List of Publications by Year in descending order

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		126907	95266
116	6,134	33	68
papers	citations	h-index	g-index
123	123	123	9553
all docs	docs citations	times ranked	citing authors

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#	Article	IF	CITATIONS
1	Lower probability of persistence of total anti-SARS-CoV-2 antibodies after COVID-19 among people living with HIV. Clinical Microbiology and Infection, 2022, , .	6.0	4
2	New insights into the genetic etiology of Alzheimer's disease and related dementias. Nature Genetics, 2022, 54, 412-436.	21.4	700
3	<i>IFNL4</i> genotype influences the rate of HIV-1 seroconversion in men who have sex with men. Virulence, 2022, 13, 757-763.	4.4	0
4	Association of Rare <i>APOE</i> Missense Variants V236E and R251G With Risk of Alzheimer Disease. JAMA Neurology, 2022, 79, 652.	9.0	31
5	High efficacy of glecaprevir/pibrentasvir for HCV-infected individuals with active drug use. Journal of Infection, 2022, 85, 322-326.	3.3	4
6	Human Immunodeficiency Virus (HIV) Infection Is Associated With Lower Risk of Hepatocellular Carcinoma After Sustained Virological Response to Direct-acting Antivirals in Hepatitis C Infected Patients With Advanced Fibrosis. Clinical Infectious Diseases, 2021, 73, e2109-e2116.	5.8	10
7	Liver Stiffness–Based Strategies Predict Absence of Variceal Bleeding in Cirrhotic Hepatitis C Virus–Infected Patients With and Without Human Immunodeficiency Virus Coinfection After Sustained Virological Response. Clinical Infectious Diseases, 2021, 72, e96-e102.	5.8	4
8	Long runs of homozygosity are associated with Alzheimer's disease. Translational Psychiatry, 2021, 11, 142.	4.8	6
9	Impact of recent drug use on the efficacy of elbasvir/grazoprevir for HCVâ€infected people on opioid agonist therapy. Journal of Viral Hepatitis, 2021, 28, 878-886.	2.0	0
10	Multiomics integrative analysis identifies APOE allele-specific blood biomarkers associated to Alzheimer's disease etiopathogenesis. Aging, 2021, 13, 9277-9329.	3.1	15
11	Similar incidence of coronavirus disease 2019 (COVID-19) in patients with rheumatic diseases with and without hydroxychloroquine therapy. PLoS ONE, 2021, 16, e0249036.	2.5	7
12	Kinetics of emergence of liver complications in hepatitis C virus infected patients and advanced fibrosis, with and without HIV-coinfection, after sustained virological response. Aids, 2021, 35, 2119-2127.	2.2	4
13	Common variants in Alzheimer's disease and risk stratification by polygenic risk scores. Nature Communications, 2021, 12, 3417.	12.8	140
14	A Genome-Wide Association Study on Liver Stiffness Changes during Hepatitis C Virus Infection Cure. Diagnostics, 2021, 11, 1501.	2.6	2
15	Increased risk of severe clinical course of COVID-19 in carriers of HLA-C*04:01. EClinicalMedicine, 2021, 40, 101099.	7.1	52
16	CD46 Genetic Variability and HIV-1 Infection Susceptibility. Cells, 2021, 10, 3094.	4.1	3
17	Genomic Characterization of Host Factors Related to SARS-CoV-2 Infection in People with Dementia and Control Populations: The GR@ACE/DEGESCO Study. Journal of Personalized Medicine, 2021, 11, 1318.	2.5	7
18	A polygenic risk score for mosaic loss of chromosome Y susceptibility is associated with higher risk of MCI to AD conversion Alzheimer's and Dementia, 2021, 17 Suppl 3, e053745.	0.8	0

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19	Association of complement C3d receptor 2 genotypes with the acquisition of HIV infection in a trial of recombinant glycoprotein 120 vaccine. Aids, 2020, 34, 25-32.	2.2	9
20	Effects of first-line antiretroviral therapy on the CD4/CD8 ratio and CD8 cell counts in CoRIS: a prospective multicentre cohort study. Lancet HIV,the, 2020, 7, e565-e573.	4.7	42
21	The Ubiquitin Proteasome System in Neuromuscular Disorders: Moving Beyond Movement. International Journal of Molecular Sciences, 2020, 21, 6429.	4.1	17
22	<scp>CCR</scp> 5 deficiency impairs <scp>CD</scp> 4 ⁺ Tâ€cell memory responses and antigenic sensitivity through increased ceramide synthesis. EMBO Journal, 2020, 39, e104749.	7.8	17
23	Genomeâ€wide association analysis of dementia and its clinical endophenotypes reveal novel loci associated with Alzheimer's disease and three causality networks: The GR@ACE project. Alzheimer's and Dementia, 2019, 15, 1333-1347.	0.8	111
24	A genomeâ€wide association study on low susceptibility to hepatitis C virus infection (GEHEP012 study). Liver International, 2019, 39, 1918-1926.	3.9	4
25	MicroRNA Profile of HCV Spontaneous Clarified Individuals, Denotes Previous HCV Infection. Journal of Clinical Medicine, 2019, 8, 849.	2.4	11
26	Genetic markers of lipid metabolism genes associated with low susceptibility to HCV infection. Scientific Reports, 2019, 9, 9054.	3.3	2
27	Galectin-3, a novel endogenous TREM2 ligand, detrimentally regulates inflammatory response in Alzheimer's disease. Acta Neuropathologica, 2019, 138, 251-273.	7.7	187
28	Genetic meta-analysis of diagnosed Alzheimer's disease identifies new risk loci and implicates Aβ, tau, immunity and lipid processing. Nature Genetics, 2019, 51, 414-430.	21.4	1,962
29	HIV infection does not increase the risk of liver complications in hepatitis C virus-infected patient with advanced fibrosis, after sustained virological response with direct-acting antivirals. Aids, 2019, 33, 1167-1174.	2.2	15
30	Genetic Association Studies in Host–Pathogen Interaction Analysis. Methods in Molecular Biology, 2018, 1734, 1-11.	0.9	0
31	HERC1 Ubiquitin Ligase Is Required for Normal Axonal Myelination in the Peripheral Nervous System. Molecular Neurobiology, 2018, 55, 8856-8868.	4.0	14
32	Baseline resistance-guided therapy does not enhance the response to interferon-free treatment of HCV infection in real life. Scientific Reports, 2018, 8, 14905.	3.3	4
33	Sustained virological response to direct-acting antiviral regimens reduces the risk of hepatocellular carcinoma in HIV/HCV-coinfected patients with cirrhosis. Journal of Antimicrobial Chemotherapy, 2018, 73, 2435-2443.	3.0	16
34	Fat mass and obesityâ€associated gene variations are related to fatty liver disease in <scp>HIV</scp> â€infected patients. HIV Medicine, 2017, 18, 546-554.	2.2	6
35	Low incidence of acute hepatitis C virus infection among Southern Spanish HIV-infected individuals. Journal of Infection, 2017, 74, 514-517.	3.3	1
36	High frequency of potential interactions between directâ€acting antivirals and concomitant therapy in <scp>HIV</scp> /hepatitis C virusâ€coinfected patients in clinical practice. HIV Medicine, 2017, 18, 445-451.	2.2	11

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37	Changes in Liver Steatosis After Switching From Efavirenz to Raltegravir Among Human Immunodeficiency Virus–Infected Patients With Nonalcoholic Fatty Liver Disease. Clinical Infectious Diseases, 2017, 65, 1012-1019.	5.8	42
38	Non-Alcoholic Fatty Liver Disease in HIV Infection. AIDS Reviews, 2017, 19, 35-46.	1.0	12
39	The PNPLA3 Genetic Variant rs738409 Influences the Progression to Cirrhosis in HIV/Hepatitis C Virus Coinfected Patients. PLoS ONE, 2016, 11, e0168265.	2.5	15
40	Changes in liver steatosis evaluated by transient elastography with the controlled attenuation parameter in <scp>HIV</scp> â€infected patients. HIV Medicine, 2016, 17, 766-773.	2.2	23
41	Common haplotypes in CD209 promoter and susceptibility to HIV-1 infection in intravenous drug users. Infection, Genetics and Evolution, 2016, 45, 20-25.	2.3	7
42	Changes in the response to treatment against chronic hepatitis C between 1999 and 2015. European Journal of Gastroenterology and Hepatology, 2016, 28, 1253-1257.	1.6	1
43	Impact of genetic polymorphisms associated with nonalcoholic fatty liver disease on HIV-infected individuals. Aids, 2015, 29, 1927-1935.	2.2	19
44	IFNL4 rs368234815 polymorphism is associated with innate resistance to HIV-1 infection. Aids, 2015, 29, 1895-1897.	2.2	19
45	Association of complement receptor 2 polymorphisms with innate resistance to HIV-1 infection. Genes and Immunity, 2015, 16, 134-141.	4.1	13
46	Hepatitis C virus reinfection after sustained virological response in HIV-infected patients with chronic hepatitis C. Journal of Infection, 2015, 71, 571-577.	3.3	42
47	IFNL4 ss469415590 Variant Shows Similar Performance to rs12979860 as Predictor of Response to Treatment against Hepatitis C Virus Genotype 1 or 4 in Caucasians. PLoS ONE, 2014, 9, e95515.	2.5	24
48	A Colorectal Cancer Susceptibility New Variant at 4q26 in the Spanish Population Identified by Genome-Wide Association Analysis. PLoS ONE, 2014, 9, e101178.	2.5	26
49	A Regulatory Polymorphism in HAVCR2 Modulates Susceptibility to HIV-1 Infection. PLoS ONE, 2014, 9, e106442.	2.5	13
50	Core amino acid variation at position 110 is associated with sustained virological response in Caucasian patients with chronic hepatitis C virus 1b infection. Archives of Virology, 2014, 159, 3345-3351.	2.1	3
51	ATP5H/KCTD2 locus is associated with Alzheimer's disease risk. Molecular Psychiatry, 2014, 19, 682-687.	7.9	62
52	A polymorphism linked to <i><scp>RRAS</scp></i> , <i><scp>SCAF</scp>1</i> , <i><scp>IRF</scp>3</i> and <i><scp>BCL</scp>2L12</i> genes is associated with cirrhosis in hepatitis C virus carriers. Liver International, 2014, 34, 558-566.	3.9	9
53	Association of low-density lipoprotein receptor genotypes with hepatitis C viral load. Genes and Immunity, 2014, 15, 16-24.	4.1	16
54	Incidence and natural history of hepatitis E virus coinfection among HIV-infected patients. Aids, 2014, 28, 1931-1937.	2.2	52

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55	Reassessment of Genotype 1 Hepatitis C Virus Subtype Misclassification by LiPA 2.0: Implications for Direct-Acting Antiviral Treatment. Journal of Clinical Microbiology, 2014, 52, 4027-4029.	3.9	23
56	Exploratory analysis of seven Alzheimer's disease genes: disease progression. Neurobiology of Aging, 2013, 34, 1310.e1-1310.e7.	3.1	15
57	Genetic analysis of candidate SNPs for metabolic syndrome in obstructive sleep apnea (OSA). Gene, 2013, 521, 150-154.	2.2	10
58	Variations at multiple genes improve interleukin 28B genotype predictive capacity for response to therapy against hepatitis C infection. Aids, 2013, 27, 2715-2724.	2.2	7
59	Genetic Study of Neurexin and Neuroligin Genes in Alzheimer's Disease. Journal of Alzheimer's Disease, 2013, 35, 403-412.	2.6	38
60	Molecular evaluation of human Ubiquilin 2 gene PXX domain in familial frontotemporal dementia patients. Journal of Neurology, 2012, 259, 2488-2490.	3.6	2
61	Estrogen receptor alpha gene variants are associated with Alzheimer's disease. Neurobiology of Aging, 2012, 33, 198.e15-198.e24.	3.1	36
62	Genome-Wide Association Study of Multiple Sclerosis Confirms a Novel Locus at 5p13.1. PLoS ONE, 2012, 7, e36140.	2.5	46
63	The membrane-spanning 4-domains, subfamily A (MS4A) gene cluster contains a common variant associated with Alzheimer's disease. Genome Medicine, 2011, 3, 33.	8.2	81
64	Genetic association of complement receptor 1 polymorphism rs3818361 in Alzheimer's disease. Alzheimer's and Dementia, 2011, 7, e124-9.	0.8	16
65	The MTHFD1L Gene rs11754661 Marker is Not Associated with Alzheimer's Disease in a Sample of the Spanish Population. Journal of Alzheimer's Disease, 2011, 25, 47-50.	2.6	5
66	Identification of genetic factors associated with susceptibility to angiotensin-converting enzyme inhibitors-induced cough. Pharmacogenetics and Genomics, 2011, 21, 10-17.	1.5	45
67	<i>Calpain 10</i> gene and laryngeal cancer: A survival analysis. Head and Neck, 2011, 33, 72-76.	2.0	13
68	Activation of PKR Causes Amyloid ß-Peptide Accumulation via De-Repression of BACE1 Expression. PLoS ONE, 2011, 6, e21456.	2.5	50
69	CALHM1 P86L Polymorphism is Associated with Late-Onset Alzheimer's Disease in a Recessive Model. Journal of Alzheimer's Disease, 2010, 20, 247-251.	2.6	38
70	Genetic Structure of the Spanish Population. BMC Genomics, 2010, 11, 326.	2.8	49
71	WWOX gene is associated with HDL cholesterol and triglyceride levels. BMC Medical Genetics, 2010, 11, 148.	2.1	24
72	Whole-genome conditional two-locus analysis identifies novel candidate genes for late-onset Parkinson's disease. Neurogenetics, 2009, 10, 173-181.	1.4	13

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73	GOLPH2 Gene Markers are Not Associated with Alzheimer's Disease in a Sample of the Spanish Population. Journal of Alzheimer's Disease, 2009, 18, 751-754.	2.6	7
74	Analysis of HLA class I expression in progressing and regressing metastatic melanoma lesions after immunotherapy. Immunogenetics, 2008, 60, 439-447.	2.4	119
75	Regressing and progressing metastatic lesions: resistance to immunotherapy is predetermined by irreversible HLA class I antigen alterations. Cancer Immunology, Immunotherapy, 2008, 57, 1727-1733.	4.2	56
76	A method for detecting epistasis in genome-wide studies using case-control multi-locus association analysis. BMC Genomics, 2008, 9, 360.	2.8	76
77	CAPN10 alleles modify laryngeal cancer risk in the Spanish population. European Journal of Surgical Oncology, 2008, 34, 94-99.	1.0	10
78	A Digenic Combination of Polymorphisms Within ESR1 and ESR2 Genes Are Associated With Age at Menarche in the Spanish Population. Reproductive Sciences, 2008, 15, 305-311.	2.5	15
79	Identification of a 2244 base pair interstitial deletion within the human ESR1 gene in the Spanish population. Journal of Medical Genetics, 2008, 45, 420-424.	3.2	7
80	Analysis of the ERalpha germline Pvull marker in breast cancer risk. Medical Science Monitor, 2008, 14, CR136-43.	1.1	13
81	Genetic analysis of the GRM1 gene in human melanoma susceptibility. European Journal of Human Genetics, 2007, 15, 1176-1182.	2.8	17
82	Identification of a protective haplogenotype within CAPN10 gene influencing colorectal cancer susceptibility. Journal of Gastroenterology and Hepatology (Australia), 2007, 22, 2298-2302.	2.8	14
83	HLA class I expression in metastatic melanoma correlates with tumor development during autologous vaccination. Cancer Immunology, Immunotherapy, 2007, 56, 709-717.	4.2	78
84	Pyrosequencing protocol requiring a unique biotinylated primer. Clinical Chemistry and Laboratory Medicine, 2006, 44, 435-41.	2.3	5
85	Bone morphogenetic protein 15 (BMP15) alleles predict over-response to recombinant follicle stimulation hormone and iatrogenic ovarian hyperstimulation syndrome (OHSS). Pharmacogenetics and Genomics, 2006, 16, 485-495.	1.5	58
86	Lack of Association Between NOS3 Glu298Asp and Breast Cancer Risk: a Case–ontrol Study. Breast Cancer Research and Treatment, 2006, 100, 331-333.	2.5	21
87	Specific haplotypes of the CALPAIN-5 gene are associated with polycystic ovary syndrome. Human Reproduction, 2006, 21, 943-951.	0.9	20
88	Association of genetic markers within the KIT and KITLG genes with human male infertility. Human Reproduction, 2006, 21, 3185-3192.	0.9	40
89	Exploring allelic imbalance within paraffin-embedded tumor biopsies using pyrosequencing technology. Clinical Chemistry and Laboratory Medicine, 2006, 44, 1076-81.	2.3	7
90	Genetic analysis of CAV1 gene in hypertension and metabolic syndrome. Thrombosis and Haemostasis, 2006, 95, 696-701.	3.4	9

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91	Genetic analysis of caveolin-1 and eNOS genes in colorectal cancer. Oncology Reports, 2006, 16, 353-9.	2.6	18
92	Preliminary molecular genetic analysis of the Receptor Interacting Protein 140 (RIP140) in women affected by endometriosis. Journal of Experimental & Clinical Assisted Reproduction, 2005, 2, 11.	0.4	15
93	Re: Polymorphisms Associated With Circulating Sex Hormone Levels in Postmenopausal Women. Journal of the National Cancer Institute, 2005, 97, 152-153.	6.3	13
94	Multilocus analyses of estrogen-related genes reveal involvement of the ESR1 gene in male infertility and the polygenic nature of the pathology. Fertility and Sterility, 2005, 84, 910-918.	1.0	47
95	Pharmacogenetics of controlled ovarian hyperstimulation. Pharmacogenomics, 2005, 6, 629-637.	1.3	25
96	Association of NOS3 gene with metabolic syndrome in hypertensive patients. Thrombosis and Haemostasis, 2004, 92, 413-418.	3.4	50
97	Absence of de novo Y-chromosome microdeletions in male children conceived through intracytoplasmic sperm injection. Fertility and Sterility, 2004, 82, 1679-1680.	1.0	8
98	Human controlled ovarian hyperstimulation outcome is a polygenic trait. Pharmacogenetics and Genomics, 2004, 14, 285-293.	5.7	109
99	Role of follicle-stimulating hormone receptor Ser680Asn polymorphism in the efficacy of follicle-stimulating hormone. Fertility and Sterility, 2003, 80, 571-576.	1.0	122
100	Scanning of Y-chromosome azoospermia factors loci using real-time polymerase chain reaction and melting curve analysis. Fertility and Sterility, 2003, 80, 907-913.	1.0	19
101	CCR5 Expression Influences the Progression of Human Breast Cancer in a p53-dependent Manner. Journal of Experimental Medicine, 2003, 198, 1381-1389.	8.5	129
102	Detection of Pvull Polymorphism within Intron 1 of ESR1 Gene by Real-Time PCR. Clinical Chemistry and Laboratory Medicine, 2003, 41, 392-3.	2.3	10
103	Specific CAPN10 Gene Haplotypes Influence the Clinical Profile of Polycystic Ovary Patients. Journal of Clinical Endocrinology and Metabolism, 2003, 88, 5529-5536.	3.6	63
104	CAPN10 Alleles Are Associated with Polycystic Ovary Syndrome. Journal of Clinical Endocrinology and Metabolism, 2002, 87, 3971-3976.	3.6	76
105	CAPN10 Alleles Are Associated with Polycystic Ovary Syndrome. Journal of Clinical Endocrinology and Metabolism, 2002, 87, 3971-3976.	3.6	30
106	Detection of Nucleotide c985 A→G Mutation of Medium-Chain Acyl-CoA Dehydrogenase Gene by Real-Time PCR. Clinical Chemistry, 2001, 47, 958-959.	3.2	5
107	Multiple mechanisms of immune evasion can coexist in melanoma tumor cell lines derived from the same patient. Cancer Immunology, Immunotherapy, 2001, 49, 621-628.	4.2	45
108	Microsatellite instability analysis in tumors with different mechanisms for total loss of HLA expression. Cancer Immunology, Immunotherapy, 2000, 48, 684-690.	4.2	21

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109	Looking for HLA-G expression in human tumours. Journal of Reproductive Immunology, 1999, 43, 263-273.	1.9	13
110	Differential effect on U937 cell differentiation by targeting transcriptional factors implicated in tissue- or stage-specific induced integrin expression. Experimental Hematology, 1999, 27, 353-364.	0.4	37
111	Expression of HLA G in human tumors is not a frequent event. , 1999, 81, 512-518.		65
112	Chromosome loss is the most frequent mechanism contributing to HLA haplotype loss in human tumors. , 1999, 83, 91-97.		104
113	Chromosome loss is the most frequent mechanism contributing to HLA haplotype loss in human tumors. International Journal of Cancer, 1999, 83, 91-97.	5.1	3
114	In vivo and in vitro generation of a new altered HLA phenotype in melanoma-tumour-cell variants expressing a single HLA-class-I allele. , 1998, 75, 317-323.		31
115	Unresponsiveness to interferon associated with STAT1 protein deficiency in a gastric adenocarcinoma cell line. Cancer Immunology, Immunotherapy, 1998, 47, 113-120.	4.2	62
116	In vivo and in vitro generation of a new altered HLA phenotype in melanomaâ€ŧumourâ€cell variants expressing a single HLAâ€classâ€I allele. International Journal of Cancer, 1998, 75, 317-323.	5.1	1