Sayeh Ezzikouri

List of Publications by Year
in descending order

85	1,339	331670 21	454955 30
papers	citations	h-index	g-index

86	86	86	2044
all docs	docs citations	times ranked	citing authors

#	Article	IF	Citations
1	Prevalence and risk factors of hepatitis B and C virus infections among the general population and blood donors in Morocco. BMC Public Health, 2013, 13, 50.	2.9	78
2	Global, regional, and national sex-specific burden and control of the HIV epidemic, 1990â€“2019, for 204 countries and territories: the Global Burden of Diseases Study 2019. Lancet HIV,the, 2021, 8, e633-e651.	4.7	56
3	How artificial intelligence may help the Covidâ€19 pandemic: Pitfalls and lessons for the future. Reviews in Medical Virology, 2021, 31, 1-11.	8.3	53
4	Saliva specimens for detection of severe acute respiratory syndrome coronavirus 2 in Kuwait: A cross-sectional study. Journal of Clinical Virology, 2020, 132, 104652.	3.1	49
5	The Pro variant of the p53 codon 72 polymorphism is associated with hepatocellular carcinoma in Moroccan population. Hepatology Research, 2007, 37, 748-754.	3.4	46
6	Polymorphisms in antioxidant defence genes and susceptibility to hepatocellular carcinoma in a Moroccan population. Free Radical Research, 2010, 44, 208-216.	3.3	40
7	Hepatitis B genotypes/subgenotypes and MHR variants among Moroccan chronic carriers. Journal of Infection, 2011, 63, 66-75.	3.3	40
8	First multicenter study for risk factors for hepatocellular carcinoma development in North Africa. World Journal of Hepatology, 2011, 3, 24.	2.0	35
9	Variability in the Precore and Core Promoter Regions of HBV Strains in Morocco: Characterization and Impact on Liver Disease Progression. PLoS ONE, 2012, 7, e42891.	2.5	33
10	Genetic Variation in the Interleukin-28B Gene Is Associated with Spontaneous Clearance and Progression of Hepatitis C Virus in Moroccan Patients. PLoS ONE, 2013, 8, e54793.	2.5	33
11	Genetic variations in tollâ€like receptors 7 and 8 modulate natural hepatitis C outcomes and liver disease progression. Liver International, 2018, 38, 432-442.	3.9	33
12	MDM2 SNP309T>G polymorphism and risk of hepatocellular carcinoma: A caseâ€“control analysis in a Moroccan population. Cancer Detection and Prevention, 2009, 32, 380-385.	2.1	32
13	Subnational mapping of HIV incidence and mortality among individuals aged 15â€“49 years in sub-Saharan Africa, 2000â€“18: a modelling study. Lancet HIV,the, 2021, 8, e363-e375.	4.7	32
14	Genetic polymorphism in the manganese superoxide dismutase gene is associated with an increased risk for hepatocellular carcinoma in HCV-infected Moroccan patients. Mutation Research - Genetic Toxicology and Environmental Mutagenesis, 2008, 649, 1-6.	1.7	30
15	Genotype determination in Moroccan hepatitis B chronic carriers. Infection, Genetics and Evolution, 2008, 8, 306-312.	2.3	29
16	Morocco underwent a drift of circulating hepatitis C virus subtypes in recent decades. Archives of Virology, 2012, 157, 515-520.	2.1	29
17	Hepatitis B virus in the Maghreb Region: from epidemiology to prospective research. Liver International, 2013, 33, 811-819.	3.9	29
18	Single nucleotide polymorphism in DNMT3B promoter and its association with hepatocellular carcinoma in a Moroccan population. Infection, Genetics and Evolution, 2009, 9, 877-881.	2.3	28

#	Article	IF	Citations
19	HCV genotypes in Morocco. , 1997, 52, 396-398.		23
20	Polymorphic <scp>APOBEC</scp>3 modulates chronic hepatitis <scp>B</scp> in <scp>M</scp>oroccan population. Journal of Viral Hepatitis, 2013, 20, 678-686.	2.0	23
21	The prevalence of resistance-associated mutations to protease and reverse transcriptase inhibitors in treatment-naÃ¯ve (HIV1)-infected individuals in Casablanca, Morocco. Journal of Infection in Developing Countries, 2009, 3, 380-91.	1.2	23
22	Hepatitis C virus infection in the Maghreb region. Journal of Medical Virology, 2013, 85, 1542-1549.	5.0	22
23	Recent insights into hepatitis B virusâ€“host interactions. Journal of Medical Virology, 2014, 86, 925-932.	5.0	22
24	Occult <scp>HBV</scp> infection in Morocco: from chronic hepatitis to hepatocellular carcinoma. Liver International, 2014, 34, e144-50.	3.9	21
25	Property of hepatitis B virus replication in Tupaia belangeri hepatocytes. Biochemical and Biophysical Research Communications, 2016, 469, 229-235.	2.1	21
26	The adiponutrin I148M variant is a risk factor for HCV-associated liver cancer in North-African patients. Infection, Genetics and Evolution, 2014, 21, 179-183.	2.3	20
27	Prevalence of Common HFE and SERPINA1 Mutations in Patients with Hepatocellular Carcinoma in a Moroccan Population. Archives of Medical Research, 2008, 39, 236-241.	3.3	19
28	Inhibitory effects of PycnogenolÂ® on hepatitis C virus replication. Antiviral Research, 2015, 113, 93-102.	4.1	19
29	Oxidative Stress and Immune Responses During Hepatitis C Virus Infection in Tupaia belangeri. Scientific Reports, 2017, 7, 9848.	3.3	18
30	Lack of Ser267Phe variant of sodium taurocholate cotransporting polypeptide among Moroccans regardless of hepatitis B virus infection status. BMC Infectious Diseases, 2017, 17, 99.	2.9	17
31	Interferon-Î² response is impaired by hepatitis B virus infection in Tupaia belangeri. Virus Research, 2017, 237, 47-57.	2.2	17
32	MicroRNAs as Important Players in Host-hepatitis B Virus Interactions. Journal of Clinical and Translational Hepatology, 2015, 3, 149-61.	1.4	17
33	Coronavirus disease 2019â€”Historical context, virology, pathogenesis, immunotherapy, and vaccine development. Human Vaccines and Immunotherapeutics, 2020, 16, 2992-3000.	3.3	16
34	Genomic stability prevails in North-African hepatocellular carcinomas. Digestive and Liver Disease, 2007, 39, 671-677.	0.9	15
35	TP53 R72P polymorphism modulates DNA methylation in hepatocellular carcinoma. Molecular Cancer, 2015, 14, 74.	19.2	14
36	Serum DHCR24 Auto-antibody as a new Biomarker for Progression of Hepatitis C. EBioMedicine, 2015, 2, 604-612.	6.1	14

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37	Toll-like receptor 9 polymorphisms and Hepatitis B virus clearance in Moroccan chronic carriers. Gene, 2019, 687, 212-218.	2.2	14
38	Targeting Host Innate and Adaptive Immunity to Achieve the Functional Cure of Chronic Hepatitis B. Vaccines, 2020, 8, 216.	4.4	14
39	Impact of TP53 Codon 72 and MDM2 Promoter 309 Allelic Dosage in a Moroccan Population with Hepatocellular Carcinoma. International Journal of Biological Markers, 2011, 26, 229-233.	1.8	12
40	Associations of genetic variants in the transcriptional coactivators EP300 and PCAF with hepatocellular carcinoma. Cancer Epidemiology, 2012, 36, e300-e305.	1.9	12
41	A variant in the promoter of MBL2 is associated with protection against visceral leishmaniasis in Morocco. Infection, Genetics and Evolution, 2013, 13, 162-167.	2.3	12
42	The -94Ins/DelATTG polymorphism in NFÎºB1 promoter modulates chronic hepatitis C and liver disease progression. Infection, Genetics and Evolution, 2016, 39, 141-146.	2.3	12
43	Co-infections with hepatitis B and C viruses in human immunodeficiency virus-infected patients in Morocco. Clinical Microbiology and Infection, 2013, 19, E454-E457.	6.0	11
44	Programmed cell deathâ€1 3â€²â€untranslated region polymorphism is associated with spontaneous clearance of hepatitis B virus infection. Journal of Medical Virology, 2018, 90, 1730-1738.	5.0	11
45	Effect of MBOAT7 variant on hepatitis B and C infections in Moroccan patients. Scientific Reports, 2018, 8, 12247.	3.3	10
46	Moving towards hepatitis B elimination in Gulf Health Council states: From commitment to action. Journal of Infection and Public Health, 2020, 13, 221-227.	4.1	10
47	Serological evidence of West Nile virus infection in human populations and domestic birds in the Northwest of Morocco. Comparative Immunology, Microbiology and Infectious Diseases, 2021, 76, 101646.	1.6	9
48	Common polymorphic effectors of immunity against hepatitis B and C modulate susceptibility to infection and spontaneous clearance in a Moroccan population. Infection, Genetics and Evolution, 2014, 26, 1-7.	2.3	8
49	MDM2 285G>C and 344T>A gene variants and their association with hepatocellular carcinoma: a Moroccan caseâ€“control study. Infectious Agents and Cancer, 2014, 9, 11.	2.6	8
50	Molecular epidemiological study of adenovirus infecting western lowland gorillas and humans in and around Moukalaba-Doudou National Park (Gabon). Virus Genes, 2016, 52, 671-678.	1.6	8
51	Myxovirus resistance 1 gene polymorphisms and outcomes of viral hepatitis B and C infections in Moroccan patients. Journal of Medical Virology, 2017, 89, 647-652.	5.0	8
52	Nanobodies: an unexplored opportunity to combat COVID-19. Journal of Biomolecular Structure and Dynamics, 2022, 40, 3129-3131.	3.5	8
53	The Human papillomavirus among women living with Human Immunodeficiency Virus in Morocco A prospective cross-sectional study. Journal of Infection in Developing Countries, 2018, 12, 477-484.	1.2	8
54	Human genetic variation and the risk of hepatocellular carcinoma development. Hepatology International, 2013, 7, 820-831.	4.2	7

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55	Establishment of an intermittent cold stress model using <i>Tupaia belangeri</i> and evaluation of compound C737 targeting neuron-restrictive silencer factor. Experimental Animals, 2016, 65, 285-292.	1.1	7
56	Hepatitis B virus, hepatitis C virus and human immunodeficiency virus infections among people who inject drugs in Kuwait: A cross-sectional study. Scientific Reports, 2019, 9, 6292.	3.3	7
57	Immuno-informatics-based Identification of Novel Potential B Cell and T Cell Epitopes to Fight Zika Virus Infections. Infectious Disorders - Drug Targets, 2021, 21, 572-581.	0.8	7
58	Virtual Screening in Hepatitis B Virus Drug Discovery: Current Stateof- the-Art and Future Perspectives. Current Medicinal Chemistry, 2018, 25, 2709-2721.	2.4	7
59	Interleukin 28B rs12979860 genotype and Human Immunodeficiency Virus type 1: Susceptibility, AIDS development and therapeutic outcome. Human Immunology, 2018, 79, 70-75.	2.4	6
60	Diagnostic Performance of Automated SARS-CoV-2 Antigen Assay in Nasal Swab during COVID-19 Vaccination Campaign. Diagnostics, 2021, 11, 2110.	2.6	6
61	The allele 4 of neck region liver-lymph node-specific ICAM-3-grabbing integrin variant is associated with spontaneous clearance of hepatitis C virus and decrease of viral loads. Clinical Microbiology and Infection, 2014, 20, O325-O332.	6.0	5
62	Control of progression towards liver fibrosis and hepatocellular carcinoma by SOCS3 polymorphisms in chronic HCV-infected patients. Infection, Genetics and Evolution, 2018, 66, 1-8.	2.3	5
63	Non-primate hepacivirus transmission and prevalence: Novel findings of virus circulation in horses and dogs in Morocco. Infection, Genetics and Evolution, 2021, 93, 104975.	2.3	5
64	TP53 R72P Polymorphism and Susceptibility to Human Papillomavirus Infection Among Women With Human Immunodeficiency Virus in Morocco: A Case-control Study. Journal of Cancer Prevention, 2017, 22, 248-253.	2.0	5
65	Anti-SARS-CoV-2 Antibody Responses 5 Months Post Complete Vaccination of Moroccan Healthcare Workers. Vaccines, 2022, 10, 465.	4.4	5
66	Virus-associated human cancers in Moroccan population: From epidemiology to prospective research. Infection, Genetics and Evolution, 2019, 75, 103990.	2.3	4
67	An assessment of toll-like receptor 7 and 8 gene polymorphisms with susceptibility to HIV-1 infection, AIDS development and response to antiretroviral therapy. Immunology Letters, 2020, 227, 88-95.	2.5	4
68	An Integrative Gene Expression Microarray Meta-analysis Identifies Host Factors and Key Signatures Involved in Hepatitis B Virus Infection. Infectious Disorders - Drug Targets, 2020, 20, 698-707.	0.8	4
69	Design of a multi-epitope Zika virus vaccine candidate â€“ an in-silico study. Journal of Biomolecular Structure and Dynamics, 2023, 41, 3762-3771.	3.5	4
70	Reverse vaccinology-based prediction of a multi-epitope SARS-CoV-2 vaccine and its tailoring to new coronavirus variants. Journal of Biomolecular Structure and Dynamics, 2022, , 1-22.	3.5	4
71	Influence of mutation of the HFE gene on the progression of chronic viral hepatitis B and C in Moroccan patients. Journal of Medical Virology, 2011, 83, 2096-2102.	5.0	3
72	Amino acid substitutions in the Hepatitis C virus core region of genotype 1b in Moroccan patients. Infection, Genetics and Evolution, 2013, 14, 102-104.	2.3	3

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73	Gene expression analysis during acute hepatitis C virus infection associates dendritic cell activation with viral clearance. Journal of Medical Virology, 2016, 88, 843-851.	5.0	3
74	Blocking neddylation elicits antiviral effect against hepatitis B virus replication. Molecular Biology Reports, 2022, 49, 403-412.	2.3	3
75	Structure-guided discovery approach identifies potential lead compounds targeting Mpro of SARS-CoV-2. VirusDisease, 2020, 31, 549-553.	2.0	2
76	Effect of Peroxisome Proliferator-Activated Receptor-Î³ Coactivator-1 Alpha Variants on Spontaneous Clearance and Fibrosis Progression during Hepatitis C Virus Infection in Moroccan Patients. Virologica Sinica, 2020, 35, 566-574.	3.0	2
77	Programmed cell death-1 single-nucleotide polymorphism rs10204525 is associated with human immunodeficiency virus type 1 RNA viral load in HIV-1-infected Moroccan subjects. Medical Microbiology and Immunology, 2021, 210, 187-196.	4.8	2
78	IFNL4 rs12979860 polymorphism influences HBV DNA viral loads but not the outcome of HBV infection in Moroccan patients. Microbes and Infection, 2021, 23, 104802.	1.9	2
79	Prostate-specific Antigen Levels in Moroccan Diabetic Males: A Cross-sectional Study. Current Diabetes Reviews, 2018, 14, 286-290.	1.3	2
80	World Society for Virology first international conference: Tackling global virus epidemics. Virology, 2022, 566, 114-121.	2.4	2
81	Severe acute respiratory syndrome coronavirus 2 seroprevalence survey among 10,256 workers in Kuwait. Journal of Clinical Virology Plus, 2021, 1, 100017.	1.0	1
82	Supplementing Conventional Treatment with PycnogenolÂ® May Improve Hepatitis C Virusâ€“Associated Type 2 Diabetes: A Mini Review. Journal of Clinical and Translational Hepatology, 2016, 4, 228-233.	1.4	1
83	Genetic variability of Hepatitis C Virus in Moroccan population. Retrovirology, 2012, 9, .	2.0	0
84	Lack of Association between IFNL3 Polymorphism and Human Papillomavirus Infection and Their Progression in HIV-Infected Women Receiving Antiretroviral Treatment. Pathobiology, 2020, 87, 262-267.	3.8	0
85	Molecular and computational analysis of natural drug resistance mutations among Moroccan chronic hepatitis B carriers. Gene Reports, 2021, 23, 101197.	0.8	0