	Molecular epidemiology of hepatitis delta virus (HDV) in Iran: A preliminary report. <i>Journal of Medical Virology</i> , 2008, 80, 2092-2099.	5.0	17
27	Molecular characterization of hepatitis B virus (HBV) strains circulating in the northern coast of the Persian Gulf and its comparison with worldwide distribution of HBV subgenotype D1. <i>Journal of Medical Virology</i> , 2014, 86, 745-757.	5.0	15
28	Prevalence of viral hepatitis and molecular analysis of HBV among voluntary blood donors in west Iran. <i>New Microbiologica</i> , 2009, 32, 193-8.	0.1	15
29	A novel accurate ACRS-PCR method with a digestion internal control for identification of wild type and YMDD mutants of hepatitis B virus strains. <i>Journal of Virological Methods</i> , 2006, 137, 298-303.	2.1	9
30	A novel accurate amplification created restriction site method for determination of the wild type and the precore mutant hepatitis B virus variants. <i>Journal of Virological Methods</i> , 2005, 127, 19-23.	2.1	8
31	15 year fulminant hepatitis B follow-up in Belgium: Viral evolution and signature of demographic change. <i>Infection, Genetics and Evolution</i> , 2017, 49, 221-225.	2.3	8
32	Epidemiological history and genomic characterization of non-D1 HBV strains identified in Iran. <i>Journal of Clinical Virology</i> , 2015, 63, 38-41.	3.1	7
33	HBV Subgenotypes D1, D2, D-del! Are "Old" Genotyping Methods Interpreted Correctly?. <i>Hepatitis Monthly</i> , 2013, 13, e13048.	0.2	5
34	A rare case of HBV genotype fluctuation (shifting and reversion) after liver transplantation. <i>Journal of Clinical Virology</i> , 2015, 71, 93-97.	3.1	5
35	Hepatitis E vaccine: current status and future prospects. <i>Future Virology</i> , 2009, 4, 143-154.	1.8	4
36	Why comprehensive datasets matter when inferring epidemic links or subgenotyping. <i>Infection, Genetics and Evolution</i> , 2018, 65, 350-351.	2.3	4

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37	Reconstruction of the origin and dispersal of the worldwide dominant Hepatitis B Virus subgenotype D1. <i>Virus Evolution</i> , 2022, 8, .	4.9	4
38	Hepatitis E Virus Infection in Macaca Mulatta. <i>Hepatitis Monthly</i> , 2011, 11, 852-853.	0.2	1
39	LETTER TO THE EDITOR: Associations between Hepatitis B Virus Core Gene Mutations, Immune Escape and Clinical Presentation. <i>Clinical Laboratory</i> , 2013, 59, 1097-9; author reply 1100-1.	0.5	0