

Leonid Padyukov

List of Publications by Year in Descending Order

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The third column is the impact factor (IF) of the journal, and the fourth column is the number of citations of the article.

255
papers

21,595
citations

68
h-index

143
g-index

295
ext. papers

25,116
ext. citations

9
avg, IF

5.79
L-index

#	Paper	IF	Citations
255	Genetics of rheumatoid arthritis.. <i>Seminars in Immunopathology</i> , 2022 , 44, 47	12	1
254	Four Systemic Lupus Erythematosus Subgroups, Defined by Autoantibodies Status, Differ Regarding HLA-DRB1 Genotype Associations and Immunological and Clinical Manifestations. <i>ACR Open Rheumatology</i> , 2021 ,	3.5	2
253	Clinical characteristics of Vietnamese patients with idiopathic inflammatory myopathies and autoantibodies to aminoacyl-transfer RNA synthetases. <i>International Journal of Rheumatic Diseases</i> , 2021 , 24, 663-670	2.3	
252	Age at menarche, age at natural menopause, and risk of rheumatoid arthritis - a Mendelian randomization study. <i>Arthritis Research and Therapy</i> , 2021 , 23, 108	5.7	2
251	HLA-B*08 Identified as the Most Prominently Associated Major Histocompatibility Complex Locus for Anti-Carbamylated Protein Antibody-Positive/Anti-Cyclic Citrullinated Peptide-Negative Rheumatoid Arthritis. <i>Arthritis and Rheumatology</i> , 2021 , 73, 963-969	9.5	6
250	POS0355 ASSOCIATIONS BETWEEN HLA-DRB1 SHARED EPITOPES ALLELES AND ANTI-RA33 ANTIBODIES IN DIFFERENT SUBSETS OF RHEUMATOID ARTHRITIS IN MALAYSIAN POPULATION. <i>Annals of the Rheumatic Diseases</i> , 2021 , 80, 408.1-408	2.4	
249	POS0348 GENETIC SUSCEPTIBILITY VARIANTS FOR RHEUMATOID ARTHRITIS ARE NOT ASSOCIATED WITH EARLY REMISSION; A MULTI-COHORT STUDY. <i>Annals of the Rheumatic Diseases</i> , 2021 , 80, 403.1-404	2.4	
248	POS0457 CHANGES OF RF ISOTYPE PROFILE IN PATIENTS WITH RHEUMATOID ARTHRITIS: DATA FROM 10 YEARS FOLLOW-UP STUDY. <i>Annals of the Rheumatic Diseases</i> , 2021 , 80, 459-460	2.4	
247	OP0021 IDENTIFICATION OF DIFFERENTIALLY EXPRESSED GENES IN EARLY RHEUMATOID ARTHRITIS PATIENTS RESPONDING TO TOCILIZUMAB. <i>Annals of the Rheumatic Diseases</i> , 2021 , 80, 12.2-12 ⁴		
246	Early Prediction of Clinical Response to Anti-TNF Treatment using Multi-omics and Machine Learning in Rheumatoid Arthritis. <i>Rheumatology</i> , 2021 ,	3.9	6
245	Shared genetic risk between eating disorder- and substance-use-related phenotypes: Evidence from genome-wide association studies. <i>Addiction Biology</i> , 2021 , 26, e12880	4.6	12
244	Molecular pathways in patients with systemic lupus erythematosus revealed by gene-centred DNA sequencing. <i>Annals of the Rheumatic Diseases</i> , 2021 , 80, 109-117	2.4	10
243	Obesity-Related Traits and the Development of Rheumatoid Arthritis: Evidence From Genetic Data. <i>Arthritis and Rheumatology</i> , 2021 , 73, 203-211	9.5	9
242	The spectrum of association in HLA region with rheumatoid arthritis in a diverse Asian population: evidence from the MyEIRA case-control study. <i>Arthritis Research and Therapy</i> , 2021 , 23, 46	5.7	3
241	Contribution of rare genetic variation to disease susceptibility in a large Scandinavian myositis cohort. <i>Arthritis and Rheumatology</i> , 2021 ,	9.5	2
240	Reply. <i>Arthritis and Rheumatology</i> , 2021 , 73, 1944-1945	9.5	
239	Haplotype-Specific Expression Analysis of MHC Class II Genes in Healthy Individuals and Rheumatoid Arthritis Patients. <i>Frontiers in Immunology</i> , 2021 , 12, 707217	8.4	1

238	HLA-B*27 is significantly enriched in Nordic patients with psoriatic arthritis mutilans. <i>Clinical and Experimental Rheumatology</i> , 2021 , 39, 775-780	2.2	1
237	FLT3 stop mutation increases FLT3 ligand level and risk of autoimmune thyroid disease. <i>Nature</i> , 2020 , 584, 619-623	50.4	23
236	The importance of differences; On environment and its interactions with genes and immunity in the causation of rheumatoid arthritis. <i>Journal of Internal Medicine</i> , 2020 , 287, 514-533	10.8	16
235	Dimensionality reduction reveals fine-scale structure in the Japanese population with consequences for polygenic risk prediction. <i>Nature Communications</i> , 2020 , 11, 1569	17.4	22
234	Soluble urokinase plasminogen activator receptor (suPAR) levels predict damage accrual in patients with recent-onset systemic lupus erythematosus. <i>Journal of Autoimmunity</i> , 2020 , 106, 102340	15.5	15
233	Cigarette smoking patterns preceding primary Sjögren's syndrome. <i>RMD Open</i> , 2020 , 6,	5.9	2
232	Distinct HLA Associations with Rheumatoid Arthritis Subsets Defined by Serological Subphenotype. <i>American Journal of Human Genetics</i> , 2019 , 105, 616-624	11	16
231	Osteopontin and Disease Activity in Patients with Recent-onset Systemic Lupus Erythematosus: Results from the SLICC Inception Cohort. <i>Journal of Rheumatology</i> , 2019 , 46, 492-500	4.1	7
230	Focused HLA analysis in Caucasians with myositis identifies significant associations with autoantibody subgroups. <i>Annals of the Rheumatic Diseases</i> , 2019 , 78, 996-1002	2.4	48
229	Association of response to TNF inhibitors in rheumatoid arthritis with quantitative trait loci for and CD39. <i>Annals of the Rheumatic Diseases</i> , 2019 , 78, 1055-1061	2.4	12
228	Sequencing of the MHC region defines as the major genetic risk for seropositive rheumatoid arthritis in Han Chinese population. <i>Annals of the Rheumatic Diseases</i> , 2019 , 78, 773-780	2.4	16
227	Molecular mimicry between Anoctamin 2 and Epstein-Barr virus nuclear antigen 1 associates with multiple sclerosis risk. <i>Proceedings of the National Academy of Sciences of the United States of America</i> , 2019 , 116, 16955-16960	11.5	53
226	Genetic overlap between autoimmune diseases and non-Hodgkin lymphoma subtypes. <i>Genetic Epidemiology</i> , 2019 , 43, 844-863	2.6	15
225	Genome-wide association study identifies eight risk loci and implicates metabo-psychiatric origins for anorexia nervosa. <i>Nature Genetics</i> , 2019 , 51, 1207-1214	36.3	303
224	GWAS for systemic sclerosis identifies multiple risk loci and highlights fibrotic and vasculopathy pathways. <i>Nature Communications</i> , 2019 , 10, 4955	17.4	46
223	A Gene-Environment Interaction Between Smoking and Gene polymorphisms Provides a High Risk of Two Subgroups of Sarcoidosis. <i>Scientific Reports</i> , 2019 , 9, 18633	4.9	10
222	Genome-wide meta-analysis reveals shared new in systemic seropositive rheumatic diseases. <i>Annals of the Rheumatic Diseases</i> , 2019 , 78, 311-319	2.4	41
221	Clinical phenotype, autoantibody profile and HLA-DR-type in Vietnamese patients with idiopathic inflammatory myopathies. <i>Rheumatology</i> , 2019 , 58, 361-363	3.9	4

220	T cells are influenced by a long non-coding RNA in the autoimmune associated PTPN2 locus. <i>Journal of Autoimmunity</i> , 2018 , 90, 28-38	15.5	16
219	DNA methylation mapping identifies gene regulatory effects in patients with systemic lupus erythematosus. <i>Annals of the Rheumatic Diseases</i> , 2018 , 77, 736-743	2.4	76
218	Systematic approach demonstrates enrichment of multiple interactions between non- risk variants and risk alleles in rheumatoid arthritis. <i>Annals of the Rheumatic Diseases</i> , 2018 , 77, 1454-1462	2.4	17
217	Analysis of shared heritability in common disorders of the brain. <i>Science</i> , 2018 , 360,	33.3	666
216	T-cell transcriptomics from peripheral blood highlights differences between polymyositis and dermatomyositis patients. <i>Arthritis Research and Therapy</i> , 2018 , 20, 188	5.7	10
215	Genome-wide association study of response to methotrexate in early rheumatoid arthritis patients. <i>Pharmacogenomics Journal</i> , 2018 , 18, 528-538	3.5	25
214	Dense genotyping of immune-related loci identifies HLA variants associated with increased risk of collagenous colitis. <i>Gut</i> , 2017 , 66, 421-428	19.2	33
213	VAV1 regulates experimental autoimmune arthritis and is associated with anti-CCP negative rheumatoid arthritis. <i>Genes and Immunity</i> , 2017 , 18, 48-56	4.4	9
212	Immune-Array Analysis in Sporadic Inclusion Body Myositis Reveals HLA-DRB1 Amino Acid Heterogeneity Across the Myositis Spectrum. <i>Arthritis and Rheumatology</i> , 2017 , 69, 1090-1099	9.5	24
211	A combined large-scale meta-analysis identifies COG6 as a novel shared risk locus for rheumatoid arthritis and systemic lupus erythematosus. <i>Annals of the Rheumatic Diseases</i> , 2017 , 76, 286-294	2.4	39
210	DNA methylation mediates genotype and smoking interaction in the development of anti-citrullinated peptide antibody-positive rheumatoid arthritis. <i>Arthritis Research and Therapy</i> , 2017 , 19, 71	5.7	36
209	Genetic variation and epigenetic modification of the prodynorphin gene in peripheral blood cells in alcoholism. <i>Progress in Neuro-Psychopharmacology and Biological Psychiatry</i> , 2017 , 76, 195-203	5.5	13
208	Possible Interaction Between Cigarette Smoking and HLA-DRB1 Variation in the Risk of Follicular Lymphoma. <i>American Journal of Epidemiology</i> , 2017 , 185, 681-687	3.8	7
207	Anticollagen type II antibodies are associated with an acute onset rheumatoid arthritis phenotype and prognosticate lower degree of inflammation during 5 years follow-up. <i>Annals of the Rheumatic Diseases</i> , 2017 , 76, 1529-1536	2.4	18
206	Depressed serum IgM levels in SLE are restricted to defined subgroups. <i>Clinical Immunology</i> , 2017 , 183, 304-315	9	15
205	Cutting Edge: Genetic Association between IFI16 Single Nucleotide Polymorphisms and Resistance to Genital Herpes Correlates with IFI16 Expression Levels and HSV-2-Induced IFN- γ Expression. <i>Journal of Immunology</i> , 2017 , 199, 2613-2617	5.3	13
204	Common variants of T-cells contribute differently to phenotypic variation in sarcoidosis. <i>Scientific Reports</i> , 2017 , 7, 5623	4.9	7
203	Novel risk genes for systemic lupus erythematosus predicted by random forest classification. <i>Scientific Reports</i> , 2017 , 7, 6236	4.9	31

202	Discovery of new candidate genes for rheumatoid arthritis through integration of genetic association data with expression pathway analysis. <i>Arthritis Research and Therapy</i> , 2017 , 19, 19	5.7	15
201	Differences in the Spectrum of Anti-Citrullinated Protein Antibody Fine Specificities Between Malaysian and Swedish Patients With Rheumatoid Arthritis: Implications for Disease Pathogenesis. <i>Arthritis and Rheumatology</i> , 2017 , 69, 58-69	9.5	9
200	SNP Variants in Major Histocompatibility Complex Are Associated with Sarcoidosis Susceptibility-A Joint Analysis in Four European Populations. <i>Frontiers in Immunology</i> , 2017 , 8, 422	8.4	20
199	FRI0596 Rheumatoid Arthritis Risk Alleles, Hla-Drb1 Haplotypes, and Response To TNFi Therapy □ Results from A Swedish Cohort Study. <i>Annals of the Rheumatic Diseases</i> , 2016 , 75, 658.1-658	2.4	
198	Crowdsourced assessment of common genetic contribution to predicting anti-TNF treatment response in rheumatoid arthritis. <i>Nature Communications</i> , 2016 , 7, 12460	17.4	54
197	Conserved 33-kb haplotype in the MHC class III region regulates chronic arthritis. <i>Proceedings of the National Academy of Sciences of the United States of America</i> , 2016 , 113, E3716-24	11.5	14
196	Brief Report: IRF4 Newly Identified as a Common Susceptibility Locus for Systemic Sclerosis and Rheumatoid Arthritis in a Cross-Disease Meta-Analysis of Genome-Wide Association Studies. <i>Arthritis and Rheumatology</i> , 2016 , 68, 2338-44	9.5	35
195	High-Density Genetic Mapping Identifies New Susceptibility Variants in Sarcoidosis Phenotypes and Shows Genomic-driven Phenotypic Differences. <i>American Journal of Respiratory and Critical Care Medicine</i> , 2016 , 193, 1008-22	10.2	44
194	An ImmunoChip-based interaction study of contrasting interaction effects with smoking in ACPA-positive versus ACPA-negative rheumatoid arthritis. <i>Rheumatology</i> , 2016 , 55, 149-55	3.9	21
193	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. <i>Annals of the Rheumatic Diseases</i> , 2016 , 75, 1558-66	2.4	85
192	Gene copy-number variations (CNVs) of complement C4 and C4A deficiency in genetic risk and pathogenesis of juvenile dermatomyositis. <i>Annals of the Rheumatic Diseases</i> , 2016 , 75, 1599-606	2.4	27
191	Integration of known DNA, RNA and protein biomarkers provides prediction of anti-TNF response in rheumatoid arthritis: results from the COMBINE study. <i>Molecular Medicine</i> , 2016 , 22, 322-328	6.2	28
190	Autoantibody Specificities and Type I Interferon Pathway Activation in Idiopathic Inflammatory Myopathies. <i>Scandinavian Journal of Immunology</i> , 2016 , 84, 100-9	3.4	18
189	Occupational exposure to textile dust increases the risk of rheumatoid arthritis: results from a Malaysian population-based case-control study. <i>Annals of the Rheumatic Diseases</i> , 2016 , 75, 997-1002	2.4	57
188	A genetic risk score composed of rheumatoid arthritis risk alleles, HLA-DRB1 haplotypes, and response to TNFi therapy - results from a Swedish cohort study. <i>Arthritis Research and Therapy</i> , 2016 , 18, 288	5.7	4
187	HLA Associations Distinguish Collagenous From Lymphocytic Colitis. <i>American Journal of Gastroenterology</i> , 2016 , 111, 1211-3	0.7	16
186	Reply to Liu et al.: Translation of rat congenic data to humans on a conserved MHC-III haplotype associated with rheumatoid arthritis. <i>Proceedings of the National Academy of Sciences of the United States of America</i> , 2016 , 113, E6323-E6324	11.5	2
185	High-density genotyping of immune loci in Koreans and Europeans identifies eight new rheumatoid arthritis risk loci. <i>Annals of the Rheumatic Diseases</i> , 2015 , 74, e13	2.4	76

184	A genome-wide association study of rheumatoid arthritis without antibodies against citrullinated peptides. <i>Annals of the Rheumatic Diseases</i> , 2015 , 74, e15	2.4	49
183	Association of CLEC16A with human common variable immunodeficiency disorder and role in murine B cells. <i>Nature Communications</i> , 2015 , 6, 6804	17.4	53
182	Histological antiphospholipid-associated nephropathy versus lupus nephritis in patients with systemic lupus erythematosus: an observational cross-sectional study with longitudinal follow-up. <i>Arthritis Research and Therapy</i> , 2015 , 17, 109	5.7	23
181	Genome-wide association study identifies HLA 8.1 ancestral haplotype alleles as major genetic risk factors for myositis phenotypes. <i>Genes and Immunity</i> , 2015 , 16, 470-80	4.4	75
180	New data and an old puzzle: the negative association between schizophrenia and rheumatoid arthritis. <i>International Journal of Epidemiology</i> , 2015 , 44, 1706-21	7.8	43
179	Shared immunological targets in the lungs and joints of patients with rheumatoid arthritis: identification and validation. <i>Annals of the Rheumatic Diseases</i> , 2015 , 74, 1772-7	2.4	86
178	Serum RANKL levels associate with anti-citrullinated protein antibodies in early untreated rheumatoid arthritis and are modulated following methotrexate. <i>Arthritis Research and Therapy</i> , 2015 , 17, 239	5.7	36
177	Genetic vectors as a tool in association studies: definitions and application for study of rheumatoid arthritis. <i>International Journal of Genomics</i> , 2015 , 2015, 256818	2.5	
176	Genome-wide meta-analysis identifies multiple novel associations and ethnic heterogeneity of psoriasis susceptibility. <i>Nature Communications</i> , 2015 , 6, 6916	17.4	115
175	Gene-gene interaction and RNA splicing profiles of MAP2K4 gene in rheumatoid arthritis. <i>Clinical Immunology</i> , 2015 , 158, 19-28	9	10
174	Identification of Immune-Relevant Factors Conferring Sarcoidosis Genetic Risk. <i>American Journal of Respiratory and Critical Care Medicine</i> , 2015 , 192, 727-36	10.2	66
173	Identification of secreted phosphoprotein 1 gene as a new rheumatoid arthritis susceptibility gene. <i>Annals of the Rheumatic Diseases</i> , 2015 , 74, e19	2.4	16
172	Development of autoantibodies against muscle-specific FHL1 in severe inflammatory myopathies. <i>Journal of Clinical Investigation</i> , 2015 , 125, 4612-24	15.9	27
171	TYK2 protein-coding variants protect against rheumatoid arthritis and autoimmunity, with no evidence of major pleiotropic effects on non-autoimmune complex traits. <i>PLoS ONE</i> , 2015 , 10, e0122271	3.7	77
170	IgA measurements in over 12 000 Swedish twins reveal sex differential heritability and regulatory locus near CD30L. <i>Human Molecular Genetics</i> , 2014 , 23, 4177-84	5.6	9
169	GeMes, clusters of DNA methylation under genetic control, can inform genetic and epigenetic analysis of disease. <i>American Journal of Human Genetics</i> , 2014 , 94, 485-95	11	76
168	Genetics of rheumatoid arthritis contributes to biology and drug discovery. <i>Nature</i> , 2014 , 506, 376-81	50.4	1426
167	A 129-kb deletion on chromosome 12 confers substantial protection against rheumatoid arthritis, implicating the gene SLC2A3. <i>Human Mutation</i> , 2014 , 35, 248-56	4.7	19

166	Thyroxin substitution and the risk of developing rheumatoid arthritis; results from the Swedish population-based EIRA study. <i>Annals of the Rheumatic Diseases</i> , 2014 , 73, 1096-100	2.4	15
165	Fine mapping seronegative and seropositive rheumatoid arthritis to shared and distinct HLA alleles by adjusting for the effects of heterogeneity. <i>American Journal of Human Genetics</i> , 2014 , 94, 522-32	11	132
164	ImmunoChip analysis identifies multiple susceptibility loci for systemic sclerosis. <i>American Journal of Human Genetics</i> , 2014 , 94, 47-61	11	151
163	THU0478 Genome-Wide Association Analysis of Pain Reduction in Rheumatoid Arthritis Patients Treated with TNF Inhibitors. <i>Annals of the Rheumatic Diseases</i> , 2014 , 73, 348.3-348	2.4	
162	FRI0520 Association Study of the BAFF Genetic Variations in Two Independent Cohorts with Idiopathic Inflammatory Myopathies. <i>Annals of the Rheumatic Diseases</i> , 2014 , 73, 575.3-576	2.4	
161	Genetic Vectors Approach in a Study of Fine Structure of Interaction Between Risk Haplotype of HTR2A and HLA-DRB1 Shared Epitope Alleles in Rheumatoid Arthritis 2014 , 151-174		
160	Integration of sequence data from a Consanguineous family with genetic data from an outbred population identifies PLB1 as a candidate rheumatoid arthritis risk gene. <i>PLoS ONE</i> , 2014 , 9, e87645	3.7	24
159	Functional Studies of Gene-Gene Interaction of Autoimmune Diseases 2014 , 137-150		
158	Gene-Gene and Gene-Environment Interaction in Rheumatoid Arthritis 2014 , 85-100		
157	A role for noncoding variation in schizophrenia. <i>Cell Reports</i> , 2014 , 9, 1417-29	10.6	174
156	Variability in the CIITA gene interacts with HLA in multiple sclerosis. <i>Genes and Immunity</i> , 2014 , 15, 162-74.4	9	
155	HLA-alleles associated with increased risk for extra-pulmonary involvement in sarcoidosis. <i>Tissue Antigens</i> , 2014 , 83, 267-72		20
154	Genotyping of immune-related genetic variants identifies TYK2 as a novel associated locus for idiopathic inflammatory myopathies. <i>Annals of the Rheumatic Diseases</i> , 2014 , 73, 1750-2	2.4	21
153	Non-HLA genes PTPN22, CDK6 and PADI4 are associated with specific autoantibodies in HLA-defined subgroups of rheumatoid arthritis. <i>Arthritis Research and Therapy</i> , 2014 , 16, 414	5.7	21
152	Genetic evidence for involvement of adaptive immunity in the development of IgA nephropathy: MHC class II alleles are protective in a Caucasian population. <i>Human Immunology</i> , 2013 , 74, 957-60	2.3	6
151	Coding variants at hexa-allelic amino acid 13 of HLA-DRB1 explain independent SNP associations with follicular lymphoma risk. <i>American Journal of Human Genetics</i> , 2013 , 93, 167-72	11	26
150	Genome-wide association analysis of anti-TNF drug response in patients with rheumatoid arthritis. <i>Annals of the Rheumatic Diseases</i> , 2013 , 72, 1375-81	2.4	79
149	Epigenome-wide association data implicate DNA methylation as an intermediary of genetic risk in rheumatoid arthritis. <i>Nature Biotechnology</i> , 2013 , 31, 142-7	44.5	691

148	Genetic variation in the serotonin receptor gene affects immune responses in rheumatoid arthritis. <i>Genes and Immunity</i> , 2013 , 14, 83-9	4.4	15
147	Rare, low-frequency, and common variants in the protein-coding sequence of biological candidate genes from GWASs contribute to risk of rheumatoid arthritis. <i>American Journal of Human Genetics</i> , 2013 , 92, 15-27	11	72
146	Dense genotyping of immune-related disease regions identifies nine new risk loci for primary sclerosing cholangitis. <i>Nature Genetics</i> , 2013 , 45, 670-5	36.3	267
145	Genome-wide association study of dermatomyositis reveals genetic overlap with other autoimmune disorders. <i>Arthritis and Rheumatism</i> , 2013 , 65, 3239-47		94
144	Genetic and environmental determinants for disease risk in subsets of rheumatoid arthritis defined by the anticitrullinated protein/peptide antibody fine specificity profile. <i>Annals of the Rheumatic Diseases</i> , 2013 , 72, 652-8	2.4	127
143	Human genetics in rheumatoid arthritis guides a high-throughput drug screen of the CD40 signaling pathway. <i>PLoS Genetics</i> , 2013 , 9, e1003487	6	45
142	Variants in RUNX3 contribute to susceptibility to psoriatic arthritis, exhibiting further common ground with ankylosing spondylitis. <i>Arthritis and Rheumatism</i> , 2013 , 65, 1224-31		56
141	Genome-wide association study and gene expression analysis identifies CD84 as a predictor of response to etanercept therapy in rheumatoid arthritis. <i>PLoS Genetics</i> , 2013 , 9, e1003394	6	127
140	Implication of IL-2/IL-21 region in systemic sclerosis genetic susceptibility. <i>Annals of the Rheumatic Diseases</i> , 2013 , 72, 1233-8	2.4	28
139	HLA-DRB1*04/*13 alleles are associated with vascular disease and antiphospholipid antibodies in systemic lupus erythematosus. <i>Annals of the Rheumatic Diseases</i> , 2013 , 72, 1018-25	2.4	40
138	Crowdsourcing genetic prediction of clinical utility in the Rheumatoid Arthritis Responder Challenge. <i>Nature Genetics</i> , 2013 , 45, 468-9	36.3	20
137	Genetic association with ERAP1 in psoriasis is confined to disease onset after puberty and not dependent on HLA-C*06. <i>Journal of Investigative Dermatology</i> , 2013 , 133, 411-7	4.3	40
136	Contribution of IKBKE and IFIH1 gene variants to SLE susceptibility. <i>Genes and Immunity</i> , 2013 , 14, 217-22	4.4	22
135	Genes identified in Asian SLE GWASs are also associated with SLE in Caucasian populations. <i>European Journal of Human Genetics</i> , 2013 , 21, 994-9	5.3	62
134	OP0208 High density fine mapping in rheumatoid arthritis identifies 14 new loci. <i>Annals of the Rheumatic Diseases</i> , 2013 , 71, 126.1-126	2.4	
133	OP0052 A Dense Mapping of HLA Region for Study of Interaction with Smoking in the Development of Rheumatoid Arthritis. <i>Annals of the Rheumatic Diseases</i> , 2013 , 72, A67.2-A67	2.4	
132	AB0233 Type I interferon and clinical phenotype in idiopathic inflammatory myopathies. <i>Annals of the Rheumatic Diseases</i> , 2013 , 71, 650.16-650	2.4	
131	SAT0031 Anti-RO52 autoantibody epitope mapping in european cohort of myositis patients. <i>Annals of the Rheumatic Diseases</i> , 2013 , 71, 481.2-481	2.4	

130	THU0514 Thyroxin Substitution and the Risk of Developing Rheumatoid Arthritis; Results from the Swedish Population-Based Eira Study. <i>Annals of the Rheumatic Diseases</i> , 2013 , 72, A337.3-A338	2.4	
129	A research study of the association between maternal microchimerism and systemic lupus erythematosus in adults: a comparison between patients and healthy controls based on single-nucleotide polymorphism using quantitative real-time PCR. <i>PLoS ONE</i> , 2013 , 8, e74534	3.7	9
128	High-density genetic mapping identifies new susceptibility loci for rheumatoid arthritis. <i>Nature Genetics</i> , 2012 , 44, 1336-40	36.3	436
127	Age-dependent variation of genotypes in MHC II transactivator gene (CIITA) in controls and association to type 1 diabetes. <i>Genes and Immunity</i> , 2012 , 13, 632-40	4.4	19
126	Meta-analysis identifies nine new loci associated with rheumatoid arthritis in the Japanese population. <i>Nature Genetics</i> , 2012 , 44, 511-6	36.3	232
125	Polymorphisms in Toll-like receptor 3 confer natural resistance to human herpes simplex virus type 2 infection. <i>Journal of General Virology</i> , 2012 , 93, 1717-1724	4.9	35
124	The balance of expression of PTPN22 splice forms is significantly different in rheumatoid arthritis patients compared with controls. <i>Genome Medicine</i> , 2012 , 4, 2	14.4	20
123	Polymorphisms in peptidylarginine deiminase associate with rheumatoid arthritis in diverse Asian populations: evidence from MyEIRA study and meta-analysis. <i>Arthritis Research and Therapy</i> , 2012 , 14, R250	5.7	35
122	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). <i>Arthritis Research and Therapy</i> , 2012 , 14, R89	5.7	49
121	A replication study confirms the association of dendritic cell immunoreceptor (DCIR) polymorphisms with ACPA - negative RA in a large Asian cohort. <i>PLoS ONE</i> , 2012 , 7, e41228	3.7	20
120	Five amino acids in three HLA proteins explain most of the association between MHC and seropositive rheumatoid arthritis. <i>Nature Genetics</i> , 2012 , 44, 291-6	36.3	607
119	Variants within STAT genes reveal association with anticitrullinated protein antibody-negative rheumatoid arthritis in 2 European populations. <i>Journal of Rheumatology</i> , 2012 , 39, 1509-16	4.1	21
118	CIITA gene variants are associated with rheumatoid arthritis in Scandinavian populations. <i>Genes and Immunity</i> , 2012 , 13, 431-6	4.4	18
117	Very high levels of anti-citrullinated protein antibodies are associated with HLA-DRB1*15 non-shared epitope allele in patients with rheumatoid arthritis. <i>Arthritis and Rheumatism</i> , 2012 , 64, 2078-84		31
116	Use of a multiethnic approach to identify rheumatoid- arthritis-susceptibility loci, 1p36 and 17q12. <i>American Journal of Human Genetics</i> , 2012 , 90, 524-32	11	60
115	Novel genetic association of the VTCN1 region with rheumatoid arthritis. <i>Annals of the Rheumatic Diseases</i> , 2012 , 71, 567-71	2.4	11
114	High-density SNP mapping of the HLA region identifies multiple independent susceptibility loci associated with selective IgA deficiency. <i>PLoS Genetics</i> , 2012 , 8, e1002476	6	46
113	STAT4 regulates antiviral gamma interferon responses and recurrent disease during herpes simplex virus 2 infection. <i>Journal of Virology</i> , 2012 , 86, 9409-15	6.6	11

112	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. <i>Annals of the Rheumatic Diseases</i> , 2012 , 71, 961-5	2.4	75
111	The heterogeneity of asthma phenotypes in children and young adults. <i>Journal of Allergy</i> , 2012 , 2012, 163089		10
110	Influence of FCGR3A genotype on the therapeutic response to rituximab in rheumatoid arthritis: an observational cohort study. <i>BMJ Open</i> , 2012 , 2,	3	26
109	A novel sarcoidosis risk locus for Europeans on chromosome 11q13.1. <i>American Journal of Respiratory and Critical Care Medicine</i> , 2012 , 186, 877-85	10.2	34
108	Anti-Ro52 epitope mapping in inflammatory myopathies. <i>Annals of the Rheumatic Diseases</i> , 2012 , 71, A50.1-A50	2.4	
107	Smoking interacts with HLA-DRB1 shared epitope in the development of ACPA-positive rheumatoid arthritis: a case-control study from Malaysian epidemiological investigation of rheumatoid arthritis (MyEIRA). <i>Annals of the Rheumatic Diseases</i> , 2012 , 71, A57.1-A57	2.4	
106	Genetic variation in the serotonin receptor gene affects immune responses. <i>Annals of the Rheumatic Diseases</i> , 2012 , 71, A93-A93	2.4	
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