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List of Publications by Year
in descending order

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65 papers	1,264 citations	471509 17 h-index	414414 32 g-index
72 all docs	72 docs citations	72 times ranked	2141 citing authors

#	Article	IF	Citations
1	Artemisinin kills malaria parasites by damaging proteins and inhibiting the proteasome. Nature Communications, 2018, 9, 3801.	12.8	193
2	Identification of copy number variations and common deletion polymorphisms in cattle. BMC Genomics, 2010, 11, 232.	2.8	126
3	Aminoacyl-tRNA synthetases as drug targets in eukaryotic parasites. International Journal for Parasitology: Drugs and Drug Resistance, 2014, 4, 1-13.	3.4	116
4	Genome-wide association study of aspirin-exacerbated respiratory disease in a Korean population. Human Genetics, 2013, 132, 313-321.	3.8	69
5	Inhibition of Resistance-Refractory P. falciparum Kinase PKG Delivers Prophylactic, Blood Stage, and Transmission-Blocking Antiplasmodial Activity. Cell Chemical Biology, 2020, 27, 806-816.e8.	5.2	56
6	Chemogenomics identifies acetyl-coenzyme A synthetase as a target for malaria treatment and prevention. Cell Chemical Biology, 2022, 29, 191-201.e8.	5.2	39
7	Association study of genetic variations in microRNAs with the risk of hepatitis B-related liver diseases. Digestive and Liver Disease, 2012, 44, 849-854.	0.9	37
8	Prioritization of Molecular Targets for Antimalarial Drug Discovery. ACS Infectious Diseases, 2021, 7, 2764-2776.	3.8	35
9	Selective inhibition of apicoplast tryptophanyl-tRNA synthetase causes delayed death in Plasmodium falciparum. Scientific Reports, 2016, 6, 27531.	3.3	34
10	Targeting Protein Translation in Organelles of the Apicomplexa. Trends in Parasitology, 2016, 32, 953-965.	3.3	31
11	Association of SLC6A12 variants with aspirinâ€intolerant asthma in a Korean population. Annals of Human Genetics, 2010, 74, 326-334.	0.8	29
12	An integrated platform for genome engineering and gene expression perturbation in Plasmodium falciparum. Scientific Reports, 2021, 11, 342.	3.3	29
13	The antimalarial MMV688533 provides potential for single-dose cures with a high barrier to Plasmodium falciparum parasite resistance. Science Translational Medicine, 2021, 13, .	12.4	25
14	Reaction hijacking of tyrosine tRNA synthetase as a new whole-of-life-cycle antimalarial strategy. Science, 2022, 376, 1074-1079.	12.6	25
15	Association of CACNG6 polymorphisms with aspirin-intolerance asthmatics in a Korean population. BMC Medical Genetics, 2010, 11, 138.	2.1	23
16	A possible association of EMID2 polymorphisms with aspirin hypersensitivity in asthma. Immunogenetics, 2011, 63, 13-21.	2.4	21
17	Phosphatidylinositol 3-phosphate and Hsp70 protect Plasmodium falciparum from heat-induced cell death. ELife, 2020, 9, .	6.0	20
18	The Genetic Effect of Copy Number Variations on the Risk of Type 2 Diabetes in a Korean Population. PLoS ONE, 2011, 6, e19091.	2.5	19

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19	Association analysis of UBE3C polymorphisms in Korean aspirin-intolerant asthmatic patients. Annals of Allergy, Asthma and Immunology, 2010, 105, 307-312.e1.	1.0	16
20	Positive Association between Aspirin-Intolerant Asthma and Genetic Polymorphisms of FSIP1: a Case-Case Study. BMC Pulmonary Medicine, 2010, 10, 34.	2.0	14
21	HLA-DRA Polymorphisms associated with Risk of Nasal Polyposis in Asthmatic Patients. American Journal of Rhinology and Allergy, 2012, 26, 12-17.	2.0	14
22	Possible role of EMID2 on nasal polyps pathogenesis in Korean asthma patients. BMC Medical Genetics, 2012, 13, 2.	2.1	14
23	The Plasmodium falciparum ABC transporter ABCI3 confers parasite strain-dependent pleiotropic antimalarial drug resistance. Cell Chemical Biology, 2022, 29, 824-839.e6.	5.2	14
24	UBE3C genetic variations as potent markers of nasal polyps in Korean asthma patients. Journal of Human Genetics, 2011, 56, 797-800.	2.3	13
25	DCBLD2 Gene Variations Correlate with Nasal Polyposis in Korean Asthma Patients. Lung, 2012, 190, 199-207.	3.3	13
26	Preclinical characterization and target validation of the antimalarial pantothenamide MMV693183. Nature Communications, 2022, 13, 2158.	12.8	13
27	WDR46is a Genetic Risk Factor for Aspirin-Exacerbated Respiratory Disease in a Korean Population. Allergy, Asthma and Immunology Research, 2012, 4, 199.	2.9	12
28	Genetic association analysis of ERBB4 polymorphisms with the risk of schizophrenia and SPEM abnormality in a Korean population. Brain Research, 2012, 1466, 146-151.	2.2	12
29	Functional genomics of RAP proteins and their role in mitoribosome regulation in Plasmodium falciparum. Nature Communications, 2022, 13, 1275.	12.8	12
30	<i>TGFBR3</i> Polymorphisms and Its Haplotypes Associated with Chronic Hepatitis B Virus Infection and Age of Hepatocellular Carcinoma Occurrence. Digestive Diseases, 2011, 29, 278-283.	1.9	11
31	Lack of association of RAD51 genetic variations with hepatitis B virus clearance and occurrence of hepatocellular carcinoma in a Korean population. Journal of Medical Virology, 2011, 83, 1892-1899.	5.0	11
32	Genetic association analysis of CIITA variations with nasal polyp pathogenesis in asthmatic patients. Molecular Medicine Reports, 2013, 7, 927-934.	2.4	11
33	Neuregulin 3does not confer risk for schizophrenia and smooth pursuit eye movement abnormality in a Korean population. Genes, Brain and Behavior, 2011, 10, 828-833.	2.2	10
34	The Genetic Effect of Copy Number Variations on the Risk of Alcoholism in a Korean Population. Alcoholism: Clinical and Experimental Research, 2012, 36, 35-42.	2.4	10
35	Lack of Associations of Neuregulin 1 Variations with Schizophrenia and Smooth Pursuit Eye Movement Abnormality in a Korean Population. Journal of Molecular Neuroscience, 2012, 46, 476-482.	2.3	10
36	Genetic association analysis of TAP1 and TAP2 polymorphisms with aspirin exacerbated respiratory disease and its FEV1 decline. Journal of Human Genetics, 2011, 56, 652-659.	2.3	9

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37	Association analysis of C6 genetic variations and aspirin hypersensitivity in Korean asthmatic patients. Human Immunology, 2011, 72, 973-978.	2.4	8
38	Polymorphisms ofATF6BAre Potentially Associated With FEV1 Decline by Aspirin Provocation in Asthmatics. Allergy, Asthma and Immunology Research, 2014, 6, 142.	2.9	8
39	Genome-wide association analysis of copy number variations in subarachnoid aneurysmal hemorrhage. Journal of Human Genetics, 2010, 55, 726-730.	2.3	7
40	Association of the variants in AGT gene with modified drug response in Korean aspirin-intolerant asthma patients. Pulmonary Pharmacology and Therapeutics, 2011, 24, 595-601.	2.6	7
41	Association analysis of DTD1 gene variations with aspirin-intolerance in asthmatics. International Journal of Molecular Medicine, 2011, 28, 129-37.	4.0	7
42	Possible Association of SLC22A2 Polymorphisms with Aspirin-Intolerant Asthma. International Archives of Allergy and Immunology, 2011, 155, 395-402.	2.1	7
43	Contribution of the OBSCN Nonsynonymous Variants to Aspirin Exacerbated Respiratory Disease Susceptibility in Korean Population. DNA and Cell Biology, 2012, 31, 1001-1009.	1.9	7
44	Potential Association ofDCBLD2Polymorphisms with Fall Rates of FEV₁by Aspirin Provocation in Korean Asthmatics. Journal of Korean Medical Science, 2012, 27, 343.	2.5	7
45	Genome-Wide Profiling of Structural Genomic Variations in Korean HapMap Individuals. PLoS ONE, 2010, 5, e11417.	2.5	6
46	Lack of Association betweenCD58Genetic Variations and Aspirin-Exacerbated Respiratory Disease in a Korean Population. Journal of Asthma, 2011, 48, 539-545.	1.7	6
47	Association of FANCC polymorphisms with FEV1 decline in aspirin exacerbated respiratory disease. Molecular Biology Reports, 2012, 39, 2385-2394.	2.3	5
48	Selective expression of variant surface antigens enables Plasmodium falciparum to evade immune clearance in vivo. Nature Communications, 2022, 13, .	12.8	5
49	Potential Association Between ANXA4 Polymorphisms and Aspirin-exacerbated Respiratory Disease. Diagnostic Molecular Pathology, 2012, 21, 164-171.	2.1	4
50	Lack of association of the RTN4R genetic variations with risk of schizophrenia and SPEM abnormality in a Korean population. Psychiatry Research, 2011, 189, 312-314.	3.3	3
51	No associations of polymorphisms in ADPRT with hepatitis B virus clearance and hepatocellular carcinoma occurrence in a Korean population. Hepatology Research, 2011, 41, 250-257.	3.4	3
52	Genetic Analysis of Complement Component 9 (C9) Polymorphisms with Clearance of Hepatitis B Virus Infection. Digestive Diseases and Sciences, 2011, 56, 2735-2741.	2.3	3
53	Effect of Diffuse Panbronchiolitis Critical Region 1 Polymorphisms on the Risk of Aspirin-Exacerbated Respiratory Disease in Korean Asthmatics. Respiratory Care, 2012, 57, 758-763.	1.6	3
54	Association study between TRIM26 polymorphisms and risk of aspirin-exacerbated respiratory disease. International Journal of Molecular Medicine, 2012, 29, 927-33.	4.0	2

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55	CD55 polymorphisms and risk of aspirin-exacerbated respiratory disease. Molecular Medicine Reports, 2012, 6, 1087-1092.	2.4	2
56	Genetic variations in KIFC1 and the risk of aspirin exacerbated respiratory disease in a Korean population: an association analysis. Molecular Biology Reports, 2012, 39, 5913-5919.	2.3	2
57	A newly characterized malaria antigen on erythrocyte and merozoite surfaces induces parasite inhibitory antibodies. Journal of Experimental Medicine, 2021, 218, .	8.5	2
58	Lack of Association betweenPRNPM129V Polymorphism and Multiple Sclerosis, Mild Cognitive Impairment, Alcoholism and Schizophrenia in a Korean Population. Disease Markers, 2010, 28, 315-321.	1.3	2
59	Lack of association between proline dehydrogenase (oxidase) 1 polymorphisms and schizophrenia in a Korean population. Psychiatric Genetics, 2012, 22, 153-154.	1.1	1
60	Potential Association ofDDR1Genetic Variant with FEV1Decline by Aspirin Provocation in Asthmatics. Journal of Asthma, 2012, 49, 237-242.	1.7	1
61	Association Analysis BetweenFILIP1Polymorphisms and Aspirin Hypersensitivity in Korean Asthmatics. Allergy, Asthma and Immunology Research, 2013, 5, 34.	2.9	1
62	No association of TF gene polymorphisms with hepatitis B virus Clearance and hepatocellular carcinoma occurrence in a Korean population. Genes and Genomics, 2011, 33, 209-215.	1.4	0
63	Lack of association between FOS polymorphisms and clearance of HBV infection as well as HCC occurrence. Genes and Genomics, 2011, 33, 327-333.	1.4	0
64	Lack of association of HLA-DRA polymorphisms with aspirin exacerbated respiratory disease in a Korean population. Genes and Genomics, 2011, 33, 613-620.	1.4	0
65	Genetic analysis between FGD6 and aspirin exacerbated respiratory disease in a Korean population. Genes and Genomics, 2011, 33, 557-564.	1.4	0