Leonid Padyukov

List of Publications by Year
in descending order

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		9775	6643
273	27,439	73	156
papers	citations	h-index	g-index

295	295	295	29912
all docs	docs citations	times ranked	citing authors

#	Article	IF	Citations
1	Genetics of rheumatoid arthritis contributes to biology and drug discovery. Nature, 2014, 506, 376-381.	13.7	1,974
2	A new model for an etiology of rheumatoid arthritis: Smoking may trigger HLAâ€“DR (shared) Tj ETQq0 0 0 rgBT /Overlock 10 Tf 50 707 Rheumatism, 2006, 54, 38-46.	6.7	1,233
3	Genome-wide association study meta-analysis identifies seven new rheumatoid arthritis risk loci. Nature Genetics, 2010, 42, 508-514.	9.4	1,132
4	Analysis of shared heritability in common disorders of the brain. Science, 2018, 360, .	6.0	1,085
5	STAT4and the Risk of Rheumatoid Arthritis and Systemic Lupus Erythematosus. New England Journal of Medicine, 2007, 357, 977-986.	13.9	914
6	Epigenome-wide association data implicate DNA methylation as an intermediary of genetic risk in rheumatoid arthritis. Nature Biotechnology, 2013, 31, 142-147.	9.4	874
7	Five amino acids in three HLA proteins explain most of the association between MHC and seropositive rheumatoid arthritis. Nature Genetics, 2012, 44, 291-296.	9.4	768
8	TRAF1â€“C5as a Risk Locus for Rheumatoid Arthritis â€” A Genomewide Study. New England Journal of Medicine, 2007, 357, 1199-1209.	13.9	729
9	A large-scale replication study identifies TNIP1, PRDM1, JAZF1, UHRF1BP1 and IL10 as risk loci for systemic lupus erythematosus. Nature Genetics, 2009, 41, 1228-1233.	9.4	729
10	Genome-wide association study identifies eight risk loci and implicates metabo-psychiatric origins for anorexia nervosa. Nature Genetics, 2019, 51, 1207-1214.	9.4	641
11	Genome-wide association identifies multiple ulcerative colitis susceptibility loci. Nature Genetics, 2010, 42, 332-337.	9.4	572
12	High-density genetic mapping identifies new susceptibility loci for rheumatoid arthritis. Nature Genetics, 2012, 44, 1336-1340.	9.4	558
13	A gene-environment interaction between smoking and shared epitope genes in HLA-DR provides a high risk of seropositive rheumatoid arthritis. Arthritis and Rheumatism, 2004, 50, 3085-3092.	6.7	546
14	Two independent alleles at 6q23 associated with risk of rheumatoid arthritis. Nature Genetics, 2007, 39, 1477-1482.	9.4	497
15	Replication of Putative Candidate-Gene Associations with Rheumatoid Arthritis in >4,000 Samples from North America and Sweden: Association of Susceptibility with PTPN22, CTLA4, and PADI4. American Journal of Human Genetics, 2005, 77, 1044-1060.	2.6	494
16	Common variants at CD40 and other loci confer risk of rheumatoid arthritis. Nature Genetics, 2008, 40, 1216-1223.	9.4	476
17	Immunity to Citrullinated Proteins in Rheumatoid Arthritis. Annual Review of Immunology, 2008, 26, 651-675.	9.5	400
18	Gene-Gene and Gene-Environment Interactions Involving HLA-DRB1, PTPN22, and Smoking in Two Subsets of Rheumatoid Arthritis. American Journal of Human Genetics, 2007, 80, 867-875.	2.6	374

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19	Dense genotyping of immune-related disease regions identifies nine new risk loci for primary sclerosing cholangitis. Nature Genetics, 2013, 45, 670-675.	9.4	339
20	Smoking is a major preventable risk factor for rheumatoid arthritis: estimations of risks after various exposures to cigarette smoke. Annals of the Rheumatic Diseases, 2011, 70, 508-511.	0.5	309
21	Genetic variants at CD28, PRDM1 and CD2/CD58 are associated with rheumatoid arthritis risk. Nature Genetics, 2009, 41, 1313-1318.	9.4	306
22	Meta-analysis identifies nine new loci associated with rheumatoid arthritis in the Japanese population. Nature Genetics, 2012, 44, 511-516.	9.4	285
23	Specific interaction between genotype, smoking and autoimmunity to citrullinated Î±-enolase in the etiology of rheumatoid arthritis. Nature Genetics, 2009, 41, 1319-1324.	9.4	282
24	MHC2TA is associated with differential MHC molecule expression and susceptibility to rheumatoid arthritis, multiple sclerosis and myocardial infarction. Nature Genetics, 2005, 37, 486-494.	9.4	276
25	A combination of autoantibodies to cyclic citrullinated peptide (CCP) and HLA-DRB1 locus antigens is strongly associated with future onset of rheumatoid arthritis. Arthritis Research, 2004, 6, R303.	2.0	243
26	A genome-wide association study suggests contrasting associations in ACPA-positive versus ACPA-negative rheumatoid arthritis. Annals of the Rheumatic Diseases, 2011, 70, 259-265.	0.5	238
27	A Candidate Gene Approach Identifies the TRAF1/C5 Region as a Risk Factor for Rheumatoid Arthritis. PLoS Medicine, 2007, 4, e278.	3.9	232
28	Mapping of multiple susceptibility variants within the MHC region for 7 immune-mediated diseases. Proceedings of the National Academy of Sciences of the United States of America, 2009, 106, 18680-18685.	3.3	231
29	A Role for Noncoding Variation in Schizophrenia. Cell Reports, 2014, 9, 1417-1429.	2.9	225
30	Identification of Novel Genetic Markers Associated with Clinical Phenotypes of Systemic Sclerosis through a Genome-Wide Association Strategy. PLoS Genetics, 2011, 7, e1002178.	1.5	201
31	Smoking, citrullination and genetic variability in the immunopathogenesis of rheumatoid arthritis. Seminars in Immunology, 2011, 23, 92-98.	2.7	195
32	Immunochip Analysis Identifies Multiple Susceptibility Loci for Systemic Sclerosis. American Journal of Human Genetics, 2014, 94, 47-61.	2.6	182
33	Genetic markers for the efficacy of tumour necrosis factor blocking therapy in rheumatoid arthritis. Annals of the Rheumatic Diseases, 2003, 62, 526-529.	0.5	175
34	Multiple antibody reactivities to citrullinated antigens in sera from patients with rheumatoid arthritis: association with HLA-DRB1 alleles. Annals of the Rheumatic Diseases, 2009, 68, 736-743.	0.5	175
35	Association of a haplotype in the promoter region of the interferon regulatory factor 5 gene with rheumatoid arthritis. Arthritis and Rheumatism, 2007, 56, 2202-2210.	6.7	174
36	Mechanisms of Disease: genetic susceptibility and environmental triggers in the development of rheumatoid arthritis. Nature Clinical Practice Rheumatology, 2006, 2, 425-433.	3.2	170

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37	Alcohol consumption is associated with decreased risk of rheumatoid arthritis: results from two Scandinavian caseâ€“control studies. Annals of the Rheumatic Diseases, 2009, 68, 222-227.	0.5	166
38	Smoking as a trigger for inflammatory rheumatic diseases. Current Opinion in Rheumatology, 2007, 19, 49-54.	2.0	162
39	Fine Mapping Seronegative and Seropositive Rheumatoid Arthritis to Shared and Distinct HLA Alleles by Adjusting for the Effects of Heterogeneity. American Journal of Human Genetics, 2014, 94, 522-532.	2.6	156
40	Genome-wide meta-analysis identifies multiple novel associations and ethnic heterogeneity of psoriasis susceptibility. Nature Communications, 2015, 6, 6916.	5.8	154
41	Genes, environment and immunity in the development of rheumatoid arthritis. Current Opinion in Immunology, 2006, 18, 650-655.	2.4	153
42	Additive effects of the major risk alleles of IRF5 and STAT4 in primary SjÃ¶gren's syndrome. Genes and Immunity, 2009, 10, 68-76.	2.2	152
43	Association of the PD-1.3A allele of thePDCD1gene in patients with rheumatoid arthritis negative for rheumatoid factor and the shared epitope. Arthritis and Rheumatism, 2004, 50, 1770-1773.	6.7	146
44	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
45	Increased expression of the novel proinflammatory cytokine high mobility group box chromosomal protein 1 in skin lesions of patients with lupus erythematosus. Arthritis and Rheumatism, 2005, 52, 3639-3645.	6.7	137
46	Genetic and environmental determinants for disease risk in subsets of rheumatoid arthritis defined by the anticitrullinated protein/peptide antibody fine specificity profile. Annals of the Rheumatic Diseases, 2013, 72, 652-658.	0.5	137
47	Protection against antiâ€“citrullinated protein antibodyâ€“positive rheumatoid arthritis is predominantly associated with HLAâ€“DRB1*1301: A metaâ€analysis of HLAâ€“DRB1 associations with antiâ€“citrullinated protein antibodyâ€“positive and antiâ€“citrullinated protein antibodyâ€“negative rheumatoid arthritis in four European populations. Arthritis and Rheumatism, 2010, 62, 1236-1245.	6.7	135
48	DNA methylation mapping identifies gene regulatory effects in patients with systemic lupus erythematosus. Annals of the Rheumatic Diseases, 2018, 77, 736-743.	0.5	135
49	Geneâ€“environment interaction between the DRB1 shared epitope and smoking in the risk of antiâ€“citrullinated protein antibodyâ€“positive rheumatoid arthritis: All alleles are important. Arthritis and Rheumatism, 2009, 60, 1597-1603.	6.7	129
50	Dense genotyping of immune-related loci in idiopathic inflammatory myopathies confirms HLA alleles as the strongest genetic risk factor and suggests different genetic background for major clinical subgroups. Annals of the Rheumatic Diseases, 2016, 75, 1558-1566.	0.5	127
51	Molecular mimicry between Anoctamin 2 and Epstein-Barr virus nuclear antigen 1 associates with multiple sclerosis risk. Proceedings of the National Academy of Sciences of the United States of America, 2019, 116, 16955-16960.	3.3	120
52	TYK2 Protein-Coding Variants Protect against Rheumatoid Arthritis and Autoimmunity, with No Evidence of Major Pleiotropic Effects on Non-Autoimmune Complex Traits. PLoS ONE, 2015, 10, e0122271.	1.1	120
53	Copy number, linkage disequilibrium and disease association in the FCGR locus. Human Molecular Genetics, 2010, 19, 3282-3294.	1.4	119
54	Different patterns of associations with antiâ€“citrullinated protein antibodyâ€“positive and antiâ€“citrullinated protein antibodyâ€“negative rheumatoid arthritis in the extended major histocompatibility complex region. Arthritis and Rheumatism, 2009, 60, 30-38.	6.7	113

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55	Genomeâ€Wide Association Study of Dermatomyositis Reveals Genetic Overlap With Other Autoimmune Disorders. Arthritis and Rheumatism, 2013, 65, 3239-3247.	6.7	113
56	Shared immunological targets in the lungs and joints of patients with rheumatoid arthritis: identification and validation. Annals of the Rheumatic Diseases, 2015, 74, 1772-1777.	0.5	112
57	Genome-wide association study identifies HLA 8.1 ancestral haplotype alleles as major genetic risk factors for myositis phenotypes. Genes and Immunity, 2015, 16, 470-480.	2.2	103
58	Interaction of HLA-DRB1*03 and smoking for the development of anti-Jo-1 antibodies in adult idiopathic inflammatory myopathies: a European-wide case study. Annals of the Rheumatic Diseases, 2012, 71, 961-965.	0.5	100
59	High-density genotyping of immune loci in Koreans and Europeans identifies eight new rheumatoid arthritis risk loci. Annals of the Rheumatic Diseases, 2015, 74, e13-e13.	0.5	100
60	GWAS for systemic sclerosis identifies multiple risk loci and highlights fibrotic and vasculopathy pathways. Nature Communications, 2019, 10, 4955.	5.8	100
61	Rheumatoid arthritis risk allele PTPRC is also associated with response to antiâ€“tumor necrosis factor Î± therapy. Arthritis and Rheumatism, 2010, 62, 1849-1861.	6.7	95
62	Genome-wide association analysis of anti-TNF drug response in patients with rheumatoid arthritis. Annals of the Rheumatic Diseases, 2013, 72, 1375-1381.	0.5	94
63	Identification of Immune-Relevant Factors Conferring Sarcoidosis Genetic Risk. American Journal of Respiratory and Critical Care Medicine, 2015, 192, 727-736.	2.5	94
64	Association of arthritis with a gene complex encoding Câ€type lectinâ€“like receptors. Arthritis and Rheumatism, 2007, 56, 2620-2632.	6.7	93
65	GWAS of Follicular Lymphoma Reveals Allelic Heterogeneity at 6p21.32 and Suggests Shared Genetic Susceptibility with Diffuse Large B-cell Lymphoma. PLoS Genetics, 2011, 7, e1001378.	1.5	93
66	GeMes, Clusters of DNA Methylation under Genetic Control, Can Inform Genetic and Epigenetic Analysis of Disease. American Journal of Human Genetics, 2014, 94, 485-495.	2.6	93
67	Genes identified in Asian SLE GWASs are also associated with SLE in Caucasian populations. European Journal of Human Genetics, 2013, 21, 994-999.	1.4	90
68	Polymorphism in the P2X7 receptor gene and survival in chronic lymphocytic leukaemia. Lancet, The, 2002, 360, 1935-1939.	6.3	88
69	A case-control study of rheumatoid arthritis identifies an associated single nucleotide polymorphism in the NCF4 gene, supporting a role for the NADPH-oxidase complex in autoimmunity. Arthritis Research and Therapy, 2007, 9, R98.	1.6	84
70	Rare, Low-Frequency, and Common Variants in the Protein-Coding Sequence of Biological Candidate Genes from GWASs Contribute to Risk of Rheumatoid Arthritis. American Journal of Human Genetics, 2013, 92, 15-27.	2.6	83
71	Focused HLA analysis in Caucasians with myositis identifies significant associations with autoantibody subgroups. Annals of the Rheumatic Diseases, 2019, 78, 996-1002.	0.5	81
72	Genome-wide meta-analysis reveals shared new loci in systemic seropositive rheumatic diseases. Annals of the Rheumatic Diseases, 2019, 78, 311-319.	0.5	81

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73	FLT3 stop mutation increases FLT3 ligand level and risk of autoimmune thyroid disease. Nature, 2020, 584, 619-623.	13.7	81
74	Association between occupational exposure to mineral oil and rheumatoid arthritis: results from the Swedish EIRA case-control study. Arthritis Research and Therapy, 2005, 7, R1296.	1.6	80
75	Association of soluble CD89 levels with disease progression but not susceptibility in IgA nephropathy. Kidney International, 2010, 78, 1281-1287.	2.6	79
76	Occupational exposure to textile dust increases the risk of rheumatoid arthritis: results from a Malaysian population-based caseâ€“control study. Annals of the Rheumatic Diseases, 2016, 75, 997-1002.	0.5	78
77	Specific association of type 1 diabetes mellitus with antiâ€“cyclic citrullinated peptideâ€“positive rheumatoid arthritis. Arthritis and Rheumatism, 2009, 60, 653-660.	6.7	76
78	Association of theâ˜1087 IL 10 gene polymorphism with severe chronic periodontitis in Swedish Caucasians. Journal of Clinical Periodontology, 2003, 30, 249-254.	2.3	75
79	Crowdsourced assessment of common genetic contribution to predicting anti-TNF treatment response in rheumatoid arthritis. Nature Communications, 2016, 7, 12460.	5.8	73
80	Use of a Multiethnic Approach to Identify Rheumatoid- Arthritis-Susceptibility Loci, 1p36 and 17q12. American Journal of Human Genetics, 2012, 90, 524-532.	2.6	69
81	A STAT4 risk allele is associated with ischaemic cerebrovascular events and anti-phospholipid antibodies in systemic lupus erythematosus. Annals of the Rheumatic Diseases, 2010, 69, 834-840.	0.5	68
82	High-Density Genetic Mapping Identifies New Susceptibility Variants in Sarcoidosis Phenotypes and Shows Genomic-driven Phenotypic Differences. American Journal of Respiratory and Critical Care Medicine, 2016, 193, 1008-1022.	2.5	68
83	Opposing effects of HLAâ€“DRB1*13 alleles on the risk of developing antiâ€“citrullinated protein antibodyâ€“positive and antiâ€“citrullinated protein antibodyâ€“negative rheumatoid arthritis. Arthritis and Rheumatism, 2009, 60, 924-930.	6.7	64
84	Variants in RUNX3 Contribute to Susceptibility to Psoriatic Arthritis, Exhibiting Further Common Ground With Ankylosing Spondylitis. Arthritis and Rheumatism, 2013, 65, 1224-1231.	6.7	63
85	Association of CLEC16A with human common variable immunodeficiency disorder and role in murine B cells. Nature Communications, 2015, 6, 6804.	5.8	63
86	A genome-wide association study of rheumatoid arthritis without antibodies against citrullinated peptides. Annals of the Rheumatic Diseases, 2015, 74, e15-e15.	0.5	62
87	Association of the -159 CD14 gene polymorphism and lack of association of the -308 TNFA and Q551R IL-4RA polymorphisms with severe chronic periodontitis in Swedish Caucasians. Journal of Clinical Periodontology, 2005, 32, 474-479.	2.3	61
88	Smoking interacts with HLA-DRB1 shared epitope in the development of anti-citrullinated protein antibody-positive rheumatoid arthritis: results from the Malaysian Epidemiological Investigation of Rheumatoid Arthritis (MyEIRA). Arthritis Research and Therapy, 2012, 14, R89.	1.6	61
89	FcÎ³ receptor type IIIA genotype and response to tumor necrosis factor Î±â€“blocking agents in patients with rheumatoid arthritis. Arthritis and Rheumatism, 2007, 56, 448-452.	6.7	60
90	A candidate gene study of the type I interferon pathway implicates IKBKE and IL8 as risk loci for SLE. European Journal of Human Genetics, 2011, 19, 479-484.	1.4	58

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91	A combined large-scale meta-analysis identifies COG6 as a novel shared risk locus for rheumatoid arthritis and systemic lupus erythematosus. Annals of the Rheumatic Diseases, 2017, 76, 286-294.	0.5	58
92	Dimensionality reduction reveals fine-scale structure in the Japanese population with consequences for polygenic risk prediction. Nature Communications, 2020, 11, 1569.	5.8	58
93	Genetics of rheumatoid arthritis. Seminars in Immunopathology, 2022, 44, 47-62.	2.8	57
94	HLA-DRB1* alleles and symptoms associated with Heerfordt's syndrome in sarcoidosis. European Respiratory Journal, 2011, 38, 1151-1157.	3.1	56
95	Novel risk genes for systemic lupus erythematosus predicted by random forest classification. Scientific Reports, 2017, 7, 6236.	1.6	54
96	High-Density SNP Mapping of the HLA Region Identifies Multiple Independent Susceptibility Loci Associated with Selective IgA Deficiency. PLoS Genetics, 2012, 8, e1002476.	1.5	53
97	New data and an old puzzle: the negative association between schizophrenia and rheumatoid arthritis. International Journal of Epidemiology, 2015, 44, 1706-1721.	0.9	53
98	A Role for VAV1 in Experimental Autoimmune Encephalomyelitis and Multiple Sclerosis. Science Translational Medicine, 2009, 1, 10ra21.	5.8	52
99	Human Genetics in Rheumatoid Arthritis Guides a High-Throughput Drug Screen of the CD40 Signaling Pathway. PLoS Genetics, 2013, 9, e1003487.	1.5	52
100	A Novel Sarcoidosis Risk Locus for Europeans on Chromosome 11q13.1. American Journal of Respiratory and Critical Care Medicine, 2012, 186, 877-885.	2.5	51
101	Haplotypes of the interleukin-4 receptor alpha chain gene associate with susceptibility to and severity of atopic asthma. Clinical and Experimental Allergy, 2004, 34, 1570-1575.	1.4	50
102	HLA-DRB1 and month of birth in multiple sclerosis. Neurology, 2009, 73, 2107-2111.	1.5	50
103	Dense genotyping of immune-related loci identifies HLA variants associated with increased risk of collagenous colitis. Gut, 2017, 66, 421-428.	6.1	50
104	HLA-DRB1*04/*13 alleles are associated with vascular disease and antiphospholipid antibodies in systemic lupus erythematosus. Annals of the Rheumatic Diseases, 2013, 72, 1018-1025.	0.5	49
105	DNA methylation mediates genotype and smoking interaction in the development of anti-citrullinated peptide antibody-positive rheumatoid arthritis. Arthritis Research and Therapy, 2017, 19, 71.	1.6	48
106	Genetic Association with ERAP1 in Psoriasis Is Confined to Disease Onset after Puberty and Not Dependent on HLA-C*06. Journal of Investigative Dermatology, 2013, 133, 411-417.	0.3	47
107	Brief Report: IRF4 Newly Identified as a Common Susceptibility Locus for Systemic Sclerosis and Rheumatoid Arthritis in a Crossâ€Disease Metaâ€Analysis of Genomeâ€Wide Association Studies. Arthritis and Rheumatology, 2016, 68, 2338-2344.	2.9	46
108	Effect of interactions of glutathione Sâ€transferase T1, M1, and P1 and HMOX1 gene promoter polymorphisms with heavy smoking on the risk of rheumatoid arthritis. Arthritis and Rheumatism, 2010, 62, 3196-3210.	6.7	45

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109	Polymorphisms in peptidylarginine deiminase (PADI) associate with rheumatoid arthritis in diverse Asian populations: evidence from MyEIRA study and meta-analysis. Arthritis Research and Therapy, 2012, 14, R250.	1.6	45
110	Serum RANKL levels associate with anti- citrullinated protein antibodies in early untreated rheumatoid arthritis and are modulated following methotrexate. Arthritis Research and Therapy, 2015, 17, 239.	1.6	45
111	Different allelic frequencies of several cytokine genes in Hong Kong Chinese and Swedish Caucasians. Genes and Immunity, 2001, 2, 280-283.	2.2	44
112	Fine mapping the TAGAP risk locus in rheumatoid arthritis. Genes and Immunity, 2011, 12, 314-318.	2.2	44
113	Genome-wide association study of response to methotrexate in early rheumatoid arthritis patients. Pharmacogenomics Journal, 2018, 18, 528-538.	0.9	42
114	Shared Epitope Alleles Remain A Risk Factor for Anti-Citrullinated Proteins Antibody (ACPA) â€“ Positive Rheumatoid Arthritis in Three Asian Ethnic Groups. PLoS ONE, 2011, 6, e21069.	1.1	42
115	Polymorphisms in Toll-like receptor 3 confer natural resistance to human herpes simplex virus type 2 infection. Journal of General Virology, 2012, 93, 1717-1724.	1.3	41
116	Immuneâ€Array Analysis in Sporadic Inclusion Body Myositis Reveals HLAâ€“DRB1 Amino Acid Heterogeneity Across the Myositis Spectrum. Arthritis and Rheumatology, 2017, 69, 1090-1099.	2.9	41
117	Very high levels of antiâ€“citrullinated protein antibodies are associated with HLAâ€“DRB1*15 nonâ€“shared epitope allele in patients with rheumatoid arthritis. Arthritis and Rheumatism, 2012, 64, 2078-2084.	6.7	40
118	Histological antiphospholipid-associated nephropathy versus lupus nephritis in patients with systemic lupus erythematosus: an observational cross-sectional study with longitudinal follow-up. Arthritis Research and Therapy, 2015, 17, 109.	1.6	40
119	Endometriosis and autoimmune disease: association of susceptibility to moderate/severe endometriosis with CCL21 and HLA-DRB1. Fertility and Sterility, 2011, 95, 437-440.	0.5	39
120	Integration of Known DNA, RNA and Protein Biomarkers Provides Prediction of Anti-TNF Response in Rheumatoid Arthritis: Results from the COMBINE Study. Molecular Medicine, 2016, 22, 322-328.	1.9	39
121	Polymorphisms of the ITGAM Gene Confer Higher Risk of Discoid Cutaneous Than of Systemic Lupus Erythematosus. PLoS ONE, 2010, 5, e14212.	1.1	39
122	Patterns of interaction between genetic and nongenetic attributes and methotrexate efficacy in rheumatoid arthritis. Pharmacogenetics and Genomics, 2012, 22, 1-9.	0.7	38
123	What precedes development of rheumatoid arthritis?. Annals of the Rheumatic Diseases, 2004, 63, ii28-ii31.	0.5	36
124	Gene copy-number variations (CNVs) of complement C4 and C4A deficiency in genetic risk and pathogenesis of juvenile dermatomyositis. Annals of the Rheumatic Diseases, 2016, 75, 1599-1606.	0.5	36
125	The importance of differences; On environment and its interactions with genes and immunity in the causation of rheumatoid arthritis. Journal of Internal Medicine, 2020, 287, 514-533.	2.7	36
126	Molecular pathways in patients with systemic lupus erythematosus revealed by gene-centred DNA sequencing. Annals of the Rheumatic Diseases, 2021, 80, 109-117.	0.5	35

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127	Cytokines in the Placenta of Pakistani Newborns With and Without Intrauterine Growth Retardation. Pediatric Research, 2006, 59, 254-258.	1.1	34
128	Integration of Sequence Data from a Consanguineous Family with Genetic Data from an Outbred Population Identifies PLB1 as a Candidate Rheumatoid Arthritis Risk Gene. PLoS ONE, 2014, 9, e87645.	1.1	34
129	A Geneâ€"Environment Interaction Between Smoking and Gene polymorphisms Provides a High Risk of Two Subgroups of Sarcoidosis. Scientific Reports, 2019, 9, 18633.	1.6	34
130	Association of the proinflammatory haplotype (MICA5.1/TNF2/TNFa2/DRB1*03) with polymyositis and dermatomyositis. Arthritis and Rheumatism, 2004, 50, 1013-1015.	6.7	33
131	Influence ofFCGR3Agenotype on the therapeutic response to rituximab in rheumatoid arthritis: an observational cohort study. BMJ Open, 2012, 2, e001524.	0.8	33
132	Development of autoantibodies against muscle-specific FHL1 in severe inflammatory myopathies. Journal of Clinical Investigation, 2015, 125, 4612-4624.	3.9	33
133	Genetic variation in the transforming growth factor-Â1 gene is associated with susceptibility to IgA nephropathy. Nephrology Dialysis Transplantation, 2009, 24, 3061-3067.	0.4	31
134	<scp>HLA</scp>â€alleles associated with increased risk for extraâ€pulmonary involvement in sarcoidosis. Tissue Antigens, 2014, 83, 267-272.	1.0	31
135	SNP Variants in Major Histocompatibility Complex Are Associated with Sarcoidosis Susceptibilityâ€"A Joint Analysis in Four European Populations. Frontiers in Immunology, 2017, 8, 422.	2.2	31
136	Cellular distribution of the C-type II lectin dendritic cell immunoreceptor (DCIR) and its expression in the rheumatic joint: identification of a subpopulation of DCIR+ T cells. Annals of the Rheumatic Diseases, 2008, 67, 1742-1749.	0.5	30
137	Genetic variations in the serotonin 5-HT2A receptor gene (HTR2A) are associated with rheumatoid arthritis. Annals of the Rheumatic Diseases, 2008, 67, 1111-1115.	0.5	30
138	Analysis of Neuropeptide S Receptor Gene (NPSR1) Polymorphism in Rheumatoid Arthritis. PLoS ONE, 2010, 5, e9315.	1.1	30
139	The balance of expression of PTPN22 splice forms is significantly different in rheumatoid arthritis patients compared with controls. Genome Medicine, 2012, 4, 2.	3.6	30
140	Implication of IL-2/IL-21 region in systemic sclerosis genetic susceptibility. Annals of the Rheumatic Diseases, 2013, 72, 1233-1238.	0.5	30
141	Autoantibody Specificities and Type I Interferon Pathway Activation in Idiopathic Inflammatory Myopathies. Scandinavian Journal of Immunology, 2016, 84, 100-109.	1.3	30
142	Anticollagen type II antibodies are associated with an acute onset rheumatoid arthritis phenotype and prognosticate lower degree of inflammation during 5â€…years follow-up. Annals of the Rheumatic Diseases, 2017, 76, 1529-1536.	0.5	30
143	T cells are influenced by a long non-coding RNA in the autoimmune associated PTPN2 locus. Journal of Autoimmunity, 2018, 90, 28-38.	3.0	29
144	Contribution of IKBKE and IFIH1 gene variants to SLE susceptibility. Genes and Immunity, 2013, 14, 217-222.	2.2	28

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145	Genetic overlap between autoimmune diseases and nonâ€Hodgkin lymphoma subtypes. Genetic Epidemiology, 2019, 43, 844-863.	0.6	28
146	Shared genetic risk between eating disorderâ€•and substanceâ€useâ€related phenotypes: Evidence from genomeâ€wide association studies. Addiction Biology, 2021, 26, e12880.	1.4	28
147	The A-1087IL-10 allele is associated with cardiovascular disease in SLE. Atherosclerosis, 2004, 177, 409-414.	0.4	27
148	Distinct HLA Associations with Rheumatoid Arthritis Subsets Defined by Serological Subphenotype. American Journal of Human Genetics, 2019, 105, 616-624.	2.6	27
149	Sequencing of the MHC region defines HLA-DQA1 as the major genetic risk for seropositive rheumatoid arthritis in Han Chinese population. Annals of the Rheumatic Diseases, 2019, 78, 773-780.	0.5	27
150	Soluble urokinase plasminogen activator receptor (suPAR) levels predict damage accrual in patients with recent-onset systemic lupus erythematosus. Journal of Autoimmunity, 2020, 106, 102340.	3.0	27
151	Coding Variants at Hexa-allelic Amino Acid 13 of HLA-DRB1 Explain Independent SNP Associations with Follicular Lymphoma Risk. American Journal of Human Genetics, 2013, 93, 167-172.	2.6	26
152	An Immunochip-based interaction study of contrasting interaction effects with smoking in ACPA-positive versus ACPA-negative rheumatoid arthritis. Rheumatology, 2016, 55, 149-155.	0.9	26
153	Multiomics analysis of rheumatoid arthritis yields sequence variants that have large effects on risk of the seropositive subset. Annals of the Rheumatic Diseases, 2022, 81, 1085-1095.	0.5	26
154	Genotyping of immune-related genetic variants identifiesTYK2as a novel associated locus for idiopathic inflammatory myopathies. Annals of the Rheumatic Diseases, 2014, 73, 1750-1752.	0.5	25
155	Discovery of new candidate genes for rheumatoid arthritis through integration of genetic association data with expression pathway analysis. Arthritis Research and Therapy, 2017, 19, 19.	1.6	25
156	Association of response to TNF inhibitors in rheumatoid arthritis with quantitative trait loci for CD40 and CD39. Annals of the Rheumatic Diseases, 2019, 78, 1055-1061.	0.5	25
157	Four Systemic Lupus Erythematosus Subgroups, Defined by Autoantibodies Status, Differ Regarding HLAâ€DRB1 Genotype Associations and Immunological and Clinical Manifestations. ACR Open Rheumatology, 2022, 4, 27-39.	0.9	25
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