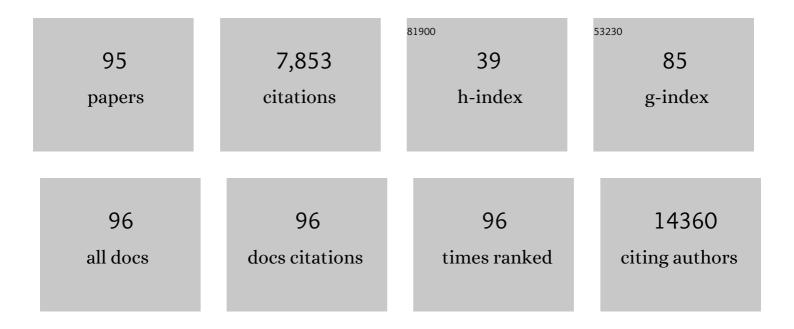


## List of Publications by Year in descending order

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#	Article	IF	CITATIONS
1	Interferon Control of Human Coronavirus Infection and Viral Evasion: Mechanistic Insights and Implications for Antiviral Drug and Vaccine Development. Journal of Molecular Biology, 2022, 434, 167438.	4.2	7
2	APOL1 variant alleles associate with reduced risk for opportunistic infections in HIV infection. Communications Biology, 2021, 4, 284.	4.4	4
3	The Epigenetic Modulation of Cancer and Immune Pathways in Hepatitis B Virus-Associated Hepatocellular Carcinoma: The Influence of HBx and miRNA Dysregulation. Frontiers in Immunology, 2021, 12, 661204.	4.8	28
4	Impacts of the SOAT1 genetic variants and protein expression on HBV-related hepatocellular carcinoma. BMC Cancer, 2021, 21, 615.	2.6	6
5	The Multiple Roles of Hepatitis B Virus X Protein (HBx) Dysregulated MicroRNA in Hepatitis B Virus-Associated Hepatocellular Carcinoma (HBV-HCC) and Immune Pathways. Viruses, 2020, 12, 746.	3.3	30
6	Dysregulated microRNAs in Hepatitis B Virus-Related Hepatocellular Carcinoma: Potential as Biomarkers and Therapeutic Targets. Frontiers in Oncology, 2020, 10, 1271.	2.8	41
7	Impact of APOL1 Genetic Variants on HIV-1 Infection and Disease Progression. Frontiers in Immunology, 2019, 10, 53.	4.8	13
8	Host Genetics of Cytomegalovirus Pathogenesis. Frontiers in Genetics, 2019, 10, 616.	2.3	47
9	The Regulatory Role of MicroRNA in Hepatitis-B Virus-Associated Hepatocellular Carcinoma (HBV-HCC) Pathogenesis. Cells, 2019, 8, 1504.	4.1	63
10	Editorial: Host Genetics in Viral Pathogenesis and Control. Frontiers in Genetics, 2019, 10, 1038.	2.3	3
11	Genetically Regulated Bilirubin and Risk of Non-alcoholic Fatty Liver Disease: A Mendelian Randomization Study. Frontiers in Genetics, 2018, 9, 662.	2.3	16
12	Host and Viral Genetic Variation in HBV-Related Hepatocellular Carcinoma. Frontiers in Genetics, 2018, 9, 261.	2.3	112
13	The Loss-of-Function S267F Variant in HBV Receptor NTCP Reduces Human Risk for HBV Infection and Disease Progression. Journal of Infectious Diseases, 2018, 218, 1404-1410.	4.0	20
14	IFITM Genes, Variants, and Their Roles in the Control and Pathogenesis of Viral Infections. Frontiers in Microbiology, 2018, 9, 3228.	3.5	129
15	Elevated CO <sub>2</sub> concentrations affect the growth patterns of dominant C <sub>3</sub> and C <sub>4</sub> shrub species differently in the Mu Us Sandy Land of Inner Mongolia. Botany, 2017, 95, 869-877.	1.0	6
16	Application of silica gel in removing high concentrations toluene vapor by adsorption and desorption process. Journal of the Taiwan Institute of Chemical Engineers, 2017, 74, 218-224.	5.3	67
17	Exendin-4 Upregulates Adiponectin Level in Adipocytes via Sirt1/Foxo-1 Signaling Pathway. PLoS ONE, 2017, 12, e0169469.	2.5	35
18	Different implications of OSL and radiocarbon ages in archaeological sites in the Qaidam Basin, Qinghai-Tibetan Plateau. Geochronometria, 2016, 43, 188-200.	0.8	7

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19	Association of Polymorphisms in the Regulatory Region of the Cyclophilin a Gene (PPIA) with Gene Expression and HIV/AIDS Disease Progression. Journal of Acquired Immune Deficiency Syndromes (1999), 2016, 72, 465-473.	2.1	8
20	BMI, leisure-time physical activity, and physical fitness in adults in China: results from a series of national surveys, 2000–14. Lancet Diabetes and Endocrinology,the, 2016, 4, 487-497.	11.4	180
21	Role of APOBEC3F Gene Variation in HIV-1 Disease Progression and Pneumocystis Pneumonia. PLoS Genetics, 2016, 12, e1005921.	3.5	17
22	Telomerase reverse transcriptase promoter mutations in hepatitis B virus-associated hepatocellular carcinoma. Oncotarget, 2016, 7, 27838-27847.	1.8	25
23	Low-frequency and rare exome chip variants associate with fasting glucose and type 2 diabetes susceptibility. Nature Communications, 2015, 6, 5897.	12.8	173
24	Sequencing rare and common APOL1 coding variants to determine kidney disease risk. Kidney International, 2015, 88, 754-763.	5.2	30
25	Distribution of plant species and species-soil relationship in the east central Gurbantunggut Desert, China. Journal of Chinese Geography, 2015, 25, 101-112.	3.9	12
26	Regulatory Variation in HIV-1 Dependency Factor <i>ZNRD1</i> Associates with Host Resistance to HIV-1 Acquisition. Journal of Infectious Diseases, 2014, 210, 1539-1548.	4.0	11
27	Comparative multidimensional molecular analyses of pediatric diffuse intrinsic pontine glioma reveals distinct molecular subtypes. Acta Neuropathologica, 2014, 127, 881-895.	7.7	91
28	Genome-wide association studies identified novel loci for non-high-density lipoprotein cholesterol and its postprandial lipemic response. Human Genetics, 2014, 133, 919-930.	3.8	10
29	ZnO nanoparticles embedded in polyethylene-glycol (PEG) matrix as sensitive strain gauge elements. Journal of Nanoparticle Research, 2014, 16, 1.	1.9	5
30	Identification of a Novel Polyomavirus in a Pancreatic Transplant Recipient With Retinal Blindness and Vasculitic Myopathy. Journal of Infectious Diseases, 2014, 210, 1595-1599.	4.0	129
31	Genome-wide association study identifies common loci influencing circulating glycated hemoglobin (HbA1c) levels in non-diabetic subjects: The Long Life Family Study (LLFS). Metabolism: Clinical and Experimental, 2014, 63, 461-468.	3.4	22
32	Persistence of four dominant psammophyte species in central Inner Mongolia of China under continual drought. Journal of Arid Land, 2013, 5, 331-339.	2.3	4
33	Isoflurane post-conditioning protects primary cultures of cortical neurons against oxygen and glucose deprivation injury via upregulation of Slit2/Robo1. Brain Research, 2013, 1537, 283-289.	2.2	15
34	Identifying Darwinian Selection Acting on Different Human APOL1 Variants among Diverse African Populations. American Journal of Human Genetics, 2013, 93, 191.	6.2	1
35	Propofol Increases Expression of Basic Fibroblast Growth Factor After Transient Cerebral Ischemia in Rats. Neurochemical Research, 2013, 38, 530-537.	3.3	16
36	Identifying Darwinian Selection Acting on Different Human APOL1 Variants among Diverse African Populations. American Journal of Human Genetics, 2013, 93, 54-66.	6.2	91

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37	Focal segmental glomerulosclerosis is associated with a <i>PDSS2</i> haplotype and, independently, with a decreased content of coenzyme Q <sub>10</sub> . American Journal of Physiology - Renal Physiology, 2013, 305, F1228-F1238.	2.7	36
38	Multicohort Genomewide Association Study Reveals a New Signal of Protection Against HIV-1 Acquisition. Journal of Infectious Diseases, 2012, 205, 1155-1162.	4.0	22
39	The Effect of CYP7A1 Polymorphisms on Lipid Responses to Fenofibrate. Journal of Cardiovascular Pharmacology, 2012, 59, 254-259.	1.9	23
40	Alternative Splicing of a Novel Inducible Exon Diversifies the CASK Guanylate Kinase Domain. Journal of Nucleic Acids, 2012, 2012, 1-15.	1.2	12
41	A genome-wide approach accounting for body mass index identifies genetic variants influencing fasting glycemic traits and insulin resistance. Nature Genetics, 2012, 44, 659-669.	21.4	762
42	Recombinant Human Angiopoietin-1 Ameliorates the Expressions of ZO-1, Occludin, VE-cadherin, and PKCα Signaling after Focal Cerebral Ischemia/Reperfusion in Rats. Journal of Molecular Neuroscience, 2012, 46, 236-247.	2.3	53
43	Association of gene variants with lipid levels in response to fenofibrate is influenced by metabolic syndrome status. Atherosclerosis, 2011, 215, 435-439.	0.8	19
44	Screening Low-Frequency SNPS From Genome-Wide Association Study Reveals a New Risk Allele for Progression to AIDS. Journal of Acquired Immune Deficiency Syndromes (1999), 2011, 56, 279-284.	2.1	32
45	Meta-analysis of gene-environment interaction: joint estimation of SNP and SNP × environment regression coefficients. Genetic Epidemiology, 2011, 35, 11-18.	1.3	158
46	APOL1 Genetic Variants in Focal Segmental Glomerulosclerosis and HIV-Associated Nephropathy. Journal of the American Society of Nephrology: JASN, 2011, 22, 2129-2137.	6.1	713
47	Genome-Wide Association Study Implicates PARD3B-Based AIDS Restriction. Journal of Infectious Diseases, 2011, 203, 1491-1502.	4.0	49
48	Modelling seasonal evapotranspiration of arid lands in China. Hydrology Research, 2011, 42, 40-49.	2.7	2
49	Association of polymorphisms in the LEDGF/p75 gene (PSIP1) with susceptibility to HIV-1 infection and disease progression. Aids, 2011, 25, 1711-1719.	2.2	22
50	Role of Exonic Variation in Chemokine Receptor Genes on AIDS: CCRL2 F167Y Association with Pneumocystis Pneumonia. PLoS Genetics, 2011, 7, e1002328.	3.5	19
51	A Common HLA–DPA1 Variant is a Major Determinant of Hepatitis B Virus Clearance in Han Chinese. Journal of Infectious Diseases, 2011, 203, 943-947.	4.0	76
52	Host genes associated with HIV/AIDS: advances in gene discovery. Trends in Genetics, 2010, 26, 119-131.	6.7	99
53	Elevated CO2 accelerates net assimilation rate and enhance growth of dominant shrub species in a sand dune in central Inner Mongolia. Environmental and Experimental Botany, 2010, 68, 31-36.	4.2	11
54	Worldwide Distribution of the MYH9 Kidney Disease Susceptibility Alleles and Haplotypes: Evidence of Historical Selection in Africa. PLoS ONE, 2010, 5, e11474.	2.5	33

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55	Multipleâ€Cohort Genetic Association Study Reveals CXCR6 as a New Chemokine Receptor Involved in Longâ€Term Nonprogression to AIDS. Journal of Infectious Diseases, 2010, 202, 908-915.	4.0	82
56	Multistage Genomewide Association Study Identifies a Locus at 1q41 Associated with Rate of HIVâ€1 Disease Progression to Clinical AIDS. Journal of Infectious Diseases, 2010, 201, 618-626.	4.0	67
57	Reply to Itaya et al. Journal of Infectious Diseases, 2010, 202, 816-818.	4.0	9
58	Dense mapping of MYH9 localizes the strongest kidney disease associations to the region of introns 13 to 15. Human Molecular Genetics, 2010, 19, 1805-1815.	2.9	58
59	Putative Primo-vascular System in Mesentery of Rats. JAMS Journal of Acupuncture and Meridian Studies, 2010, 3, 232-240.	0.7	9
60	New genetic loci implicated in fasting glucose homeostasis and their impact on type 2 diabetes risk. Nature Genetics, 2010, 42, 105-116.	21.4	1,982
61	Genetic Variants at the PDZ-Interacting Domain of the Scavenger Receptor Class B Type I Interact with Diet to Influence the Risk of Metabolic Syndrome in Obese Men and Women. Journal of Nutrition, 2009, 139, 842-848.	2.9	19
62	Developmental and Genetic Regulation of Human Surfactant Protein B in vivo. Neonatology, 2009, 95, 117-124.	2.0	24
63	Epistatic interactions of CDKN2B-TCF7L2 for risk of type 2 diabetes and of CDKN2B-JAZF1 for triglyceride/high-density lipoprotein ratio longitudinal change: evidence from the Framingham Heart Study. BMC Proceedings, 2009, 3, S71.	1.6	13
64	Novel variants at KCTD10, MVK, and MMAB genes interact with dietary carbohydrates to modulate HDL-cholesterol concentrations in the Genetics of Lipid Lowering Drugs and Diet Network Study. American Journal of Clinical Nutrition, 2009, 90, 686-694.	4.7	25
65	<i>APOBEC3B</i> Deletion and Risk of HIVâ€I Acquisition. Journal of Infectious Diseases, 2009, 200, 1054-1058.	4.0	53
66	Polyunsaturated Fatty Acids Modulate the Effect of TCF7L2 Gene Variants on Postprandial Lipemia. Journal of Nutrition, 2009, 139, 439-446.	2.9	45
67	<i>ADIPOQ</i> Polymorphisms, Monounsaturated Fatty Acids, and Obesity Risk: The GOLDN Study. Obesity, 2009, 17, 510-517.	3.0	80
68	Expression of Duffy Antigen Receptor for Chemokines (DARC) Has No Effect on HIV-1 Acquisition or Progression to AIDS in African Americans. Cell Host and Microbe, 2009, 5, 411-413.	11.0	32
69	TERC and TERT gene mutations in patients with bone marrow failure and the significance of telomere length measurements. Blood, 2009, 113, 309-316.	1.4	129
70	Responses to air temperature and soil moisture of growth of four dominant species on sand dunes of central Inner Mongolia. Journal of Plant Research, 2008, 121, 473-482.	2.4	15
71	A population-based study to investigate host genetic factors associated with hepatitis B infection and pathogenesis in the Chinese population. BMC Infectious Diseases, 2008, 8, 1.	2.9	113
72	Enuresis Is a Common and Persistent Problem Among Children and Young Adults with Sickle Cell Anemia. Urology, 2008, 72, 81-84.	1.0	26

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73	Population and Disease-Based Prevalence of the Common Mutations Associated With Surfactant Deficiency. Pediatric Research, 2008, 63, 645-649.	2.3	94
74	Regulatory Polymorphisms in the Interleukinâ€18 Promoter Are Associated with Hepatitis C Virus Clearance. Journal of Infectious Diseases, 2008, 198, 1159-1165.	4.0	50
75	Polymorphisms of CUL5 Are Associated with CD4+ T Cell Loss in HIV-1 Infected Individuals. PLoS Genetics, 2007, 3, e19.	3.5	47
76	Regulatory Polymorphisms in the Cyclophilin A Gene, PPIA, Accelerate Progression to AIDS. PLoS Pathogens, 2007, 3, e88.	4.7	58
77	Plant species as indicators of the extent of desertification in four sandy rangelands. African Journal of Ecology, 2007, 45, 94-102.	0.9	11
78	Telomerase reverse transcriptase haploinsufficiency and telomere length in individuals with 5p– syndrome. Aging Cell, 2007, 6, 689-697.	6.7	22
79	Genetic Variation in the CCL18-CCL3-CCL4 Chemokine Gene Cluster Influences HIV Type 1 Transmission and AIDS Disease Progression. American Journal of Human Genetics, 2006, 79, 120-128.	6.2	63
80	Low Frequency of Telomerase RNA Mutations Among Children With Aplastic Anemia or Myelodysplastic Syndrome. Journal of Pediatric Hematology/Oncology, 2006, 28, 450-453.	0.6	27
81	Effects of human TRIM5α polymorphisms on antiretroviral function and susceptibility to human immunodeficiency virus infection. Virology, 2006, 354, 15-27.	2.4	116
82	Disruption of estrogen receptor DNA-binding domain and related intramolecular communication restores tamoxifen sensitivity in resistant breast cancer. Cancer Cell, 2006, 10, 487-499.	16.8	68
83	Global transmission/disequilibrium tests based on haplotype sharing in multiple candidate genes. Genetic Epidemiology, 2005, 29, 323-335.	1.3	7
84	Genetic variant characterization in intron 4 of the surfactant protein B gene. Human Mutation, 2005, 26, 494-495.	2.5	24
85	A Combinatorial Code for Splicing Silencing: UAGG and GGGG Motifs. PLoS Biology, 2005, 3, e158.	5.6	177
86	The Effect of RANTES Chemokine Genetic Variants on Early HIV-1 Plasma RNA Among African American Injection Drug Users. Journal of Acquired Immune Deficiency Syndromes (1999), 2005, 38, 584-589.	2.1	24
87	Patterns of ethnic diversity among the genes that influence AIDS. Human Molecular Genetics, 2004, 13, 9R-19.	2.9	46
88	APOBEC3G Genetic Variants and Their Influence on the Progression to AIDS. Journal of Virology, 2004, 78, 11070-11076.	3.4	178
89	Dominant Effects of CCR2-CCR5 Haplotypes in HIV-1 Disease Progression. Journal of Acquired Immune Deficiency Syndromes (1999), 2004, 37, 1534-1538.	2.1	32
90	Genetic influence of CXCR6 chemokine receptor alleles on PCP-mediated AIDS progression among African Americans. Genes and Immunity, 2003, 4, 245-250.	4.1	48

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91	A Tumor Necrosis Factor–α–Inducible Promoter Variant of Interferonâ€Î³ Accelerates CD4+T Cell Depletion in Human Immunodeficiency Virus–1–Infected Individuals. Journal of Infectious Diseases, 2003, 188, 228-231.	4.0	38
92	Modulating influence on HIV/AIDS by interacting <i>RANTES</i> gene variants. Proceedings of the National Academy of Sciences of the United States of America, 2002, 99, 10002-10007.	7.1	196
93	Validation of Genetic Case-Control Studies in AIDS and Application to the CX3CR1 Polymorphism. Journal of Acquired Immune Deficiency Syndromes (1999), 2001, 26, 507-511.	2.1	23
94	Validation of Genetic Case-Control Studies in AIDS and Application to the CX3CR1 Polymorphism. Journal of Acquired Immune Deficiency Syndromes (1999), 2001, 26, 507-511.	2.1	16
95	Influence of CCR5 promoter haplotypes on AIDS progression in African–Americans. Aids, 2000, 14, 2117-2122.	2.2	70