

# Sayeh Ezzikouri

## List of Publications by Year in descending order

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Version: 2024-02-01

85  
papers

1,339  
citations

331670

21  
h-index

454955

30  
g-index

86  
all docs

86  
docs citations

86  
times ranked

2044  
citing authors

#	ARTICLE	IF	CITATIONS
1	Design of a multi-epitope Zika virus vaccine candidate – an <i>in-silico</i> study. <i>Journal of Biomolecular Structure and Dynamics</i> , 2023, 41, 3762-3771.	3.5	4
2	Nanobodies: an unexplored opportunity to combat COVID-19. <i>Journal of Biomolecular Structure and Dynamics</i> , 2022, 40, 3129-3131.	3.5	8
3	Blocking neddylation elicits antiviral effect against hepatitis B virus replication. <i>Molecular Biology Reports</i> , 2022, 49, 403-412.	2.3	3
4	World Society for Virology first international conference: Tackling global virus epidemics. <i>Virology</i> , 2022, 566, 114-121.	2.4	2
5	Anti-SARS-CoV-2 Antibody Responses 5 Months Post Complete Vaccination of Moroccan Healthcare Workers. <i>Vaccines</i> , 2022, 10, 465.	4.4	5
6	Reverse vaccinology-based prediction of a multi-epitope SARS-CoV-2 vaccine and its tailoring to new coronavirus variants. <i>Journal of Biomolecular Structure and Dynamics</i> , 2022, , 1-22.	3.5	4
7	How artificial intelligence may help the Covid-19 pandemic: Pitfalls and lessons for the future. <i>Reviews in Medical Virology</i> , 2021, 31, 1-11.	8.3	53
8	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. <i>Medical Microbiology and Immunology</i> , 2021, 210, 187-196.	4.8	2
9	IFNL4 rs12979860 polymorphism influences HBV DNA viral loads but not the outcome of HBV infection in Moroccan patients. <i>Microbes and Infection</i> , 2021, 23, 104802.	1.9	2
10	Subnational mapping of HIV incidence and mortality among individuals aged 15–49 years in sub-Saharan Africa, 2000–18: a modelling study. <i>Lancet HIV</i> , 2021, 8, e363-e375.	4.7	32
11	Serological evidence of West Nile virus infection in human populations and domestic birds in the Northwest of Morocco. <i>Comparative Immunology, Microbiology and Infectious Diseases</i> , 2021, 76, 101646.	1.6	9
12	Severe acute respiratory syndrome coronavirus 2 seroprevalence survey among 10,256 workers in Kuwait. <i>Journal of Clinical Virology Plus</i> , 2021, 1, 100017.	1.0	1
13	Molecular and computational analysis of natural drug resistance mutations among Moroccan chronic hepatitis B carriers. <i>Gene Reports</i> , 2021, 23, 101197.	0.8	0
14	Immuno-informatics-based Identification of Novel Potential B Cell and T Cell Epitopes to Fight Zika Virus Infections. <i>Infectious Disorders - Drug Targets</i> , 2021, 21, 572-581.	0.8	7
15	Non-primate hepacivirus transmission and prevalence: Novel findings of virus circulation in horses and dogs in Morocco. <i>Infection, Genetics and Evolution</i> , 2021, 93, 104975.	2.3	5
16	Global, regional, and national sex-specific burden and control of the HIV epidemic, 1990–2019, for 204 countries and territories: the Global Burden of Diseases Study 2019. <i>Lancet HIV</i> , 2021, 8, e633-e651.	4.7	56
17	Diagnostic Performance of Automated SARS-CoV-2 Antigen Assay in Nasal Swab during COVID-19 Vaccination Campaign. <i>Diagnostics</i> , 2021, 11, 2110.	2.6	6
18	Moving towards hepatitis B elimination in Gulf Health Council states: From commitment to action. <i>Journal of Infection and Public Health</i> , 2020, 13, 221-227.	4.1	10

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19	Saliva specimens for detection of severe acute respiratory syndrome coronavirus 2 in Kuwait: A cross-sectional study. <i>Journal of Clinical Virology</i> , 2020, 132, 104652.	3.1	49
20	An assessment of toll-like receptor 7 and 8 gene polymorphisms with susceptibility to HIV-1 infection, AIDS development and response to antiretroviral therapy. <i>Immunology Letters</i> , 2020, 227, 88-95.	2.5	4
21	Structure-guided discovery approach identifies potential lead compounds targeting Mpro of SARS-CoV-2. <i>VirusDisease</i> , 2020, 31, 549-553.	2.0	2
22	Coronavirus disease 2019â€”Historical context, virology, pathogenesis, immunotherapy, and vaccine development. <i>Human Vaccines and Immunotherapeutics</i> , 2020, 16, 2992-3000.	3.3	16
23	Lack of Association between IFNL3 Polymorphism and Human Papillomavirus Infection and Their Progression in HIV-Infected Women Receiving Antiretroviral Treatment. <i>Pathobiology</i> , 2020, 87, 262-267.	3.8	0
24	Targeting Host Innate and Adaptive Immunity to Achieve the Functional Cure of Chronic Hepatitis B. <i>Vaccines</i> , 2020, 8, 216.	4.4	14
25	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. <i>Virologica Sinica</i> , 2020, 35, 566-574.	3.0	2
26	An Integrative Gene Expression Microarray Meta-analysis Identifies Host Factors and Key Signatures Involved in Hepatitis B Virus Infection. <i>Infectious Disorders - Drug Targets</i> , 2020, 20, 698-707.	0.8	4
27	Virus-associated human cancers in Moroccan population: From epidemiology to prospective research. <i>Infection, Genetics and Evolution</i> , 2019, 75, 103990.	2.3	4
28	Hepatitis B virus, hepatitis C virus and human immunodeficiency virus infections among people who inject drugs in Kuwait: A cross-sectional study. <i>Scientific Reports</i> , 2019, 9, 6292.	3.3	7
29	Toll-like receptor 9 polymorphisms and Hepatitis B virus clearance in Moroccan chronic carriers. <i>Gene</i> , 2019, 687, 212-218.	2.2	14
30	Genetic variations in toll-like receptors 7 and 8 modulate natural hepatitis C outcomes and liver disease progression. <i>Liver International</i> , 2018, 38, 432-442.	3.9	33
31	Interleukin 28B rs12979860 genotype and Human Immunodeficiency Virus type 1: Susceptibility, AIDS development and therapeutic outcome. <i>Human Immunology</i> , 2018, 79, 70-75.	2.4	6
32	Effect of MBOAT7 variant on hepatitis B and C infections in Moroccan patients. <i>Scientific Reports</i> , 2018, 8, 12247.	3.3	10
33	Control of progression towards liver fibrosis and hepatocellular carcinoma by SOCS3 polymorphisms in chronic HCV-infected patients. <i>Infection, Genetics and Evolution</i> , 2018, 66, 1-8.	2.3	5
34	Programmed cell deathâ€”1 3â€”untranslated region polymorphism is associated with spontaneous clearance of hepatitis B virus infection. <i>Journal of Medical Virology</i> , 2018, 90, 1730-1738.	5.0	11
35	Virtual Screening in Hepatitis B Virus Drug Discovery: Current State-of-the-Art and Future Perspectives. <i>Current Medicinal Chemistry</i> , 2018, 25, 2709-2721.	2.4	7
36	Prostate-specific Antigen Levels in Moroccan Diabetic Males: A Cross-sectional Study. <i>Current Diabetes Reviews</i> , 2018, 14, 286-290.	1.3	2

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37	The Human papillomavirus among women living with Human Immunodeficiency Virus in Morocco A prospective cross-sectional study. <i>Journal of Infection in Developing Countries</i> , 2018, 12, 477-484.	1.2	8
38	Lack of Ser267Phe variant of sodium taurocholate cotransporting polypeptide among Moroccans regardless of hepatitis B virus infection status. <i>BMC Infectious Diseases</i> , 2017, 17, 99.	2.9	17
39	Interferon- $\gamma$ response is impaired by hepatitis B virus infection in <i>Tupaia belangeri</i> . <i>Virus Research</i> , 2017, 237, 47-57.	2.2	17
40	Oxidative Stress and Immune Responses During Hepatitis C Virus Infection in <i>Tupaia belangeri</i> . <i>Scientific Reports</i> , 2017, 7, 9848.	3.3	18
41	Myxovirus resistance 1 gene polymorphisms and outcomes of viral hepatitis B and C infections in Moroccan patients. <i>Journal of Medical Virology</i> , 2017, 89, 647-652.	5.0	8
42	TP53 R72P Polymorphism and Susceptibility to Human Papillomavirus Infection Among Women With Human Immunodeficiency Virus in Morocco: A Case-control Study. <i>Journal of Cancer Prevention</i> , 2017, 22, 248-253.	2.0	5
43	Gene expression analysis during acute hepatitis C virus infection associates dendritic cell activation with viral clearance. <i>Journal of Medical Virology</i> , 2016, 88, 843-851.	5.0	3
44	Establishment of an intermittent cold stress model using <i>Tupaia belangeri</i> and evaluation of compound C737 targeting neuron-restrictive silencer factor. <i>Experimental Animals</i> , 2016, 65, 285-292.	1.1	7
45	Molecular epidemiological study of adenovirus infecting western lowland gorillas and humans in and around Moukalaba-Doudou National Park (Gabon). <i>Virus Genes</i> , 2016, 52, 671-678.	1.6	8
46	The -94Ins/DelATTG polymorphism in NF- $\kappa$ B1 promoter modulates chronic hepatitis C and liver disease progression. <i>Infection, Genetics and Evolution</i> , 2016, 39, 141-146.	2.3	12
47	Property of hepatitis B virus replication in <i>Tupaia belangeri</i> hepatocytes. <i>Biochemical and Biophysical Research Communications</i> , 2016, 469, 229-235.	2.1	21
48	Supplementing Conventional Treatment with Pycnogenol <sup>®</sup> May Improve Hepatitis C Virus-Associated Type 2 Diabetes: A Mini Review. <i>Journal of Clinical and Translational Hepatology</i> , 2016, 4, 228-233.	1.4	1
49	Inhibitory effects of Pycnogenol <sup>®</sup> on hepatitis C virus replication. <i>Antiviral Research</i> , 2015, 113, 93-102.	4.1	19
50	TP53 R72P polymorphism modulates DNA methylation in hepatocellular carcinoma. <i>Molecular Cancer</i> , 2015, 14, 74.	19.2	14
51	Serum DHCR24 Auto-antibody as a new Biomarker for Progression of Hepatitis C. <i>EBioMedicine</i> , 2015, 2, 604-612.	6.1	14
52	MicroRNAs as Important Players in Host-hepatitis B Virus Interactions. <i>Journal of Clinical and Translational Hepatology</i> , 2015, 3, 149-61.	1.4	17
53	Occult HBV infection in Morocco: from chronic hepatitis to hepatocellular carcinoma. <i>Liver International</i> , 2014, 34, e144-50.	3.9	21
54	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. <i>Clinical Microbiology and Infection</i> , 2014, 20, O325-O332.	6.0	5

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55	The adiponutrin I148M variant is a risk factor for HCV-associated liver cancer in North-African patients. <i>Infection, Genetics and Evolution</i> , 2014, 21, 179-183.	2.3	20
56	Recent insights into hepatitis B virus–host interactions. <i>Journal of Medical Virology</i> , 2014, 86, 925-932.	5.0	22
57	Common polymorphic effectors of immunity against hepatitis B and C modulate susceptibility to infection and spontaneous clearance in a Moroccan population. <i>Infection, Genetics and Evolution</i> , 2014, 26, 1-7.	2.3	8
58	MDM2 285G>C and 344T>A gene variants and their association with hepatocellular carcinoma: a Moroccan case–control study. <i>Infectious Agents and Cancer</i> , 2014, 9, 11.	2.6	8
59	Hepatitis C virus infection in the Maghreb region. <i>Journal of Medical Virology</i> , 2013, 85, 1542-1549.	5.0	22
60	Prevalence and risk factors of hepatitis B and C virus infections among the general population and blood donors in Morocco. <i>BMC Public Health</i> , 2013, 13, 50.	2.9	78
61	Human genetic variation and the risk of hepatocellular carcinoma development. <i>Hepatology International</i> , 2013, 7, 820-831.	4.2	7
62	Amino acid substitutions in the Hepatitis C virus core region of genotype 1b in Moroccan patients. <i>Infection, Genetics and Evolution</i> , 2013, 14, 102-104.	2.3	3
63	A variant in the promoter of MBL2 is associated with protection against visceral leishmaniasis in Morocco. <i>Infection, Genetics and Evolution</i> , 2013, 13, 162-167.	2.3	12
64	Co-infections with hepatitis B and C viruses in human immunodeficiency virus-infected patients in Morocco. <i>Clinical Microbiology and Infection</i> , 2013, 19, E454-E457.	6.0	11
65	Polymorphic APOBEC3 modulates chronic hepatitis B in Moroccan population. <i>Journal of Viral Hepatitis</i> , 2013, 20, 678-686.	2.0	23
66	Hepatitis B virus in the Maghreb Region: from epidemiology to prospective research. <i>Liver International</i> , 2013, 33, 811-819.	3.9	29
67	Genetic Variation in the Interleukin-28B Gene Is Associated with Spontaneous Clearance and Progression of Hepatitis C Virus in Moroccan Patients. <i>PLoS ONE</i> , 2013, 8, e54793.	2.5	33
68	Associations of genetic variants in the transcriptional coactivators EP300 and PCAF with hepatocellular carcinoma. <i>Cancer Epidemiology</i> , 2012, 36, e300-e305.	1.9	12
69	Variability in the Precore and Core Promoter Regions of HBV Strains in Morocco: Characterization and Impact on Liver Disease Progression. <i>PLoS ONE</i> , 2012, 7, e42891.	2.5	33
70	Morocco underwent a drift of circulating hepatitis C virus subtypes in recent decades. <i>Archives of Virology</i> , 2012, 157, 515-520.	2.1	29
71	Genetic variability of Hepatitis C Virus in Moroccan population. <i>Retrovirology</i> , 2012, 9, .	2.0	0
72	First multicenter study for risk factors for hepatocellular carcinoma development in North Africa. <i>World Journal of Hepatology</i> , 2011, 3, 24.	2.0	35

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73	Impact of TP53 Codon 72 and MDM2 Promoter 309 Allelic Dosage in a Moroccan Population with Hepatocellular Carcinoma. <i>International Journal of Biological Markers</i> , 2011, 26, 229-233.	1.8	12
74	Influence of mutation of the <i>HFE</i> gene on the progression of chronic viral hepatitis B and C in Moroccan patients. <i>Journal of Medical Virology</i> , 2011, 83, 2096-2102.	5.0	3
75	Hepatitis B genotypes/subgenotypes and MHR variants among Moroccan chronic carriers. <i>Journal of Infection</i> , 2011, 63, 66-75.	3.3	40
76	Polymorphisms in antioxidant defence genes and susceptibility to hepatocellular carcinoma in a Moroccan population. <i>Free Radical Research</i> , 2010, 44, 208-216.	3.3	40
77	MDM2 SNP309T>G polymorphism and risk of hepatocellular carcinoma: A case-control analysis in a Moroccan population. <i>Cancer Detection and Prevention</i> , 2009, 32, 380-385.	2.1	32
78	Single nucleotide polymorphism in DNMT3B promoter and its association with hepatocellular carcinoma in a Moroccan population. <i>Infection, Genetics and Evolution</i> , 2009, 9, 877-881.	2.3	28
79	The prevalence of resistance-associated mutations to protease and reverse transcriptase inhibitors in treatment-naïve (HIV1)-infected individuals in Casablanca, Morocco. <i>Journal of Infection in Developing Countries</i> , 2009, 3, 380-91.	1.2	23
80	Genotype determination in Moroccan hepatitis B chronic carriers. <i>Infection, Genetics and Evolution</i> , 2008, 8, 306-312.	2.3	29
81	Prevalence of Common HFE and SERPINA1 Mutations in Patients with Hepatocellular Carcinoma in a Moroccan Population. <i>Archives of Medical Research</i> , 2008, 39, 236-241.	3.3	19
82	Genetic polymorphism in the manganese superoxide dismutase gene is associated with an increased risk for hepatocellular carcinoma in HCV-infected Moroccan patients. <i>Mutation Research - Genetic Toxicology and Environmental Mutagenesis</i> , 2008, 649, 1-6.	1.7	30
83	The Pro variant of the p53 codon 72 polymorphism is associated with hepatocellular carcinoma in Moroccan population. <i>Hepatology Research</i> , 2007, 37, 748-754.	3.4	46
84	Genomic stability prevails in North-African hepatocellular carcinomas. <i>Digestive and Liver Disease</i> , 2007, 39, 671-677.	0.9	15
85	HCV genotypes in Morocco. , 1997, 52, 396-398.		23