Pascal Pineau

List of Publications by Year
in descending order

100 papers	4,922 citations	172457 29 h-index	95266 68 g-index
103 all docs	103 docs citations	103 times ranked	7685 citing authors

#	Article	IF	Citations
1	A preoperative nomogram for predicting long-term survival after resection of large hepatocellular carcinoma (>10Âcm). Hpb, 2022, 24, 192-201.	0.3	10
2	APOBEC3C S188I Polymorphism Enhances Context-Specific Editing of Hepatitis B Virus Genome. Journal of Infectious Diseases, 2022, 226, 891-895.	4.0	1
3	Natural Occurrence of Mycotoxin-Producing Fusaria in Market-Bought Peruvian Cereals: A Food Safety Threat for Andean Populations. Toxins, 2021, 13, 172.	3.4	13
4	The Threat of Multiple Liver Carcinogens in the Population of Laos: A Review. Livers, 2021, 1, 49-59.	1.9	1
5	Global DNA hypermethylation pattern and unique gene expression signature in liver cancer from patients with Indigenous American ancestry. Oncotarget, 2021, 12, 475-492.	1.8	16
6	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
7	Metallomic profile in non-cirrhotic hepatocellular carcinoma supports a phenomenon of metal metabolism adaptation in tumor cells. Scientific Reports, 2021, 11, 14195.	3.3	5
8	ddPCR increases detection of SARS-CoV-2 RNA in patients with low viral loads. Archives of Virology, 2021, 166, 2529-2540.	2.1	10
9	Hepatitis viruses take advantage of traditional practices to increase the burden of hepatocellular carcinoma in Tunisia. Archives of Virology, 2020, 165, 33-42.	2.1	7
10	Identification of Four Immune Subtypes Characterized by Distinct Composition and Functions of Tumor Microenvironment in Intrahepatic Cholangiocarcinoma. Hepatology, 2020, 72, 965-981.	7.3	159
11	The role of hepatitis C virus genotypes and core mutations in hepatocellular carcinoma in Cameroon. Journal of Viral Hepatitis, 2020, 27, 880-885.	2.0	1
12	Virus-associated human cancers in Moroccan population: From epidemiology to prospective research. Infection, Genetics and Evolution, 2019, 75, 103990.	2.3	4
13	Enrichment in selected genotypes, basal core and precore mutations of hepatitis B virus in patients with hepatocellular carcinoma in Cameroon. Journal of Viral Hepatitis, 2019, 26, 1086-1093.	2.0	10
14	Hepatocellular carcinoma: Clinical-pathological features and HIV infection in Mozambican patients,. Cancer Treatment and Research Communications, 2019, 19, 100129.	1.7	5
15	Wide Sexual Dimorphism of Hepatocellular Carcinoma Presentation in Algeria. Gastrointestinal Tumors, 2019, 6, 122-136.	0.7	2
16	Toll-like receptor 9 polymorphisms and Hepatitis B virus clearance in Moroccan chronic carriers. Gene, 2019, 687, 212-218.	2.2	14
17	Profile of hepatocellular carcinoma in the Republic of Moldova: first-hand information on the presentation, distribution and etiologies. Romanian Journal of Internal Medicine = Revue Roumaine De Medecine Interne, 2019, 57, 37-46.	0.6	5
18	On hepatocellular carcinoma in South America and early-age onset of the disease. Clinics and Research in Hepatology and Gastroenterology, 2019, 43, 522-526.	1.5	13

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19	Increased liver injury in patients with chronic hepatitis and IgG directed against hepatitis E virus. EXCLI Journal, 2019, 18, 955-961.	0.7	3
20	Circulating Aflatoxin B1-Related TP53 Mutation Detected by Digital PCR in Tunisian Patients with and Without Hepatocellular Carcinoma. Hepatitis Monthly, 2019, In Press, .	0.2	1
21	Herbal Medicine Practices of Patients With Liver Cancer in Peru: A Comprehensive Study Toward Integrative Cancer Management. Integrative Cancer Therapies, 2018, 17, 52-64.	2.0	24
22	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
23	Effect of MBOAT7 variant on hepatitis B and C infections in Moroccan patients. Scientific Reports, 2018, 8, 12247.	3.3	10
24	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
25	A Seven-Year Retrospective Study on the Surveillance of Hepatitis B in Laos. International Journal of Hepatology, 2018, 2018, 1-11.	1.1	7
26	Liver clear cell foci and viral infection are associated with non-cirrhotic, non-fibrolamellar hepatocellular carcinoma in young patients from South America. Scientific Reports, 2018, 8, 9945.	3.3	7
27	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
28	Droplet digital PCR detects high rate of TP53 R249S mutants in cell-free DNA of middle African patients with hepatocellular carcinoma. Clinical and Experimental Medicine, 2018, 18, 421-431.	3.6	31
29	Early-onset liver cancer in South America associates with low hepatitis B virus DNA burden. Scientific Reports, 2018, 8, 12031.	3.3	23
30	Hepatitis C in Laos: A 7-Year Retrospective Study on 1765 Patients. Virologica Sinica, 2018, 33, 295-303.	3.0	3
31	Widespread geographical disparities in chronic hepatitis B virus infection in Algeria. Archives of Virology, 2017, 162, 1641-1648.	2.1	5
32	Hepatitis E virus infection as a promoting factor for hepatocellular carcinoma in Cameroon: Preliminary Observations. International Journal of Infectious Diseases, 2017, 64, 4-8.	3.3	25
33	<scp>UGT</scp>1As and predisposition to liver cancer: Still important, more elusive. Liver International, 2017, 37, 1284-1286.	3.9	1
34	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
35	Evolution of hepatocellular carcinoma epidemiology in CÃ´te dâ€™Ivoire. Bulletin Du Cancer, 2017, 104, 937-945.	1.6	3
36	A 13-Year Retrospective Study on Primary Liver Cancer in Cambodia: A Strikingly High Hepatitis C Occurrence among Hepatocellular Carcinoma Cases. Oncology, 2016, 91, 106-116.	1.9	14

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37	A prominent role of Hepatitis D Virus in liver cancers documented in Central Africa. BMC Infectious Diseases, 2016, 16, 647.	2.9	35
38	Eastern region represents a worrying cluster of active hepatitis C in Algeria in 2012. Journal of Medical Virology, 2016, 88, 1394-1403.	5.0	7
39	APOBEC3DE Antagonizes Hepatitis B Virus Restriction Factors APOBEC3F and APOBEC3G. Journal of Molecular Biology, 2016, 428, 3514-3528.	4.2	19
40	Hepatocellular carcinoma surgery outcomes in the developing world: A 20-year retrospective cohort study at the National Cancer Institute of Peru. Heliyon, 2016, 2, e00052.	3.2	20
41	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
42	A major shift of viral and nutritional risk factors affects the hepatocellular carcinoma risk among Ivorian patients: a preliminary report. Infectious Agents and Cancer, 2015, 10, 18.	2.6	6
43	Wnt/Î²-catenin signaling pathway in hepatocellular carcinomas cases from Colombia. Annals of Hepatology, 2015, 14, 64-74.	1.5	32
44	Mutation spectrum of hepatocellular carcinoma from eastern-European patients betrays the impact of a complex exposome. Journal of Exposure Science and Environmental Epidemiology, 2015, 25, 256-263.	3.9	12
45	TP53 R72P polymorphism modulates DNA methylation in hepatocellular carcinoma. Molecular Cancer, 2015, 14, 74.	19.2	14
46	Relief and worries about hepatitis C in sub-Saharan Africa. Lancet Infectious Diseases, The, 2015, 15, 754-755.	9.1	2
47	Long noncoding RNA PANDA and scaffold-attachment-factor SAFA control senescence entry and exit. Nature Communications, 2014, 5, 5323.	12.8	164
48	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
49	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
50	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
51	Influence of hepatitis viruses on clinico-pathological profiles and long-term outcome in patients undergoing surgery for hepatocellular carcinoma. Hepatobiliary and Pancreatic Diseases International, 2014, 13, 162-172.	1.3	10
52	A Peculiar Mutation Spectrum Emerging from Young Peruvian Patients with Hepatocellular Carcinoma. PLoS ONE, 2014, 9, e114912.	2.5	28
53	Hepatitis C virus infection in the Maghreb region. Journal of Medical Virology, 2013, 85, 1542-1549.	5.0	22
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	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
56	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
57	Telomere protection and TRF2 expression are enhanced by the canonical Wnt signalling pathway. EMBO Reports, 2013, 14, 356-363.	4.5	72
58	Somatic changes in primary liver cancer in Russia: A pilot study. Mutation Research - Genetic Toxicology and Environmental Mutagenesis, 2013, 755, 90-99.	1.7	11
59	Hepatitis B virus in the Maghreb Region: from epidemiology to prospective research. Liver International, 2013, 33, 811-819.	3.9	29
60	An Atypical Age-Specific Pattern of Hepatocellular Carcinoma in Peru: A Threat for Andean Populations. PLoS ONE, 2013, 8, e67756.	2.5	28
61	Insertion/deletion polymorphisms are convenient and reliable markers to assess chromosomal instability in human tumors. International Journal of Biological Markers, 2012, 27, 232-240.	1.8	2
62	Associations of genetic variants in the transcriptional coactivators EP300 and PCAF with hepatocellular carcinoma. Cancer Epidemiology, 2012, 36, e300-e305.	1.9	12
63	Morocco underwent a drift of circulating hepatitis C virus subtypes in recent decades. Archives of Virology, 2012, 157, 515-520.	2.1	29
64	p53 regulates epithelialâ€“mesenchymal transition through microRNAs targeting ZEB1 and ZEB2. Journal of Experimental Medicine, 2011, 208, 875-883.	8.5	480
65	First multicenter study for risk factors for hepatocellular carcinoma development in North Africa. World Journal of Hepatology, 2011, 3, 24.	2.0	35
66	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
67	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
68	Somatic hypermutation of human mitochondrial and nuclear DNA by APOBEC3 cytidine deaminases, a pathway for DNA catabolism. Proceedings of the National Academy of Sciences of the United States of America, 2011, 108, 4858-4863.	7.1	142
69	p53 regulates epithelialâ€“mesenchymal transition through microRNAs targeting ZEB1 and ZEB2. Journal of Cell Biology, 2011, 193, i8-i8.	5.2	0
70	Reprogramming of miRNA networks in cancer and leukemia. Genome Research, 2010, 20, 589-599.	5.5	331
71	High chromosome instability and occasional R249S TP53 Mutation characterize hepatocellular carcinoma in Romania. Cancer Genetics and Cytogenetics, 2010, 203, 73.	1.0	1
72	Massive APOBEC3 Editing of Hepatitis B Viral DNA in Cirrhosis. PLoS Pathogens, 2010, 6, e1000928.	4.7	145

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73	miR-221 overexpression contributes to liver tumorigenesis. Proceedings of the National Academy of Sciences of the United States of America, 2010, 107, 264-269.	7.1	679
74	Polymorphisms in antioxidant defence genes and susceptibility to hepatocellular carcinoma in a Moroccan population. Free Radical Research, 2010, 44, 208-216.	3.3	40
75	The dynamic DNA methylomes of double-stranded DNA viruses associated with human cancer. Genome Research, 2009, 19, 438-451.	5.5	218
76	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
77	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
78	Mutational analysis of TP53, PTEN, PIK3CA and CTNNB1/Â-catenin genes in human herpesvirus 8-associated primary effusion lymphoma. Haematologica, 2009, 94, 1170-1174.	3.5	21
79	Genotype determination in Moroccan hepatitis B chronic carriers. Infection, Genetics and Evolution, 2008, 8, 306-312.	2.3	29
80	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
81	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
82	Chromosome instability in human hepatocellular carcinoma depends on p53 status and aflatoxin exposure. Mutation Research - Genetic Toxicology and Environmental Mutagenesis, 2008, 653, 6-13.	1.7	50
83	Tbx3 Is a Downstream Target of the Wnt/Î²-Catenin Pathway and a Critical Mediator of Î²-Catenin Survival Functions in Liver Cancer. Cancer Research, 2007, 67, 901-910.	0.9	147
84	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
85	Genomic stability prevails in North-African hepatocellular carcinomas. Digestive and Liver Disease, 2007, 39, 671-677.	0.9	15
86	A Universal Primer Set for PCR Amplification of Nuclear Histone H4 Genes from All Animal Species. Molecular Biology and Evolution, 2005, 22, 582-588.	8.9	50
87	Homozygous deletion scanning in hepatobiliary tumor cell lines reveals alternative pathways for liver carcinogenesis. Hepatology, 2003, 37, 852-861.	7.3	48
88	Homozygous deletions scanning in tumor cell lines detects previously unsuspected loci. International Journal of Cancer, 2003, 106, 216-223.	5.1	10
89	Mutation analysis of novel human liver-related putative tumor suppressor gene in hepatocellular carcinoma. World Journal of Gastroenterology, 2003, 9, 89.	3.3	9
90	Identification of a Hepatitis B Virus Genome in Wild Chimpanzees (Pan troglodytes schweinfurthi) from East Africa Indicates a Wide Geographical Dispersion among Equatorial African Primates. Journal of Virology, 2002, 76, 11155-11158.	3.4	33

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91	Animal hepadnaviruses and their host species: Models for human hepatocarcinogenesis. Perspectives in Medical Virology, 2002, 6, 123-141.	0.1	1
92	Distinct chromosomal abnormality pattern in primary liver cancer of non-B, non-C patients. Oncogene, 2000, 19, 3733-3738.	5.9	107
93	Hepatocellular carcinoma occurring in nonfibrotic liver: Epidemiologic and histopathologic analysis of 80 French cases. Hepatology, 2000, 32, 200-204.	7.3	218
94	Effect of TT Virus Infection on Hepatocellular Carcinoma Development: Results of a Euroâ€Asian Survey. Journal of Infectious Diseases, 2000, 181, 1138-1142.	4.0	35
95	Studies of genetic defects in hepatocellular carcinoma: recent outcomes and new challenges. Journal of Hepatology, 2000, 33, 152-156.	3.7	10
96	Identification of three distinct regions of allelic deletions on the short arm of chromosome 8 in hepatocellular carcinoma. Oncogene, 1999, 18, 3127-3134.	5.9	107
97	Close correlation between Î²-catenin gene alterations and nuclear accumulation of the protein in human hepatocellular carcinomas. Oncogene, 1999, 18, 6583-6588.	5.9	119
98	Extensive analysis of duplicated-inverted hepatitis B virus integrations in human hepatocellular carcinoma.. Journal of General Virology, 1998, 79, 591-600.	2.9	40
99	Comprehensive allelotyping of human hepatocellular carcinoma. Oncogene, 1997, 14, 2927-2933.	5.9	273
100	Recurrent chromosomal abnormalities in hepatocellular carcinoma detected by comparative genomic hybridization. Genes Chromosomes and Cancer, 1997, 18, 59-65.	2.8	209