Stephen Eyre

List of Publications by Year
in descending order

140	26,285	34105 52	15732 125
papers	citations	h-index	g-index

144	144	144	31210
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.	27.8	8,895
2	Replication of Genome-Wide Association Signals in UK Samples Reveals Risk Loci for Type 2 Diabetes. Science, 2007, 316, 1336-1341.	12.6	2,040
3	Genetics of rheumatoid arthritis contributes to biology and drug discovery. Nature, 2014, 506, 376-381.	27.8	1,974
4	Association scan of 14,500 nonsynonymous SNPs in four diseases identifies autoimmunity variants. Nature Genetics, 2007, 39, 1329-1337.	21.4	1,298
5	Genome-wide association study meta-analysis identifies seven new rheumatoid arthritis risk loci. Nature Genetics, 2010, 42, 508-514.	21.4	1,132
6	Genome-wide association study identifies eight loci associated with blood pressure. Nature Genetics, 2009, 41, 666-676.	21.4	1,104
7	Genome-wide association study of CNVs in 16,000 cases of eight common diseases and 3,000 shared controls. Nature, 2010, 464, 713-720.	27.8	737
8	Meta-analysis and imputation refines the association of 15q25 with smoking quantity. Nature Genetics, 2010, 42, 436-440.	21.4	581
9	High-density genetic mapping identifies new susceptibility loci for rheumatoid arthritis. Nature Genetics, 2012, 44, 1336-1340.	21.4	558
10	Localization of type 1 diabetes susceptibility to the MHC class I genes HLA-B and HLA-A. Nature, 2007, 450, 887-892.	27.8	493
11	Bayesian refinement of association signals for 14 loci in 3 common diseases. Nature Genetics, 2012, 44, 1294-1301.	21.4	469
12	Rheumatoid arthritis association at 6q23. Nature Genetics, 2007, 39, 1431-1433.	21.4	361
13	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.	0.9	354
14	Dense genotyping of immune-related disease regions identifies 14 new susceptibility loci for juvenile idiopathic arthritis. Nature Genetics, 2013, 45, 664-669.	21.4	337
15	Genetic variants at CD28, PRDM1 and CD2/CD58 are associated with rheumatoid arthritis risk. Nature Genetics, 2009, 41, 1313-1318.	21.4	306
16	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.	6.2	209
17	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
18	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

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19	Widespread non-additive and interaction effects within HLA loci modulate the risk of autoimmune diseases. Nature Genetics, 2015, 47, 1085-1090.	21.4	164
20	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.	6.2	156
21	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.	6.2	144
22	Rheumatoid arthritis susceptibility loci at chromosomes 10p15, 12q13 and 22q13. Nature Genetics, 2008, 40, 1156-1159.	21.4	143
23	Genetics of rheumatoid arthritis: 2018 status. Annals of the Rheumatic Diseases, 2019, 78, 446-453.	0.9	141
24	Optimisation of methods for bacterial skin microbiome investigation: primer selection and comparison of the 454 versus MiSeq platform. BMC Microbiology, 2017, 17, 23.	3.3	133
25	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.	2.9	131
26	Study of the common genetic background for rheumatoid arthritis and systemic lupus erythematosus. Annals of the Rheumatic Diseases, 2011, 70, 463-468.	0.9	130
27	Statistical colocalization of genetic risk variants for related autoimmune diseases in the context of common controls. Nature Genetics, 2015, 47, 839-846.	21.4	128
28	Persistent inflammatory and non-inflammatory mechanisms in refractory rheumatoid arthritis. Nature Reviews Rheumatology, 2021, 17, 17-33.	8.0	118
29	Genetic Variation in Efflux Transporters Influences Outcome to Methotrexate Therapy in Patients with Psoriasis. Journal of Investigative Dermatology, 2008, 128, 1925-1929.	0.7	109
30	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.	0.9	100
31	Combined effects of three independent SNPs greatly increase the risk estimate for RA at 6q23. Human Molecular Genetics, 2009, 18, 2693-2699.	2.9	93
32	Association of the IL2RA/CD25 gene with juvenile idiopathic arthritis. Arthritis and Rheumatism, 2009, 60, 251-257.	6.7	93
33	Genetic markers of rheumatoid arthritis susceptibility in anti-citrullinated peptide antibody negative patients. Annals of the Rheumatic Diseases, 2012, 71, 1984-1990.	0.9	93
34	Overlapping genetic susceptibility variants between three autoimmune disorders: rheumatoid arthritis, type 1 diabetes and coeliac disease. Arthritis Research and Therapy, 2010, 12, R175.	3.5	92
35	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
36	Capture Hi-C identifies a novel causal gene, IL20RA, in the pan-autoimmune genetic susceptibility region 6q23. Genome Biology, 2016, 17, 212.	8.8	85

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37	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.	6.2	83
38	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.	2.9	78
39	Informed Conditioning on Clinical Covariates Increases Power in Case-Control Association Studies. PLoS Genetics, 2012, 8, e1003032.	3.5	78
40	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.	0.7	74
41	PADI4 genotype is not associated with rheumatoid arthritis in a large UK Caucasian population. Annals of the Rheumatic Diseases, 2010, 69, 666-670.	0.9	73
42	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
43	Risk-taking Behavior in Adolescents: The Paradigm. Annals of the New York Academy of Sciences, 1997, 817, 1-35.	3.8	69
44	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.	0.9	68
45	Outcomes of methotrexate therapy for psoriasis and relationship to genetic polymorphisms. British Journal of Dermatology, 2009, 160, 438-441.	1.5	64
46	Association of CD40 with rheumatoid arthritis confirmed in a large UK case-control study. Annals of the Rheumatic Diseases, 2010, 69, 813-816.	0.9	62
47	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.	21.4	62
48	Overlap of disease susceptibility loci for rheumatoid arthritis and juvenile idiopathic arthritis. Annals of the Rheumatic Diseases, 2010, 69, 1049-1053.	0.9	61
49	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.	3.5	60
50	Differential Methylation as a Biomarker of Response to Etanercept in Patients With Rheumatoid Arthritis. Arthritis and Rheumatology, 2016, 68, 1353-1360.	5.6	59
51	Genetic variants within the MAP kinase signalling network and anti-TNF treatment response in rheumatoid arthritis patients. Annals of the Rheumatic Diseases, 2011, 70, 98-103.	0.9	55
52	Predicting the Risk of Rheumatoid Arthritis and Its Age of Onset through Modelling Genetic Risk Variants with Smoking. PLoS Genetics, 2013, 9, e1003808.	3.5	55
53	A weighted genetic risk score using all known susceptibility variants to estimate rheumatoid arthritis risk. Annals of the Rheumatic Diseases, 2015, 74, 170-176.	0.9	55
54	Association of the AFF3 gene and IL2/IL21 gene region with juvenile idiopathic arthritis. Genes and Immunity, 2010, 11, 194-198.	4.1	54

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55	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.	2.9	52
56	Human Genetics in Rheumatoid Arthritis Guides a High-Throughput Drug Screen of the CD40 Signaling Pathway. PLoS Genetics, 2013, 9, e1003487.	3.5	52
57	Haplotype analysis in simplex families and novel analytic approaches in a caseâ€“control cohort reveal no evidence of association of the CTLAâ€4 gene with rheumatoid arthritis. Arthritis and Rheumatism, 2004, 50, 748-752.	6.7	50
58	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.	4.1	50
59	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.	6.2	49
60	One SNP at a Time: Moving beyond GWAS in Psoriasis. Journal of Investigative Dermatology, 2016, 136, 567-573.	0.7	48
61	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.	5.6	46
62	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
63	Investigation of association between the TRAF family genes and RA susceptibility. Annals of the Rheumatic Diseases, 2007, 66, 1322-1326.	0.9	41
64	Polymorphisms in the PTPN22 region are associated with psoriasis of early onset. British Journal of Dermatology, 2008, 158, 962-968.	1.5	41
65	Novel Rheumatoid Arthritis Susceptibility Locus at 22q12 Identified in an Extended UK Genomeâ€Wide Association Study. Arthritis and Rheumatology, 2014, 66, 24-30.	5.6	41
66	Investigation of rheumatoid arthritis susceptibility loci in juvenile idiopathic arthritis confirms high degree of overlap. Annals of the Rheumatic Diseases, 2012, 71, 1117-1121.	0.9	40
67	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
68	The potential use of expression profiling: implications for predicting treatment response in rheumatoid arthritis. Annals of the Rheumatic Diseases, 2013, 72, 1118-1124.	0.9	38
69	CD4+ and B Lymphocyte Expression Quantitative Traits at Rheumatoid Arthritis Risk Loci in Patients With Untreated Early Arthritis. Arthritis and Rheumatology, 2018, 70, 361-370.	5.6	37
70	Analysis of chromatin organization and gene expression in T cells identifies functional genes for rheumatoid arthritis. Nature Communications, 2020, 11, 4402.	12.8	37
71	Uncovering genetic mechanisms of hypertension through multi-omic analysis of the kidney. Nature Genetics, 2021, 53, 630-637.	21.4	37
72	Association of protein kinase C alpha (PRKCA) gene with multiple sclerosis in a UK population. Brain, 2004, 127, 1717-1722.	7.6	36

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73	Association of the FCRL3 gene with rheumatoid arthritis: a further example of population specificity?. Arthritis Research and Therapy, 2006, 8, R117.	3.5	36
74	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.	0.9	34
75	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
76	The bacterial skin microbiome in psoriatic arthritis, an unexplored link in pathogenesis: challenges and opportunities offered by recent technological advances. Rheumatology, 2014, 53, 777-784.	1.9	33
77	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.	5.6	33
78	Investigation of polymorphisms in the PADI4 gene in determining severity of inflammatory polyarthritis. Annals of the Rheumatic Diseases, 2005, 64, 1311-1315.	0.9	32
79	Confirmation of association of the REL locus with rheumatoid arthritis susceptibility in the UK population. Annals of the Rheumatic Diseases, 2010, 69, 1572-1573.	0.9	32
80	The role of rheumatoid arthritis genetic susceptibility markers in the prediction of erosive disease in patients with early inflammatory polyarthritis: results from the Norfolk Arthritis Register. Rheumatology, 2011, 50, 78-84.	1.9	32
81	Combined genetic analysis of juvenile idiopathic arthritis clinical subtypes identifies novel risk loci, target genes and key regulatory mechanisms. Annals of the Rheumatic Diseases, 2021, 80, 321-328.	0.9	31
82	Polymorphisms in the tumour necrosis factor gene are not associated with severity of inflammatory polyarthritis. Annals of the Rheumatic Diseases, 2004, 63, 280-284.	0.9	30
83	Investigation of genetic variation across the protein tyrosine phosphatase gene in patients with rheumatoid arthritis in the UK. Annals of the Rheumatic Diseases, 2007, 66, 683-686.	0.9	30
84	The genetics revolution in rheumatology: large scale genomic arrays and genetic mapping. Nature Reviews Rheumatology, 2017, 13, 421-432.	8.0	30
85	Rare variation at the TNFAIP3 locus and susceptibility to rheumatoid arthritis. Human Genetics, 2010, 128, 627-633.	3.8	29
86	A systematic investigation of confirmed autoimmune loci in early-onset psoriasis reveals an association with IL2/IL21. British Journal of Dermatology, 2011, 164, no-no.	1.5	28
87	Identifying Causal Genes at the Multiple Sclerosis Associated Region 6q23 Using Capture Hi-C. PLoS ONE, 2016, 11, e0166923.	2.5	28
88	Identification of the Tyrosine-Protein Phosphatase Non-Receptor Type 2 as a Rheumatoid Arthritis Susceptibility Locus in Europeans. PLoS ONE, 2013, 8, e66456.	2.5	27
89	Functional genomics atlas of synovial fibroblasts defining rheumatoid arthritis heritability. Genome Biology, 2021, 22, 247.	8.8	27
90	Association of the CCR5 gene with juvenile idiopathic arthritis. Genes and Immunity, 2010, 11, 584-589.	4.1	24

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91	Genetics of rheumatoid arthritis: GWAS and beyond. Open Access Rheumatology: Research and Reviews, 2011, 3, 31.	1.6	22
92	Investigation of theMHC2TA gene, associated with rheumatoid arthritis in a Swedish population, in a UK rheumatoid arthritis cohort. Arthritis and Rheumatism, 2006, 54, 3417-3422.	6.7	21
93	Genetic analysis of the Trichuris muris-induced model of colitis reveals QTL overlap and a novel gene cluster for establishing colonic inflammation. BMC Genomics, 2013, 14, 127.	2.8	20
94	Lymphocyte DNA methylation mediates genetic risk at shared immune-mediated disease loci. Journal of Allergy and Clinical Immunology, 2020, 145, 1438-1451.	2.9	20
95	Loci associated with N-glycosylation of human IgG are not associated with rheumatoid arthritis: a Mendelian randomisation study. Annals of the Rheumatic Diseases, 2016, 75, 317-320.	0.9	19
96	Evidence for a novel rheumatoid arthritis susceptibility locus on chromosome 6p. Arthritis and Rheumatism, 2004, 50, 3823-3830.	6.7	18
97	Correlation of C-reactive protein haplotypes with serum C-reactive protein level and response to anti-tumor necrosis factor therapy in UK rheumatoid arthritis patients: results from the Biologics in Rheumatoid Arthritis Genetics and Genomics Study Syndicate cohort. Arthritis Research and Therapy, 2012, 14, R214.	3.5	18
98	Enrichment of vitamin D response elements in RA-associated loci supports a role for vitamin D in the pathogenesis of RA. Genes and Immunity, 2013, 14, 325-329.	4.1	18
99	Polymorphisms of the equine major histocompatibility complex class II DRA locus. Tissue Antigens, 2004, 64, 173-179.	1.0	17
100	Major histocompatibility complex harbors widespread genotypic variability of non-additive risk of rheumatoid arthritis including epistasis. Scientific Reports, 2016, 6, 25014.	3.3	17
101	The type 1 diabetes susceptibility gene SUMO4 at IDDM5 is not associated with susceptibility to rheumatoid arthritis or juvenile idiopathic arthritis. Rheumatology, 2005, 44, 1390-1393.	1.9	16
102	Linkage of a marker in intron D of the estrogen synthase locus to rheumatoid arthritis. Arthritis and Rheumatism, 1999, 42, 1617-1620.	6.7	15
103	HLA-DPB1-COL11A2 and three additional xMHC loci are independently associated with RA in a UK cohort. Genes and Immunity, 2011, 12, 169-175.	4.1	15
104	A genetic marker at the OLIG3/TNFAIP3 locus associates with methotrexate continuation in early inflammatory polyarthritis: results from the Norfolk Arthritis Register. Pharmacogenomics Journal, 2012, 12, 128-133.	2.0	14
105	A re-evaluation of three putative functional single nucleotide polymorphisms in rheumatoid arthritis. Annals of the Rheumatic Diseases, 2009, 68, 1373-1375.	0.9	13
106	Association of a rheumatoid arthritis susceptibility variant at the CCL21 locus with premature mortality in inflammatory polyarthritis patients. Arthritis Care and Research, 2010, 62, 676-682.	3.4	13
107	The predictive value of serum S100A9 and response to etanercept is not confirmed in a large UK rheumatoid arthritis cohort. Rheumatology, 2017, 56, kew387.	1.9	10
108	Characterisation of the genomic architecture of human chromosome 17q and evaluation of different methods for haplotype block definition. BMC Genetics, 2005, 6, 21.	2.7	9

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109	No evidence for association of the KLF12 gene with rheumatoid arthritis in a large UK cohort. Annals of the Rheumatic Diseases, 2010, 69, 1407-1408.	0.9	9
110	Common genetic variants associated with disease from genomeâ€wide association studies are mutually exclusive in prostate cancer and rheumatoid arthritis. BJU International, 2013, 111, 1148-1155.	2.5	9
111	Genetics of RA susceptibility, what comes next?. RMD Open, 2015, 1, e000028-e000028.	3.8	9
112	Genetic susceptibility to rheumatoid arthritis and its implications for novel drug discovery. Expert Opinion on Drug Discovery, 2016, 11, 805-813.	5.0	9
113	Association with HLA-DRÎ²1 position 37 distinguishes juvenile dermatomyositis from adult-onset myositis. Human Molecular Genetics, 2022, 31, 2471-2481.	2.9	9
114	Two novel polymorphisms in the human transforming growth factor beta 2 gene. Genes and Immunity, 2001, 2, 295-296.	4.1	8
115	ASSIMILATOR: a new tool to inform selection of associated genetic variants for functional studies. Bioinformatics, 2011, 27, 144-146.	4.1	8
116	Investigation of an interleukin-6 receptor gene polymorphism (rs2228145) as a predictor of cardiovascular mortality in inflammatory polyarthritis: results from the Norfolk Arthritis Register: TableÂ1. Annals of the Rheumatic Diseases, 2014, 73, 787-788.	0.9	8
117	Immunochip Analyses of Epistasis in Rheumatoid Arthritis Confirm Multiple Interactions within MHC and Suggest Novel Non-MHC Epistatic Signals. Journal of Rheumatology, 2016, 43, 839-845.	2.0	8
118	Examining the overlap between genome-wide rare variant association signals and linkage peaks in rheumatoid arthritis. Arthritis and Rheumatism, 2011, 63, 1522-1526.	6.7	7
119	The skin microbiome in psoriatic arthritis: methodology development and pilot data. Lancet, The, 2015, 385, S27.	13.7	7
120	No Association between Polymorphisms in the Interleukin-15 Gene and Early-Onset Psoriasis in a UK Cohort Suggests Heterogeneity for this Susceptibility Locus Identified in Chinese Psoriasis Patients. Journal of Investigative Dermatology, 2008, 128, 2904-2905.	0.7	6
121	Investigating CD11c expression as a potential genomic biomarker of response to TNF inhibitor biologics in whole blood rheumatoid arthritis samples. Arthritis Research and Therapy, 2015, 17, 359.	3.5	6
122	Take Your PICS: Moving from GWAS to Immune Function. Immunity, 2014, 41, 883-885.	14.3	5
123	Monogenic disorders as mimics of juvenile idiopathic arthritis. Pediatric Rheumatology, 2022, 20, .	2.1	4
124	Exploring the overlap between rheumatoid arthritis susceptibility loci and long non-coding RNA annotations. PLoS ONE, 2020, 15, e0223939.	2.5	2
125	Association of the FCRL3 gene with rheumatoid arthritis: a further example of population specificity?. Arthritis Research and Therapy, 2008, 10, 405.	3.5	0
126	Combined effects of three independent SNPs greatly increase the risk estimate for RA at 6q23. Human Molecular Genetics, 2010, 19, 4544-4544.	2.9	0

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127	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, 2010, 19, 4543-4543.	2.9	0
128	Case Study on Rheumatoid Arthritis. , 2011, , 307-323.		0
129	The rheumatoid arthritis and juvenile idiopathic arthritis associated major (A) allele of rs2104286 is a loss of expression variant of IL2RA. Annals of the Rheumatic Diseases, 2011, 70, A6-A6.	0.9	0
130	257.â€ƒIncorporating Genotypic Variability Mapping Enhances Discovery of Risk Loci for Rheumatoid Arthritis. Rheumatology, 2015, , .	1.9	0
131	05.10â€…Comparison of cd4+ and b lymphocyte expression quantitative trait associations at ra risk loci in untreated early arthritis patients. , 2017, , .		0
132	Genetics of Rheumatic Diseases. , 2017, , 327-343.		0
133	SAT0055â€…JOINT SPECIFIC TNF RESPONSE OF SYNOVIAL FIBROBLASTS IN RHEUMATOID ARTHRITIS. , 2019, , .		0
134	O11â€ƒLymphocyte DNA methylation mediates genetic risk at RA risk loci that are shared with other immune mediated diseases. Rheumatology, 2020, 59, .	1.9	0
135	O01â€ƒGenetic risk factors associated with increased risk of uveitis in patients with juvenile idiopathic arthritis. Rheumatology, 2021, 60, .	1.9	0
136	No evidence that genetic predictors of susceptibility predict changes in core outcomes in JIA. Rheumatology, 2022, , .	1.9	0
137	Title is missing!. , 2020, 15, e0223939.		0
138	Title is missing!. , 2020, 15, e0223939.		0
139	Title is missing!. , 2020, 15, e0223939.		0
140	Title is missing!. , 2020, 15, e0223939.		0