Jane Worthington

List of Publications by Year
in descending order

272	38,568	9234 74	2940 189
papers	citations	h-index	g-index

280	280	280	37839
all docs	docs citations	times ranked	citing authors

#	Article	IF	Citations
1	Genome-wide association study of 14,000 cases of seven common diseases and 3,000 shared controls. Nature, 2007, 447, 661-678.	13.7	8,895
2	Replication of Genome-Wide Association Signals in UK Samples Reveals Risk Loci for Type 2 Diabetes. Science, 2007, 316, 1336-1341.	6.0	2,040
3	Genetics of rheumatoid arthritis contributes to biology and drug discovery. Nature, 2014, 506, 376-381.	13.7	1,974
4	Association scan of 14,500 nonsynonymous SNPs in four diseases identifies autoimmunity variants. Nature Genetics, 2007, 39, 1329-1337.	9.4	1,298
5	Genome-wide association study meta-analysis identifies seven new rheumatoid arthritis risk loci. Nature Genetics, 2010, 42, 508-514.	9.4	1,132
6	Genome-wide association study identifies eight loci associated with blood pressure. Nature Genetics, 2009, 41, 666-676.	9.4	1,104
7	A genome-wide association study identifies new psoriasis susceptibility loci and an interaction between HLA-C and ERAP1. Nature Genetics, 2010, 42, 985-990.	9.4	918
8	Identification of 15 new psoriasis susceptibility loci highlights the role of innate immunity. Nature Genetics, 2012, 44, 1341-1348.	9.4	848
9	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
10	Genome-wide association study of CNVs in 16,000 cases of eight common diseases and 3,000 shared controls. Nature, 2010, 464, 713-720.	13.7	737
11	Meta-analysis and imputation refines the association of 15q25 with smoking quantity. Nature Genetics, 2010, 42, 436-440.	9.4	581
12	A Genome-Wide Association Study of Psoriasis and Psoriatic Arthritis Identifies New Disease Loci. PLoS Genetics, 2008, 4, e1000041.	1.5	572
13	High-density genetic mapping identifies new susceptibility loci for rheumatoid arthritis. Nature Genetics, 2012, 44, 1336-1340.	9.4	558
14	Pervasive Sharing of Genetic Effects in Autoimmune Disease. PLoS Genetics, 2011, 7, e1002254.	1.5	540
15	Localization of type 1 diabetes susceptibility to the MHC class I genes HLA-B and HLA-A. Nature, 2007, 450, 887-892.	13.7	493
16	Common variants at CD40 and other loci confer risk of rheumatoid arthritis. Nature Genetics, 2008, 40, 1216-1223.	9.4	476
17	Bayesian refinement of association signals for 14 loci in 3 common diseases. Nature Genetics, 2012, 44, 1294-1301.	9.4	469
18	Bayesian inference analyses of the polygenic architecture of rheumatoid arthritis. Nature Genetics, 2012, 44, 483-489.	9.4	402

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19	Rheumatoid arthritis association at 6q23. Nature Genetics, 2007, 39, 1431-1433.	9.4	361
20	EULAR recommendations for terminology and research in individuals at risk of rheumatoid arthritis: report from the Study Group for Risk Factors for Rheumatoid Arthritis. Annals of the Rheumatic Diseases, 2012, 71, 638-641.	0.5	354
21	Genome-wide association study of systemic sclerosis identifies CD247 as a new susceptibility locus. Nature Genetics, 2010, 42, 426-429.	9.4	351
22	Meta-Analysis of Genome-Wide Association Studies in Celiac Disease and Rheumatoid Arthritis Identifies Fourteen Non-HLA Shared Loci. PLoS Genetics, 2011, 7, e1002004.	1.5	307
23	Genetic variants at CD28, PRDM1 and CD2/CD58 are associated with rheumatoid arthritis risk. Nature Genetics, 2009, 41, 1313-1318.	9.4	306
24	Meta-analysis identifies nine new loci associated with rheumatoid arthritis in the Japanese population. Nature Genetics, 2012, 44, 511-516.	9.4	285
25	Association between thePTPN22 gene and rheumatoid arthritis and juvenile idiopathic arthritis in a UK population: Further support thatPTPN22 is an autoimmunity gene. Arthritis and Rheumatism, 2005, 52, 1694-1699.	6.7	266
26	Association of rheumatoid factor and anti-cyclic citrullinated peptide positivity, but not carriage of shared epitope or PTPN22 susceptibility variants, with anti-tumour necrosis factor response in rheumatoid arthritis. Annals of the Rheumatic Diseases, 2009, 68, 69-74.	0.5	240
27	CD226 Gly307Ser association with multiple autoimmune diseases. Genes and Immunity, 2009, 10, 5-10.	2.2	227
28	A Role for Noncoding Variation in Schizophrenia. Cell Reports, 2014, 9, 1417-1429.	2.9	225
29	Whole-Genome Scan, in a Complex Disease, Using 11,245 Single-Nucleotide Polymorphisms: Comparison with Microsatellites. American Journal of Human Genetics, 2004, 75, 54-64.	2.6	209
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	Quantitative heritability of antiâ€“citrullinated protein antibodyâ€“positive and antiâ€“citrullinated protein antibodyâ€“negative rheumatoid arthritis. Arthritis and Rheumatism, 2009, 60, 916-923.	6.7	200
32	A functional haplotype of thePADI4 gene associated with rheumatoid arthritis in a Japanese population is not associated in a United Kingdom population. Arthritis and Rheumatism, 2004, 50, 1117-1121.	6.7	186
33	Whole-genome linkage analysis of rheumatoid arthritis susceptibility loci in 252 affected sibling pairs in the United Kingdom. Arthritis and Rheumatism, 2002, 46, 632-639.	6.7	184
34	Immunochip Analysis Identifies Multiple Susceptibility Loci for Systemic Sclerosis. American Journal of Human Genetics, 2014, 94, 47-61.	2.6	182
35	Identification of ZNF313 / RNF114 as a novel psoriasis susceptibility gene. Human Molecular Genetics, 2008, 17, 1938-1945.	1.4	176
36	Widespread non-additive and interaction effects within HLA loci modulate the risk of autoimmune diseases. Nature Genetics, 2015, 47, 1085-1090.	9.4	164

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37	Capture Hi-C reveals novel candidate genes and complex long-range interactions with related autoimmune risk loci. Nature Communications, 2015, 6, 10069.	5.8	161
38	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
39	A Large-Scale Genetic Analysis Reveals a Strong Contribution of the HLA Class II Region to Giant Cell Arteritis Susceptibility. American Journal of Human Genetics, 2015, 96, 565-580.	2.6	144
40	Rheumatoid arthritis susceptibility loci at chromosomes 10p15, 12q13 and 22q13. Nature Genetics, 2008, 40, 1156-1159.	9.4	143
41	A family-based and case-control association study of the dopamine D4 receptor gene and dopamine transporter gene in attention deficit hyperactivity disorder. Molecular Psychiatry, 2000, 5, 523-530.	4.1	141
42	Psoriasis is associated with pleiotropic susceptibility loci identified in type II diabetes and Crohn disease. Journal of Medical Genetics, 2007, 45, 114-116.	1.5	139
43	Re-evaluation of putative rheumatoid arthritis susceptibility genes in the post-genome wide association study era and hypothesis of a key pathway underlying susceptibility. Human Molecular Genetics, 2008, 17, 2274-2279.	1.4	131
44	Study of the common genetic background for rheumatoid arthritis and systemic lupus erythematosus. Annals of the Rheumatic Diseases, 2011, 70, 463-468.	0.5	130
45	Statistical colocalization of genetic risk variants for related autoimmune diseases in the context of common controls. Nature Genetics, 2015, 47, 839-846.	9.4	128
46	Investigation of association of the IL12B and IL23R genes with psoriatic arthritis. Arthritis and Rheumatism, 2008, 58, 3705-3709.	6.7	122
47	Fine-mapping and functional studies highlight potential causal variants for rheumatoid arthritis and type 1 diabetes. Nature Genetics, 2018, 50, 1366-1374.	9.4	122
48	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
49	Association of HLA-DRB1 Haplotypes With Rheumatoid Arthritis Severity, Mortality, and Treatment Response. JAMA - Journal of the American Medical Association, 2015, 313, 1645.	3.8	119
50	Association between rheumatoid arthritis and polymorphism of tumor necrosis factor receptor II, but not tumor necrosis factor receptor I, in Caucasians. Arthritis and Rheumatism, 2001, 44, 61-65.	6.7	118
51	Quantifying Missing Heritability at Known GWAS Loci. PLoS Genetics, 2013, 9, e1003993.	1.5	115
52	Examining for association between candidate gene polymorphisms in the dopamine pathway and attention-deficit hyperactivity disorder: A family-based study. American Journal of Medical Genetics Part A, 2001, 105, 464-470.	2.4	112
53	The genetics of rheumatoid arthritis: risk and protection in different stages of the evolution of RA: Table 1. Rheumatology, 2016, 55, 199-209.	0.9	112
54	Association of the tumour necrosis factor-308 variant with differential response to anti-TNF agents in the treatment of rheumatoid arthritis. Human Molecular Genetics, 2008, 17, 3532-3538.	1.4	111

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55	Association of DRD4 in children with ADHD and comorbid conduct problems. American Journal of Medical Genetics Part A, 2002, 114, 150-153.	2.4	109
56	Genetic Variation in Efflux Transporters Influences Outcome to Methotrexate Therapy in Patients with Psoriasis. Journal of Investigative Dermatology, 2008, 128, 1925-1929.	0.3	109
57	A systemic sclerosis and systemic lupus erythematosus pan-meta-GWAS reveals new shared susceptibility loci. Human Molecular Genetics, 2013, 22, 4021-4029.	1.4	104
58	Whole-genome screening for susceptibility genes in multicase families with BehÃ§et's disease. Arthritis and Rheumatism, 2005, 52, 1836-1842.	6.7	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	Identification of CSK as a systemic sclerosis genetic risk factor through Genome Wide Association Study follow-up. Human Molecular Genetics, 2012, 21, 2825-2835.	1.4	98
62	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
63	Two novel biallelic polymorphisms in the ILâ€2 gene. International Journal of Immunogenetics, 1998, 25, 419-420.	1.2	94
64	The distribution of the endogenous retroviruses HERV-K113 and HERV-K115 in health and disease. Genomics, 2005, 86, 337-341.	1.3	94
65	Combined effects of three independent SNPs greatly increase the risk estimate for RA at 6q23. Human Molecular Genetics, 2009, 18, 2693-2699.	1.4	93
66	Association of the IL2RA/CD25 gene with juvenile idiopathic arthritis. Arthritis and Rheumatism, 2009, 60, 251-257.	6.7	93
67	Genetic markers of rheumatoid arthritis susceptibility in anti-citrullinated peptide antibody negative patients. Annals of the Rheumatic Diseases, 2012, 71, 1984-1990.	0.5	93
68	Investigating the role of the HLA-Cw*06 and HLA-DRB1 genes in susceptibility to psoriatic arthritis: comparison with psoriasis and undifferentiated inflammatory arthritis. Annals of the Rheumatic Diseases, 2007, 67, 677-682.	0.5	92
69	Overlapping genetic susceptibility variants between three autoimmune disorders: rheumatoid arthritis, type 1 diabetes and coeliac disease. Arthritis Research and Therapy, 2010, 12, R175.	1.6	92
70	Evidence for linkage of the HLA-B locus in Behï¿½et's disease, obtained using the transmission disequilibrium test. Arthritis and Rheumatism, 2001, 44, 239-241.	6.7	86
71	Reevaluation of the interaction between HLAâ€“DRB1 shared epitope alleles, PTPN22, and smoking in determining susceptibility to autoantibodyâ€positive and autoantibodyâ€negative rheumatoid arthritis in a large UK Caucasian population. Arthritis and Rheumatism, 2009, 60, 2565-2576.	6.7	86
72	Capture Hi-C identifies a novel causal gene, IL20RA, in the pan-autoimmune genetic susceptibility region 6q23. Genome Biology, 2016, 17, 212.	3.8	85

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73	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
74	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
75	Genetic susceptibility to rheumatoid arthritis: An emerging picture. Arthritis and Rheumatism, 2009, 61, 1441-1446.	6.7	79
76	Identification of AF4/FMR2 family, member 3 (AFF3) as a novel rheumatoid arthritis susceptibility locus and confirmation of two further pan-autoimmune susceptibility genes. Human Molecular Genetics, 2009, 18, 2518-2522.	1.4	78
77	Informed Conditioning on Clinical Covariates Increases Power in Case-Control Association Studies. PLoS Genetics, 2012, 8, e1003032.	1.5	78
78	A rare polymorphism in the gene for Tollâ€like receptor 2 is associated with systemic sclerosis phenotype and increases the production of inflammatory mediators. Arthritis and Rheumatism, 2012, 64, 264-271.	6.7	77
79	Association of tumor necrosis factor microsatellite polymorphisms with HLA-DRB1*04â€“bearing haplotypes in rheumatoid arthritis patients. Arthritis and Rheumatism, 1996, 39, 1109-1114.	6.7	76
80	Macrophage migration inhibitory factor (MIF) gene polymorphism is associated with susceptibility to but not severity of inflammatory polyarthritis. Genes and Immunity, 2003, 4, 487-491.	2.2	76
81	Investigation of potential non-HLA rheumatoid arthritis susceptibility loci in a European cohort increases the evidence for nine markers. Annals of the Rheumatic Diseases, 2010, 69, 1548-1553.	0.5	75
82	Analysis of the influence of PTPN22 gene polymorphisms in systemic sclerosis. Annals of the Rheumatic Diseases, 2011, 70, 454-462.	0.5	75
83	Characterization of a prolactin gene polymorphism and its associations with systemic lupus erythematosus. Arthritis and Rheumatism, 2001, 44, 2358-2366.	6.7	74
84	Macrophage Migration Inhibitory Factor Gene Polymorphism is Associated with Psoriasis. Journal of Investigative Dermatology, 2004, 123, 484-487.	0.3	74
85	Polymorphisms in the IL-12Î² and IL-23R Genes Are Associated with Psoriasis of Early Onset in a UK Cohort. Journal of Investigative Dermatology, 2008, 128, 1325-1327.	0.3	74
86	A GWAS follow-up study reveals the association of the IL12RB2 gene with systemic sclerosis in Caucasian populations. Human Molecular Genetics, 2012, 21, 926-933.	1.4	74
87	PADI4 genotype is not associated with rheumatoid arthritis in a large UK Caucasian population. Annals of the Rheumatic Diseases, 2010, 69, 666-670.	0.5	73
88	Identification of a novel susceptibility locus for juvenile idiopathic arthritis by genome-wide association analysis. Arthritis and Rheumatism, 2009, 60, 258-263.	6.7	72
89	Increased DNA methylation variability in rheumatoid arthritis-discordant monozygotic twins. Genome Medicine, 2018, 10, 64.	3.6	71
90	Combined Sib-TDT and TDT Provide Evidence for Linkage of the Interleukin-1 Gene Cluster to Erosive Rheumatoid Arthritis. Human Molecular Genetics, 1999, 8, 1707-1713.	1.4	69

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91	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
92	Evidence to support IL-13 as a risk locus for psoriatic arthritis but not psoriasis vulgaris. Annals of the Rheumatic Diseases, 2011, 70, 1016-1019.	0.5	68
93	Evidence for common genetic control in pathways of inflammation for Crohn's disease and psoriatic arthritis. Arthritis and Rheumatism, 2005, 52, 3596-3602.	6.7	65
94	HLA-Cw6 and HLA-DRB1*07 together are associated with less severe joint disease in psoriatic arthritis. Annals of the Rheumatic Diseases, 2007, 66, 807-811.	0.5	64
95	Outcomes of methotrexate therapy for psoriasis and relationship to genetic polymorphisms. British Journal of Dermatology, 2009, 160, 438-441.	1.4	64
96	Linkage mapping of a novel susceptibility locus for Behï¿½et's disease to chromosome 6p22-23. Arthritis and Rheumatism, 2001, 44, 2693-2696.	6.7	63
97	Use of gene expression profiling to identify a novel glucocorticoid sensitivity determining gene, BMPRII. FASEB Journal, 2007, 21, 402-414.	0.2	63
98	A replication study confirms the association of TNFSF4 (OX40L) polymorphisms with systemic sclerosis in a large European cohort. Annals of the Rheumatic Diseases, 2011, 70, 638-641.	0.5	63
99	Attention deficit hyperactivity disorder with reading disabilities: preliminary genetic findings on the involvement of the ADRA2A gene. Journal of Child Psychology and Psychiatry and Allied Disciplines, 2005, 46, 1081-1088.	3.1	62
100	Association of CD40 with rheumatoid arthritis confirmed in a large UK case-control study. Annals of the Rheumatic Diseases, 2010, 69, 813-816.	0.5	62
101	A method to decipher pleiotropy by detecting underlying heterogeneity driven by hidden subgroups applied to autoimmune and neuropsychiatric diseases. Nature Genetics, 2016, 48, 803-810.	9.4	62
102	Overlap of disease susceptibility loci for rheumatoid arthritis and juvenile idiopathic arthritis. Annals of the Rheumatic Diseases, 2010, 69, 1049-1053.	0.5	61
103	Subtype specific genetic associations for juvenile idiopathic arthritis: ERAP1 with the enthesitis related arthritis subtype and IL23R with juvenile psoriatic arthritis. Arthritis Research and Therapy, 2011, 13, R12.	1.6	60
104	Are both genetic and reproductive associations with rheumatoid arthritis linked to prolactin?. Lancet, The, 1996, 348, 106-109.	6.3	59
105	Protective effect of noninherited maternal HLA-DR antigens on rheumatoid arthritis development. Proceedings of the National Academy of Sciences of the United States of America, 2007, 104, 19966-19970.	3.3	59
106	Differential Methylation as a Biomarker of Response to Etanercept in Patients With Rheumatoid Arthritis. Arthritis and Rheumatology, 2016, 68, 1353-1360.	2.9	59
107	Allelic markers close to prolactin are associated with HLA-DRB1 susceptibility alleles among women with rheumatoid arthritis and systemic lupus erythematosus. Arthritis and Rheumatism, 1997, 40, 1383-1386.	6.7	58
108	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

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109	A two-stage, genome-wide screen for susceptibility loci in primary Raynaud's phenomenon. Arthritis and Rheumatism, 2000, 43, 1641-1646.	6.7	56
110	HLA-DRB1 and disease outcome in multiple sclerosis. Journal of Neurology, 2001, 248, 304-310.	1.8	56
111	Confirmation ofTNIP1but notRHOBandPSORS1C1as systemic sclerosis risk factors in a large independent replication study. Annals of the Rheumatic Diseases, 2013, 72, 602-607.	0.5	56
112	Predicting the Risk of Rheumatoid Arthritis and Its Age of Onset through Modelling Genetic Risk Variants with Smoking. PLoS Genetics, 2013, 9, e1003808.	1.5	55
113	A weighted genetic risk score using all known susceptibility variants to estimate rheumatoid arthritis risk. Annals of the Rheumatic Diseases, 2015, 74, 170-176.	0.5	55
114	Polymorphisms of the human prolactin geneâ€"implications for production of lymphocyte prolactin and systemic lupus erythematosus. Lupus, 2001, 10, 676-683.	0.8	54
115	Association of the AFF3 gene and IL2/IL21 gene region with juvenile idiopathic arthritis. Genes and Immunity, 2010, 11, 194-198.	2.2	54
116	Differential contribution of CDKAL1 variants to psoriasis, Crohn's disease and type II diabetes. Genes and Immunity, 2009, 10, 654-658.	2.2	53
117	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
118	High resolution linkage and association mapping identifies a novel rheumatoid arthritis susceptibility locus homologous to one linked to two rat models of inflammatory arthritis. Human Molecular Genetics, 2001, 10, 1901-1906.	1.4	52
119	Human Genetics in Rheumatoid Arthritis Guides a High-Throughput Drug Screen of the CD40 Signaling Pathway. PLoS Genetics, 2013, 9, e1003487.	1.5	52
120	New insight on the Xq28 association with systemic sclerosis. Annals of the Rheumatic Diseases, 2013, 72, 2032-2038.	0.5	52
121	A common biological basis of obesity and nicotine addiction. Translational Psychiatry, 2013, 3, e308-e308.	2.4	51
122	Haplotype analysis in simplex families and novel analytic approaches in a case-control cohort reveal no evidence of association of the CTLA-4 gene with rheumatoid arthritis. Arthritis and Rheumatism, 2004, 50, 748-752.	6.7	50
123	A spectrum of susceptibility to rheumatoid arthritis within HLA-DRB1: stratification by autoantibody status in a large UK population. Genes and Immunity, 2012, 13, 120-128.	2.2	50
124	Autosomal Dominant (Beukes) Premature Degenerative Osteoarthropathy of the Hip Joint Maps to an 11-cM Region on Chromosome 4q35. American Journal of Human Genetics, 1999, 64, 904-908.	2.6	49
125	Additional genetic susceptibility for rheumatoid arthritis telomeric of the DRB1 locus. Arthritis and Rheumatism, 2004, 50, 763-769.	6.7	48
126	Polymorphisms in the endothelial nitric oxide synthase gene are associated with Behcì§et's disease. Rheumatology, 2005, 44, 614-617.	0.9	47

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127	FcgammaRIIIA-158V and rheumatoid arthritis: a confirmation study. British Journal of Rheumatology, 2003, 42, 528-533.	2.5	46
128	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
129	Multipoint linkage analysis of a candidate gene locus in rheumatoid arthritis demonstrates significant evidence of linkage and association with the corticotropin-releasing hormone genomic region. Arthritis and Rheumatism, 2000, 43, 1673-1678.	6.7	45
130	No evidence of association of two 5HT transporter gene polymorphisms and attention deficit hyperactivity disorder. Psychiatric Genetics, 2003, 13, 107-110.	0.6	45
131	Fine mapping the TAGAP risk locus in rheumatoid arthritis. Genes and Immunity, 2011, 12, 314-318.	2.2	44
132	Preliminary evidence of an association of tumour necrosis factor microsatellites with increased risk of multiple basal cell carcinomas. British Journal of Dermatology, 2000, 142, 441-445.	1.4	43
133	Apolipoprotein E genotype does not predict decline in intelligence in healthy older adults. Neuroscience Letters, 2002, 324, 74-76.	1.0	43
134	Influence of serotonin transporter gene polymorphisms on cognitive decline and cognitive abilities in a nondemented elderly population. Molecular Psychiatry, 2005, 10, 1133-1139.	4.1	43
135	Brief Report: Identification of BACH2 and RAD51B as Rheumatoid Arthritis Susceptibility Loci in a Metaâ€Analysis of Genomeâ€Wide Data. Arthritis and Rheumatism, 2013, 65, 3058-3062.	6.7	43
136	Linkage of cytokine genes to rheumatoid arthritis. Evidence of genetic heterogeneity. Annals of the Rheumatic Diseases, 1998, 57, 361-365.	0.5	42
137	Cathepsin D exon 2 polymorphism associated with general intelligence in a healthy older population. Molecular Psychiatry, 2003, 8, 14-18.	4.1	42
138	Genetic epidemiology: systemic sclerosis. Arthritis Research, 2002, 4, 165.	2.0	41
139	Investigation of association between the TRAF family genes and RA susceptibility. Annals of the Rheumatic Diseases, 2007, 66, 1322-1326.	0.5	41
140	Polymorphisms in the PTPN22 region are associated with psoriasis of early onset. British Journal of Dermatology, 2008, 158, 962-968.	1.4	41
141	Novel Rheumatoid Arthritis Susceptibility Locus at 22q12 Identified in an Extended UK Genomeâ€Wide Association Study. Arthritis and Rheumatology, 2014, 66, 24-30.	2.9	41
142	Influence ofTYK2in systemic sclerosis susceptibility: a newlocusin the IL-12 pathway. Annals of the Rheumatic Diseases, 2016, 75, 1521-1526.	0.5	41
143	The shared epitope hypothesis in rheumatoid arthritis: Evaluation of alternative classification criteria in a large UK Caucasian cohort. Arthritis and Rheumatism, 2008, 58, 1275-1283.	6.7	40
144	Investigation of rheumatoid arthritis susceptibility loci in juvenile idiopathic arthritis confirms high degree of overlap. Annals of the Rheumatic Diseases, 2012, 71, 1117-1121.	0.5	40

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145	Complement C4B null allele status confers risk for systemic lupus erythematosus in a Spanish population. International Journal of Immunogenetics, 1998, 25, 317-320.	1.2	39
146	Linkage of rheumatoid arthritis to insulin-dependent diabetes mellitus loci: Evidence supporting a hypothesis for the existence of common autoimmune susceptibility loci. Arthritis and Rheumatism, 2000, 43, 2771-2775.	6.7	39
147	Investigation of susceptibility loci identified in the UK rheumatoid arthritis whole-genome scan in a further series of 217 UK affected sibling pairs. Arthritis and Rheumatism, 2004, 50, 729-735.	6.7	39
148	The Systemic Lupus Erythematosus IRF5 Risk Haplotype Is Associated with Systemic Sclerosis. PLoS ONE, 2013, 8, e54419.	1.1	38
149	Investigating the genetic basis of susceptibility to rheumatoid arthritis. Journal of Autoimmunity, 2005, 25, 16-20.	3.0	37
150	Evaluation of the rheumatoid arthritis susceptibility loci HLA-DRB1, PTPN22, OLIG3/TNFAIP3, STAT4 and TRAF1/C5 in an inception cohort. Arthritis Research and Therapy, 2010, 12, R57.	1.6	37
151	A genome-wide association study follow-up suggests a possible role for PPARG in systemic sclerosis susceptibility. Arthritis Research and Therapy, 2014, 16, R6.	1.6	37
152	Association of protein kinase C alpha (PRKCA) gene with multiple sclerosis in a UK population. Brain, 2004, 127, 1717-1722.	3.7	36
153	Association of the FCRL3 gene with rheumatoid arthritis: a further example of population specificity?. Arthritis Research and Therapy, 2006, 8, R117.	1.6	36
154	A large multicentre analysis of CTGF -945 promoter polymorphism does not confirm association with systemic sclerosis susceptibility or phenotype. Annals of the Rheumatic Diseases, 2009, 68, 1618-1620.	0.5	36
155	Polymorphisms spanning the ON exon and promoter of the estrogen receptorâ€beta (ERÎ²) gene ESR2 are associated with venous ulceration. Clinical Genetics, 2008, 73, 55-61.	1.0	35
156	Investigation of genetic variants within candidate genes of the TNFRSF1B signalling pathway on the response to anti-TNF agents in a UK cohort of rheumatoid arthritis patients. Pharmacogenetics and Genomics, 2009, 19, 319-323.	0.7	35
157	Investigation of type 1 diabetes and coeliac disease susceptibility loci for association with juvenile idiopathic arthritis. Annals of the Rheumatic Diseases, 2010, 69, 2169-2172.	0.5	34
158	Influence of the IL6 Gene in Susceptibility to Systemic Sclerosis. Journal of Rheumatology, 2012, 39, 2294-2302.	1.0	34
159	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
160	Investigation of theSLC22A4 gene (associated with rheumatoid arthritis in a Japanese population) in a United Kingdom population of rheumatoid arthritis patients. Arthritis and Rheumatism, 2005, 52, 752-758.	6.7	33
161	The dinucleotide (CA) repeat polymorphism of estrogen receptor beta but not the dinucleotide (TA) repeat polymorphism of estrogen receptor alpha is associated with venous ulceration. Journal of Steroid Biochemistry and Molecular Biology, 2005, 97, 266-270.	1.2	33
162	Replication of Associations of Genetic Loci Outside the HLA Region With Susceptibility to Antiâ€“Cyclic Citrullinated Peptideâ€“Negative Rheumatoid Arthritis. Arthritis and Rheumatology, 2016, 68, 1603-1613.	2.9	33

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163	Investigation of polymorphisms in the PADI4 gene in determining severity of inflammatory polyarthritis. Annals of the Rheumatic Diseases, 2005, 64, 1311-1315.	0.5	32
164	Data for Genetic Analysis Workshop (GAW) 15 Problem 2, genetic causes of rheumatoid arthritis and associated traits. BMC Proceedings, 2007, 1, S3.	1.8	32
165	Confirmation of association of the REL locus with rheumatoid arthritis susceptibility in the UK population. Annals of the Rheumatic Diseases, 2010, 69, 1572-1573.	0.5	32
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167	Susceptibility genes for a trait measure of attention deficit hyperactivity disorder: a pilot study in a non-clinical sample of twins. Psychiatry Research, 2001, 105, 273-278.	1.7	30
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