List of Publications by Year in descending order

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CHIKASHI C TERAO

#	Article	IF	CITATIONS
1	Genome-wide association study of colorectal polyps identified highly overlapping polygenic architecture with colorectal cancer. Journal of Human Genetics, 2022, 67, 149-156.	1.1	5
2	Novel susceptibility loci for steroid-associated osteonecrosis of the femoral head in systemic lupus erythematosus. Human Molecular Genetics, 2022, 31, 1082-1095.	1.4	1
3	Usefulness of subclassification of adult diabetes mellitus among inpatients in Japan. Journal of Diabetes Investigation, 2022, 13, 706-713.	1.1	4
4	Association of Polygenic Risk Scores With Radiographic Progression in Patients With Rheumatoid Arthritis. Arthritis and Rheumatology, 2022, 74, 791-800.	2.9	12
5	A polygenic risk score improves risk stratification of coronary artery disease: a large-scale prospective Chinese cohort study. European Heart Journal, 2022, 43, 1702-1711.	1.0	58
6	Genomic analysis of familial pancreatic cancers and intraductal papillary mucinous neoplasms: A crossâ€sectional study. Cancer Science, 2022, 113, 1821-1829.	1.7	5
7	Co-occurrence of relapsing polychondritis and autoimmune thyroid diseases. Orphanet Journal of Rare Diseases, 2022, 17, 101.	1.2	2
8	Biological insights into systemic lupus erythematosus through an immune cell-specific transcriptome-wide association study. Annals of the Rheumatic Diseases, 2022, 81, 1273-1280.	0.5	9
9	Genetic Architectures Underlie Onset Age of Atopic Dermatitis. Journal of Investigative Dermatology, 2022, 142, 3337-3341.e7.	0.3	2
10	Long-term outcomes of refractory Takayasu arteritis patients treated with biologics including ustekinumab. Modern Rheumatology, 2021, 31, 678-683.	0.9	23
11	Chromosomally-integrated human herpesvirus 6 and autoimmune connective tissue diseases. Journal of Clinical Virology, 2021, 134, 104714.	1.6	0
12	Meta-analysis of 208370 East Asians identifies 113 susceptibility loci for systemic lupus erythematosus. Annals of the Rheumatic Diseases, 2021, 80, 632-640.	0.5	103
13	A case of retinitis pigmentosa homozygous for a rare CNGA1 causal variant. Scientific Reports, 2021, 11, 4681.	1.6	6
14	Comment on: Different treatment options for Takayasu arteritis patients with moderate-to-severe aortic regurgitation: long-term outcomes. Rheumatology, 2021, 60, e290-e291.	0.9	1
15	Comprehensive analysis of the major histocompatibility complex in systemic sclerosis identifies differential HLA associations by clinical and serological subtypes. Annals of the Rheumatic Diseases, 2021, 80, 1040-1047.	0.5	24
16	Lupus Susceptibility Region Containing <i>CDKN1B</i> rs34330 Mechanistically Influences Expression and Function of Multiple Target Genes, Also Linked to Proliferation and Apoptosis. Arthritis and Rheumatology, 2021, 73, 2303-2313.	2.9	11
17	Dynamic landscape of immune cell-specific gene regulation in immune-mediated diseases. Cell, 2021, 184, 3006-3021.e17.	13.5	147
18	Eight novel susceptibility loci and putative causal variants in atopic dermatitis. Journal of Allergy and Clinical Immunology, 2021, 148, 1293-1306.	1.5	32

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19	Hematopoietic mosaic chromosomal alterations increase the risk for diverse types of infection. Nature Medicine, 2021, 27, 1012-1024.	15.2	109
20	Combined landscape of single-nucleotide variants and copy number alterations in clonal hematopoiesis. Nature Medicine, 2021, 27, 1239-1249.	15.2	78
21	Genetics of autosomal mosaic chromosomal alteration (mCA). Journal of Human Genetics, 2021, 66, 879-885.	1.1	7
22	A susceptibility locus in the IL12B but not LILRA3 region is associated with vascular damage in Takayasu arteritis. Scientific Reports, 2021, 11, 13667.	1.6	11
23	Susceptibility loci and polygenic architecture highlight population specific and common genetic features in inguinal hernias. EBioMedicine, 2021, 70, 103532.	2.7	8
24	Genetic insights into biological mechanisms governing human ovarian ageing. Nature, 2021, 596, 393-397.	13.7	183
25	A cross-population atlas of genetic associations for 220 human phenotypes. Nature Genetics, 2021, 53, 1415-1424.	9.4	560
26	Factors associated with anxiety and depression in rheumatoid arthritis patients: a cross-sectional study. Advances in Rheumatology, 2021, 61, 65.	0.8	8
27	The power of genetic diversity in genome-wide association studies of lipids. Nature, 2021, 600, 675-679.	13.7	353
28	Genetic analysis of IgG4-related disease. Modern Rheumatology, 2020, 30, 17-23.	0.9	22
29	Population-specific and trans-ancestry genome-wide analyses identify distinct and shared genetic risk loci for coronary artery disease. Nature Genetics, 2020, 52, 1169-1177.	9.4	206
30	Genome-wide association study of intracranial aneurysms identifies 17 risk loci and genetic overlap with clinical risk factors. Nature Genetics, 2020, 52, 1303-1313.	9.4	163
31	Improving the trans-ancestry portability of polygenic risk scores by prioritizing variants in predicted cell-type-specific regulatory elements. Nature Genetics, 2020, 52, 1346-1354.	9.4	126
32	Functional annotation of human long noncoding RNAs via molecular phenotyping. Genome Research, 2020, 30, 1060-1072.	2.4	109
33	Endogenization and excision of human herpesvirus 6 in human genomes. PLoS Genetics, 2020, 16, e1008915.	1.5	22
34	Long-term follow-up of patients with anti-cyclic citrullinated peptide antibody-positive connective tissue disease: a retrospective observational study including information on the HLA-DRB1 allele and citrullination dependency. Arthritis Research and Therapy, 2020, 22, 248.	1.6	2
35	Response to: â€~Corresponence on â€~Shared epitope defines distinct associations of cigarette smoking with levels of anticitrullinated protein antibody and rheumatoid factor' by Ishikawa et al' by Regueiro and Gonzalez. Annals of the Rheumatic Diseases, 2020, , annrheumdis-2019-216872.	0.5	0
36	Genetics of systemic sclerosis. Journal of Scleroderma and Related Disorders, 2020, 5, 192-201.	1.0	4

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37	Using genetics to prioritize diagnoses for rheumatology outpatients with inflammatory arthritis. Science Translational Medicine, 2020, 12, .	5.8	31
38	Chromosomal alterations among age-related haematopoietic clones in Japan. Nature, 2020, 584, 130-135.	13.7	102
39	Roles of cytotoxic lymphocytes and MIC/LILR families in pathophysiology of Takayasu arteritis. Inflammation and Regeneration, 2020, 40, 9.	1.5	11
40	Large-scale genome-wide association study in a Japanese population identifies novel susceptibility loci across different diseases. Nature Genetics, 2020, 52, 669-679.	9.4	304
41	The Impact of Cigarette Smoking on Risk of Rheumatoid Arthritis: A Narrative Review. Cells, 2020, 9, 475.	1.8	63
42	Prevalence and Spectrum of Pathogenic Germline Variants in Japanese Patients With Early-Onset Colorectal, Breast, and Prostate Cancer. JCO Precision Oncology, 2020, 4, 183-191.	1.5	6
43	Polygenic Risk Score of Adolescent Idiopathic Scoliosis for Potential Clinical Use. Journal of Bone and Mineral Research, 2020, 36, 1481-1491.	3.1	5
44	Prevalence and factors associated with sarcopenia in patients with rheumatoid arthritis. Modern Rheumatology, 2019, 29, 589-595.	0.9	96
45	Shared epitope defines distinct associations of cigarette smoking with levels of anticitrullinated protein antibody and rheumatoid factor. Annals of the Rheumatic Diseases, 2019, 78, 1480-1487.	0.5	36
46	Genome-wide association study identifies 14 previously unreported susceptibility loci for adolescent idiopathic scoliosis in Japanese. Nature Communications, 2019, 10, 3685.	5.8	47
47	A frequent variant in the Japanese population determines quasi-Mendelian inheritance of rare retinal ciliopathy. Nature Communications, 2019, 10, 2884.	5.8	21
48	GWAS of mosaic loss of chromosome Y highlights genetic effects on blood cell differentiation. Nature Communications, 2019, 10, 4719.	5.8	50
49	Distinct HLA Associations with Rheumatoid Arthritis Subsets Defined by Serological Subphenotype. American Journal of Human Genetics, 2019, 105, 616-624.	2.6	27
50	PLD4 is a genetic determinant to systemic lupus erythematosus and involved in murine autoimmune phenotypes. Annals of the Rheumatic Diseases, 2019, 78, 509-518.	0.5	36
51	Mechanistic Characterization of RASGRP1 Variants Identifies an hnRNP-K-Regulated Transcriptional Enhancer Contributing to SLE Susceptibility. Frontiers in Immunology, 2019, 10, 1066.	2.2	13
52	Novel Risk Loci Identified in a Genome-Wide Association Study of Urolithiasis in a Japanese Population. Journal of the American Society of Nephrology: JASN, 2019, 30, 855-864.	3.0	25
53	Amino acid signatures of HLA Class-I and II molecules are strongly associated with SLE susceptibility and autoantibody production in Eastern Asians. PLoS Genetics, 2019, 15, e1008092.	1.5	36
54	Anti-nuclear antibody development is associated with poor treatment response to biological disease-modifying anti-rheumatic drugs in patients with rheumatoid arthritis. Seminars in Arthritis and Rheumatism, 2019, 49, 204-210.	1.6	7

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55	Genetic predisposition to mosaic Y chromosome loss in blood. Nature, 2019, 575, 652-657.	13.7	198
56	Genetic variants of calcium and vitamin D metabolism in kidney stone disease. Nature Communications, 2019, 10, 5175.	5.8	69
57	Linking of genetic risk variants to disease-specific gene expression via multi-omics studies in rheumatoid arthritis. Seminars in Arthritis and Rheumatism, 2019, 49, S49-S53.	1.6	11
58	Significant joint-destructive association of HLA-DRB1*04:05 independent of DAS28 in rheumatoid arthritis. Annals of the Rheumatic Diseases, 2019, 78, 284-286.	0.5	3
59	Clarification of the Basics in Takayasu Arteritis through Genetic and Epidemiological Approach. The Journal of the Japanese Society of Internal Medicine, 2019, 108, 791-797.	0.0	0
60	How to conduct research of rare autoimmune diseases. Modern Rheumatology, 2018, 28, 766-769.	0.9	1
61	<i>HLA-DRB1</i> Analysis Identified a Genetically Unique Subset within Rheumatoid Arthritis and Distinct Genetic Background of Rheumatoid Factor Levels from Anticyclic Citrullinated Peptide Antibodies. Journal of Rheumatology, 2018, 45, 470-480.	1.0	6
62	Integration of genetics and miRNA–target gene network identified disease biology implicated in tissue specificity. Nucleic Acids Research, 2018, 46, 11898-11909.	6.5	39
63	Genetic determinants and an epistasis of <i>LILRA3</i> and HLA-B*52 in Takayasu arteritis. Proceedings of the National Academy of Sciences of the United States of America, 2018, 115, 13045-13050.	3.3	51
64	Only rheumatoid factor-positive subset of anti-citrullinated peptide/protein antibody-negative rheumatoid arthritis may seroconvert to anti-citrullinated peptide/protein antibody-positive. International Journal of Rheumatic Diseases, 2017, 20, 731-736.	0.9	3
65	Transethnic meta-analysis identifies <i>CSDMA</i> and <i>PRDM1</i> as susceptibility genes to systemic sclerosis. Annals of the Rheumatic Diseases, 2017, 76, 1150-1158.	0.5	77
66	Genetic landscape of interactive effects of <i>HLA-DRB1</i> alleles on susceptibility to ACPA(+) rheumatoid arthritis and ACPA levels in Japanese population. Journal of Medical Genetics, 2017, 54, 853-858.	1.5	3
67	Time-averaged disease activity fits better joint destruction in rheumatoid arthritis. Scientific Reports, 2017, 7, 5856.	1.6	9
68	Anti-centromere antibody exhibits specific distribution levels among anti-nuclear antibodies and may characterize a distinct subset in rheumatoid arthritis. Scientific Reports, 2017, 7, 6911.	1.6	8
69	A novel susceptibility locus in the IL12B region is associated with the pathophysiology of Takayasu arteritis through IL-12p40 and IL-12p70 production. Arthritis Research and Therapy, 2017, 19, 197.	1.6	29
70	Smoking is associated with the concurrent presence of multiple autoantibodies in rheumatoid arthritis rather than with anti-citrullinated protein antibodies per se: a multicenter cohort study. Arthritis Research and Therapy, 2016, 18, 285.	1.6	43
71	A twin study of rheumatoid arthritis in the Japanese population. Modern Rheumatology, 2016, 26, 685-689.	0.9	32
72	Contribution of a Non-classical HLA Gene, HLA-DOA, to the Risk of Rheumatoid Arthritis. American Journal of Human Genetics, 2016, 99, 366-374.	2.6	68

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73	Genotyping of relapsing polychondritis identified novel susceptibility HLA alleles and distinct genetic characteristics from other rheumatic diseases. Rheumatology, 2016, 55, 1686-1692.	0.9	26
74	Recent Advances in Defining the Genetic Basis of Rheumatoid Arthritis. Annual Review of Genomics and Human Genetics, 2016, 17, 273-301.	2.5	44
75	The association between serious infection and disease outcome in patients with rheumatoid arthritis. Clinical Rheumatology, 2016, 35, 213-218.	1.0	10
76	Revisited HLA and non-HLA genetics of Takayasu arteritis—where are we?. Journal of Human Genetics, 2016, 61, 27-32.	1.1	60
77	Takayasu Arteritis and Ulcerative Colitis: A Frequent Association?. , 2016, , 63-77.		Ο
78	An Association Between Amino Acid Position 74 of HLA–DRB1 and Anti–Citrullinated Protein Antibody Levels in Japanese Patients With Anti–Citrullinated Protein Antibody–Positive Rheumatoid Arthritis. Arthritis and Rheumatology, 2015, 67, 2038-2045.	2.9	15
79	Rheumatoid Factor Is Associated With the Distribution of Hand Joint Destruction in Rheumatoid Arthritis. Arthritis and Rheumatology, 2015, 67, 3113-3123.	2.9	25
80	Periodontitis and Porphyromonas gingivalis in Preclinical Stage of Arthritis Patients. PLoS ONE, 2015, 10, e0122121.	1.1	48
81	Functional Impact and Evolution of a Novel Human Polymorphic Inversion That Disrupts a Gene and Creates a Fusion Transcript. PLoS Genetics, 2015, 11, e1005495.	1.5	22
82	Significant association of periodontal disease with anti-citrullinated peptide antibody in a Japanese healthy population – The Nagahama study. Journal of Autoimmunity, 2015, 59, 85-90.	3.0	36
83	Brief Report: Takayasu Arteritis and Ulcerative Colitis: High Rate of Coâ€Occurrence and Genetic Overlap. Arthritis and Rheumatology, 2015, 67, 2226-2232.	2.9	102
84	Brief Report: Main Contribution of DRB1*04:05 Among the Shared Epitope Alleles and Involvement of DRB1 Amino Acid Position 57 in Association With Joint Destruction in Anti–Citrullinated Protein Antibody–Positive Rheumatoid Arthritis. Arthritis and Rheumatology, 2015, 67, 1744-1750.	2.9	23
85	Anti-citrullinated peptide/protein antibody (ACPA)-negative RA shares a large proportion of susceptibility loci with ACPA-positive RA: a meta-analysis of genome-wide association study in a Japanese population. Arthritis Research and Therapy, 2015, 17, 104.	1.6	23
86	Increase of Hemoglobin Levels by Anti-IL-6 Receptor Antibody (Tocilizumab) in Rheumatoid Arthritis. PLoS ONE, 2014, 9, e98202.	1.1	39
87	Association Between Antinuclear Antibodies and the HLA Class II Locus and Heterogeneous Characteristics of Staining Patterns: The Nagahama Study. Arthritis and Rheumatology, 2014, 66, 3395-3403.	2.9	16
88	History of Takayasu arteritis and Dr. Mikito Takayasu. International Journal of Rheumatic Diseases, 2014, 17, 931-935.	0.9	28
89	Recent advances in <scp>T</scp> akayasu arteritis. International Journal of Rheumatic Diseases, 2014, 17, 238-247.	0.9	84
90	Genetics of rheumatoid arthritis contributes to biology and drug discovery. Nature, 2014, 506, 376-381.	13.7	1,974

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91	A Clinical, Pathological, and Genetic Characterization of Methotrexate-associated Lymphoproliferative Disorders. Journal of Rheumatology, 2014, 41, 293-299.	1.0	107
92	Effects of Smoking and Shared Epitope on the Production of Anti–Citrullinated Peptide Antibody in a Japanese Adult Population. Arthritis Care and Research, 2014, 66, 1818-1827.	1.5	61
93	Inverse Association between Air Pressure and Rheumatoid Arthritis Synovitis. PLoS ONE, 2014, 9, e85376.	1.1	24
94	<i>PLD4</i> as a novel susceptibility gene for systemic sclerosis in a Japanese population. Arthritis and Rheumatism, 2013, 65, 472-480.	6.7	62
95	Two Susceptibility Loci to Takayasu Arteritis Reveal a Synergistic Role of the IL12B and HLA-B Regions in a Japanese Population. American Journal of Human Genetics, 2013, 93, 289-297.	2.6	136
96	Association of Takayasu arteritis with HLA-B*67:01 and two amino acids in HLA-B protein. Rheumatology, 2013, 52, 1769-1774.	0.9	39
97	Genome-Wide Association Study and Gene Expression Analysis Identifies CD84 as a Predictor of Response to Etanercept Therapy in Rheumatoid Arthritis. PLoS Genetics, 2013, 9, e1003394.	1.5	146
98	Three Groups in the 28 Joints for Rheumatoid Arthritis Synovitis – Analysis Using More than 17,000 Assessments in the KURAMA Database. PLoS ONE, 2013, 8, e59341.	1.1	47
99	Functional Variants in NFKBIE and RTKN2 Involved in Activation of the NF-κB Pathway Are Associated with Rheumatoid Arthritis in Japanese. PLoS Genetics, 2012, 8, e1002949.	1.5	46
100	A Genome-Wide Association Study Identified AFF1 as a Susceptibility Locus for Systemic Lupus Eyrthematosus in Japanese. PLoS Genetics, 2012, 8, e1002455.	1.5	115
101	Serum IgG levels demonstrate seasonal change in connective tissue diseases: a large-scale, 4-year analysis in Japanese. Modern Rheumatology, 2012, 22, 426-430.	0.9	4
102	Quantitative effect of HLA-DRB1 alleles to ACPA levels in Japanese rheumatoid arthritis: no strong genetic impact of shared epitope to ACPA levels after stratification of HLA-DRB1*09:01. Annals of the Rheumatic Diseases, 2012, 71, 1095-1097.	0.5	19
103	Meta-analysis identifies nine new loci associated with rheumatoid arthritis in the Japanese population. Nature Genetics, 2012, 44, 511-516.	9.4	285
104	ACPA-Negative RA Consists of Two Genetically Distinct Subsets Based on RF Positivity in Japanese. PLoS ONE, 2012, 7, e40067.	1.1	33
105	Serum IgG levels demonstrate seasonal change in connective tissue diseases: a large-scale, 4-year analysis in Japanese. Modern Rheumatology, 2012, 22, 426-430.	0.9	3
106	A large-scale association study identified multiple HLA-DRB1 alleles associated with ACPA-negative rheumatoid arthritis in Japanese subjects. Annals of the Rheumatic Diseases, 2011, 70, 2134-2139.	0.5	42
107	Myelin Basic Protein as a Novel Genetic Risk Factor in Rheumatoid Arthritis—A Genome-Wide Study Combined with Immunological Analyses. PLoS ONE, 2011, 6, e20457.	1.1	29
108	The human AIRE gene at chromosome 21q22 is a genetic determinant for the predisposition to rheumatoid arthritis in Japanese population. Human Molecular Genetics, 2011, 20, 2680-2685.	1.4	90

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109	A regulatory variant in CCR6 is associated with rheumatoid arthritis susceptibility. Nature Genetics, 2010, 42, 515-519.	9.4	241
110	Anti-citrullinated peptide antibody-negative RA is a genetically distinct subset: a definitive study using only bone-erosive ACPA-negative rheumatoid arthritis. Rheumatology, 2010, 49, 2298-2304.	0.9	61