

Stephen Eyre

List of Publications by Citations

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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.

133
papers

22,288
citations

49
h-index

144
g-index

144
ext. papers

24,960
ext. citations

10.7
avg, IF

7.36
L-index

#	Paper	IF	Citations
133	Genome-wide association study of 14,000 cases of seven common diseases and 3,000 shared controls. <i>Nature</i> , 2007 , 447, 661-78	50.4	7801
132	Replication of genome-wide association signals in UK samples reveals risk loci for type 2 diabetes. <i>Science</i> , 2007 , 316, 1336-41	33.3	1823
131	Genetics of rheumatoid arthritis contributes to biology and drug discovery. <i>Nature</i> , 2014 , 506, 376-81	50.4	1426
130	Association scan of 14,500 nonsynonymous SNPs in four diseases identifies autoimmunity variants. <i>Nature Genetics</i> , 2007 , 39, 1329-37	36.3	1130
129	Genome-wide association study identifies eight loci associated with blood pressure. <i>Nature Genetics</i> , 2009 , 41, 666-76	36.3	970
128	Genome-wide association study meta-analysis identifies seven new rheumatoid arthritis risk loci. <i>Nature Genetics</i> , 2010 , 42, 508-14	36.3	969
127	Genome-wide association study of CNVs in 16,000 cases of eight common diseases and 3,000 shared controls. <i>Nature</i> , 2010 , 464, 713-20	50.4	639
126	Meta-analysis and imputation refines the association of 15q25 with smoking quantity. <i>Nature Genetics</i> , 2010 , 42, 436-40	36.3	521
125	High-density genetic mapping identifies new susceptibility loci for rheumatoid arthritis. <i>Nature Genetics</i> , 2012 , 44, 1336-40	36.3	436
124	Localization of type 1 diabetes susceptibility to the MHC class I genes HLA-B and HLA-A. <i>Nature</i> , 2007 , 450, 887-92	50.4	421
123	Bayesian refinement of association signals for 14 loci in 3 common diseases. <i>Nature Genetics</i> , 2012 , 44, 1294-301	36.3	347
122	Rheumatoid arthritis association at 6q23. <i>Nature Genetics</i> , 2007 , 39, 1431-3	36.3	328
121	Genetic variants at CD28, PRDM1 and CD2/CD58 are associated with rheumatoid arthritis risk. <i>Nature Genetics</i> , 2009 , 41, 1313-8	36.3	272
120	EULAR recommendations for terminology and research in individuals at risk of rheumatoid arthritis: report from the Study Group for Risk Factors for Rheumatoid Arthritis. <i>Annals of the Rheumatic Diseases</i> , 2012 , 71, 638-41	2.4	263
119	Dense genotyping of immune-related disease regions identifies 14 new susceptibility loci for juvenile idiopathic arthritis. <i>Nature Genetics</i> , 2013 , 45, 664-9	36.3	256
118	Whole-genome scan, in a complex disease, using 11,245 single-nucleotide polymorphisms: comparison with microsatellites. <i>American Journal of Human Genetics</i> , 2004 , 75, 54-64	11	198
117	Whole-genome linkage analysis of rheumatoid arthritis susceptibility loci in 252 affected sibling pairs in the United Kingdom. <i>Arthritis and Rheumatism</i> , 2002 , 46, 632-9		171

116	A functional haplotype of the PADI4 gene associated with rheumatoid arthritis in a Japanese population is not associated in a United Kingdom population. <i>Arthritis and Rheumatism</i> , 2004 , 50, 1117-21		170
115	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
114	Rheumatoid arthritis susceptibility loci at chromosomes 10p15, 12q13 and 22q13. <i>Nature Genetics</i> , 2008 , 40, 1156-9	36.3	125
113	Re-evaluation of putative rheumatoid arthritis susceptibility genes in the post-genome wide association study era and hypothesis of a key pathway underlying susceptibility. <i>Human Molecular Genetics</i> , 2008 , 17, 2274-9	5.6	121
112	Widespread non-additive and interaction effects within HLA loci modulate the risk of autoimmune diseases. <i>Nature Genetics</i> , 2015 , 47, 1085-90	36.3	112
111	Study of the common genetic background for rheumatoid arthritis and systemic lupus erythematosus. <i>Annals of the Rheumatic Diseases</i> , 2011 , 70, 463-8	2.4	107
110	Statistical colocalization of genetic risk variants for related autoimmune diseases in the context of common controls. <i>Nature Genetics</i> , 2015 , 47, 839-46	36.3	97
109	A large-scale genetic analysis reveals a strong contribution of the HLA class II region to giant cell arteritis susceptibility. <i>American Journal of Human Genetics</i> , 2015 , 96, 565-80	11	96
108	Genetic variation in efflux transporters influences outcome to methotrexate therapy in patients with psoriasis. <i>Journal of Investigative Dermatology</i> , 2008 , 128, 1925-9	4.3	88
107	Genetics of rheumatoid arthritis: 2018 status. <i>Annals of the Rheumatic Diseases</i> , 2019 , 78, 446-453	2.4	80
106	Overlapping genetic susceptibility variants between three autoimmune disorders: rheumatoid arthritis, type 1 diabetes and coeliac disease. <i>Arthritis Research and Therapy</i> , 2010 , 12, R175	5.7	79
105	Reevaluation of the interaction between HLA-DRB1 shared epitope alleles, PTPN22, and smoking in determining susceptibility to autoantibody-positive and autoantibody-negative rheumatoid arthritis in a large UK Caucasian population. <i>Arthritis and Rheumatism</i> , 2009 , 60, 2565-76		79
104	Combined effects of three independent SNPs greatly increase the risk estimate for RA at 6q23. <i>Human Molecular Genetics</i> , 2010 , 19, 4544-4544	5.6	78
103	Identification of AF4/FMR2 family, member 3 (AFF3) as a novel rheumatoid arthritis susceptibility locus and confirmation of two further pan-autoimmune susceptibility genes. <i>Human Molecular Genetics</i> , 2010 , 19, 4543-4543	5.6	78
102	Genetic markers of rheumatoid arthritis susceptibility in anti-citrullinated peptide antibody negative patients. <i>Annals of the Rheumatic Diseases</i> , 2012 , 71, 1984-90	2.4	78
101	Association of the FCRL3 gene with rheumatoid arthritis: a further example of population specificity?. <i>Arthritis Research and Therapy</i> , 2008 , 10, 405	5.7	78
100	Combined effects of three independent SNPs greatly increase the risk estimate for RA at 6q23. <i>Human Molecular Genetics</i> , 2009 , 18, 2693-9	5.6	77
99	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

98	Association of the IL2RA/CD25 gene with juvenile idiopathic arthritis. <i>Arthritis and Rheumatism</i> , 2009 , 60, 251-7		73
97	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
96	Optimisation of methods for bacterial skin microbiome investigation: primer selection and comparison of the 454 versus MiSeq platform. <i>BMC Microbiology</i> , 2017 , 17, 23	4.5	71
95	Identification of AF4/FMR2 family, member 3 (AFF3) as a novel rheumatoid arthritis susceptibility locus and confirmation of two further pan-autoimmune susceptibility genes. <i>Human Molecular Genetics</i> , 2009 , 18, 2518-22	5.6	70
94	Polymorphisms in the IL-12beta and IL-23R genes are associated with psoriasis of early onset in a UK cohort. <i>Journal of Investigative Dermatology</i> , 2008 , 128, 1325-7	4.3	66
93	Capture Hi-C identifies a novel causal gene, IL20RA, in the pan-autoimmune genetic susceptibility region 6q23. <i>Genome Biology</i> , 2016 , 17, 212	18.3	62
92	Identification of a novel susceptibility locus for juvenile idiopathic arthritis by genome-wide association analysis. <i>Arthritis and Rheumatism</i> , 2009 , 60, 258-63		60
91	Evidence to support IL-13 as a risk locus for psoriatic arthritis but not psoriasis vulgaris. <i>Annals of the Rheumatic Diseases</i> , 2011 , 70, 1016-9	2.4	60
90	Informed conditioning on clinical covariates increases power in case-control association studies. <i>PLoS Genetics</i> , 2012 , 8, e1003032	6	58
89	PADI4 genotype is not associated with rheumatoid arthritis in a large UK Caucasian population. <i>Annals of the Rheumatic Diseases</i> , 2010 , 69, 666-70	2.4	57
88	Outcomes of methotrexate therapy for psoriasis and relationship to genetic polymorphisms. <i>British Journal of Dermatology</i> , 2009 , 160, 438-41	4	55
87	Risk-taking behavior in adolescents: the paradigm. <i>Annals of the New York Academy of Sciences</i> , 1997 , 817, 1-35	6.5	55
86	Overlap of disease susceptibility loci for rheumatoid arthritis and juvenile idiopathic arthritis. <i>Annals of the Rheumatic Diseases</i> , 2010 , 69, 1049-53	2.4	53
85	Subtype specific genetic associations for juvenile idiopathic arthritis: ERAP1 with the enthesitis related arthritis subtype and IL23R with juvenile psoriatic arthritis. <i>Arthritis Research and Therapy</i> , 2011 , 13, R12	5.7	52
84	Association of CD40 with rheumatoid arthritis confirmed in a large UK case-control study. <i>Annals of the Rheumatic Diseases</i> , 2010 , 69, 813-6	2.4	49
83	Association of the AFF3 gene and IL2/IL21 gene region with juvenile idiopathic arthritis. <i>Genes and Immunity</i> , 2010 , 11, 194-8	4.4	48
82	Genetic variants within the MAP kinase signalling network and anti-TNF treatment response in rheumatoid arthritis patients. <i>Annals of the Rheumatic Diseases</i> , 2011 , 70, 98-103	2.4	48
81	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. <i>Arthritis and Rheumatism</i> , 2004 , 50, 748-52		48

80	High resolution linkage and association mapping identifies a novel rheumatoid arthritis susceptibility locus homologous to one linked to two rat models of inflammatory arthritis. <i>Human Molecular Genetics</i> , 2001 , 10, 1901-6	5.6	47
79	A spectrum of susceptibility to rheumatoid arthritis within HLA-DRB1: stratification by autoantibody status in a large UK population. <i>Genes and Immunity</i> , 2012 , 13, 120-8	4.4	46
78	Differential Methylation as a Biomarker of Response to Etanercept in Patients With Rheumatoid Arthritis. <i>Arthritis and Rheumatology</i> , 2016 , 68, 1353-60	9.5	46
77	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
76	A method to decipher pleiotropy by detecting underlying heterogeneity driven by hidden subgroups applied to autoimmune and neuropsychiatric diseases. <i>Nature Genetics</i> , 2016 , 48, 803-10	36.3	45
75	Autosomal dominant (Beukes) premature degenerative osteoarthropathy of the hip joint maps to an 11-cM region on chromosome 4q35. <i>American Journal of Human Genetics</i> , 1999 , 64, 904-8	11	44
74	A weighted genetic risk score using all known susceptibility variants to estimate rheumatoid arthritis risk. <i>Annals of the Rheumatic Diseases</i> , 2015 , 74, 170-6	2.4	43
73	Predicting the risk of rheumatoid arthritis and its age of onset through modelling genetic risk variants with smoking. <i>PLoS Genetics</i> , 2013 , 9, e1003808	6	40
72	Investigation of susceptibility loci identified in the UK rheumatoid arthritis whole-genome scan in a further series of 217 UK affected sibling pairs. <i>Arthritis and Rheumatism</i> , 2004 , 50, 729-35		37
71	Novel rheumatoid arthritis susceptibility locus at 22q12 identified in an extended UK genome-wide association study. <i>Arthritis and Rheumatology</i> , 2014 , 66, 24-30	9.5	36
70	Investigation of rheumatoid arthritis susceptibility loci in juvenile idiopathic arthritis confirms high degree of overlap. <i>Annals of the Rheumatic Diseases</i> , 2012 , 71, 1117-21	2.4	36
69	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
68	Identification of BACH2 and RAD51B as rheumatoid arthritis susceptibility loci in a meta-analysis of genome-wide data. <i>Arthritis and Rheumatism</i> , 2013 , 65, 3058-62		35
67	Investigation of association between the TRAF family genes and RA susceptibility. <i>Annals of the Rheumatic Diseases</i> , 2007 , 66, 1322-6	2.4	35
66	Polymorphisms in the PTPN22 region are associated with psoriasis of early onset. <i>British Journal of Dermatology</i> , 2008 , 158, 962-8	4	33
65	The potential use of expression profiling: implications for predicting treatment response in rheumatoid arthritis. <i>Annals of the Rheumatic Diseases</i> , 2013 , 72, 1118-24	2.4	32
64	Investigation of the SLC22A4 gene (associated with rheumatoid arthritis in a Japanese population) in a United Kingdom population of rheumatoid arthritis patients. <i>Arthritis and Rheumatism</i> , 2005 , 52, 752-8		31
63	One SNP at a Time: Moving beyond GWAS in Psoriasis. <i>Journal of Investigative Dermatology</i> , 2016 , 136, 567-573	4.3	30

62	The bacterial skin microbiome in psoriatic arthritis, an unexplored link in pathogenesis: challenges and opportunities offered by recent technological advances. <i>Rheumatology</i> , 2014 , 53, 777-84	3.9	29
61	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. <i>Rheumatology</i> , 2011 , 50, 78-84	3.9	29
60	Investigation of type 1 diabetes and coeliac disease susceptibility loci for association with juvenile idiopathic arthritis. <i>Annals of the Rheumatic Diseases</i> , 2010 , 69, 2169-72	2.4	29
59	Persistent inflammatory and non-inflammatory mechanisms in refractory rheumatoid arthritis. <i>Nature Reviews Rheumatology</i> , 2021 , 17, 17-33	8.1	29
58	Association of protein kinase C alpha (PRKCA) gene with multiple sclerosis in a UK population. <i>Brain</i> , 2004 , 127, 1717-22	11.2	28
57	Investigation of polymorphisms in the PADI4 gene in determining severity of inflammatory polyarthritis. <i>Annals of the Rheumatic Diseases</i> , 2005 , 64, 1311-5	2.4	28
56	Rare variation at the TNFAIP3 locus and susceptibility to rheumatoid arthritis. <i>Human Genetics</i> , 2010 , 128, 627-33	6.3	27
55	Association of the FCRL3 gene with rheumatoid arthritis: a further example of population specificity?. <i>Arthritis Research and Therapy</i> , 2006 , 8, R117	5.7	27
54	Investigation of genetic variation across the protein tyrosine phosphatase gene in patients with rheumatoid arthritis in the UK. <i>Annals of the Rheumatic Diseases</i> , 2007 , 66, 683-6	2.4	26
53	CD4+ and B Lymphocyte Expression Quantitative Traits at Rheumatoid Arthritis Risk Loci in Patients With Untreated Early Arthritis: Implications for Causal Gene Identification. <i>Arthritis and Rheumatology</i> , 2018 , 70, 361-370	9.5	26
52	Identification of the tyrosine-protein phosphatase non-receptor type 2 as a rheumatoid arthritis susceptibility locus in europeans. <i>PLoS ONE</i> , 2013 , 8, e66456	3.7	24
51	Replication of Associations of Genetic Loci Outside the HLA Region With Susceptibility to Anti-Cyclic Citrullinated Peptide-Negative Rheumatoid Arthritis. <i>Arthritis and Rheumatology</i> , 2016 , 68, 1603-13	9.5	24
50	Confirmation of association of the REL locus with rheumatoid arthritis susceptibility in the UK population. <i>Annals of the Rheumatic Diseases</i> , 2010 , 69, 1572-3	2.4	23
49	Association of the CCR5 gene with juvenile idiopathic arthritis. <i>Genes and Immunity</i> , 2010 , 11, 584-9	4.4	22
48	Polymorphisms in the tumour necrosis factor gene are not associated with severity of inflammatory polyarthritis. <i>Annals of the Rheumatic Diseases</i> , 2004 , 63, 280-4	2.4	21
47	A systematic investigation of confirmed autoimmune loci in early-onset psoriasis reveals an association with IL2/IL21. <i>British Journal of Dermatology</i> , 2011 , 164, 660-4	4	20
46	Investigation of the MHC2TA gene, associated with rheumatoid arthritis in a Swedish population, in a UK rheumatoid arthritis cohort. <i>Arthritis and Rheumatism</i> , 2006 , 54, 3417-22		20
45	Identifying Causal Genes at the Multiple Sclerosis Associated Region 6q23 Using Capture Hi-C. <i>PLoS ONE</i> , 2016 , 11, e0166923	3.7	20

44	The genetics revolution in rheumatology: large scale genomic arrays and genetic mapping. <i>Nature Reviews Rheumatology</i> , 2017 , 13, 421-432	8.1	18
43	Genetic analysis of the <i>Trichuris muris</i> -induced model of colitis reveals QTL overlap and a novel gene cluster for establishing colonic inflammation. <i>BMC Genomics</i> , 2013 , 14, 127	4.5	17
42	Evidence for a novel rheumatoid arthritis susceptibility locus on chromosome 6p. <i>Arthritis and Rheumatism</i> , 2004 , 50, 3823-30		17
41	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. <i>Arthritis Research and Therapy</i> , 2012 , 14, R214	5.7	16
40	Polymorphisms of the equine major histocompatibility complex class II DRA locus. <i>Tissue Antigens</i> , 2004 , 64, 173-9		16
39	Genetics of rheumatoid arthritis: GWAS and beyond. <i>Open Access Rheumatology: Research and Reviews</i> , 2011 , 3, 31-46	2.4	15
38	Linkage of a marker in intron D of the estrogen synthase locus to rheumatoid arthritis. <i>Arthritis and Rheumatism</i> , 1999 , 42, 1617-20		15
37	Enrichment of vitamin D response elements in RA-associated loci supports a role for vitamin D in the pathogenesis of RA. <i>Genes and Immunity</i> , 2013 , 14, 325-9	4.4	13
36	HLA-DPB1-COL11A2 and three additional xMHC loci are independently associated with RA in a UK cohort. <i>Genes and Immunity</i> , 2011 , 12, 169-75	4.4	13
35	A re-evaluation of three putative functional single nucleotide polymorphisms in rheumatoid arthritis. <i>Annals of the Rheumatic Diseases</i> , 2009 , 68, 1373-5	2.4	13
34	The type 1 diabetes susceptibility gene SUMO4 at IDDM5 is not associated with susceptibility to rheumatoid arthritis or juvenile idiopathic arthritis. <i>Rheumatology</i> , 2005 , 44, 1390-3	3.9	13
33	A genetic marker at the OLIG3/TNFAIP3 locus associates with methotrexate continuation in early inflammatory polyarthritis: results from the Norfolk Arthritis Register. <i>Pharmacogenomics Journal</i> , 2012 , 12, 128-33	3.5	12
32	Analysis of chromatin organization and gene expression in T cells identifies functional genes for rheumatoid arthritis. <i>Nature Communications</i> , 2020 , 11, 4402	17.4	12
31	Major histocompatibility complex harbors widespread genotypic variability of non-additive risk of rheumatoid arthritis including epistasis. <i>Scientific Reports</i> , 2016 , 6, 25014	4.9	11
30	Lymphocyte DNA methylation mediates genetic risk at shared immune-mediated disease loci. <i>Journal of Allergy and Clinical Immunology</i> , 2020 , 145, 1438-1451	11.5	10
29	Loci associated with N-glycosylation of human IgG are not associated with rheumatoid arthritis: a Mendelian randomisation study. <i>Annals of the Rheumatic Diseases</i> , 2016 , 75, 317-20	2.4	10
28	Association of a rheumatoid arthritis susceptibility variant at the CCL21 locus with premature mortality in inflammatory polyarthritis patients. <i>Arthritis Care and Research</i> , 2010 , 62, 676-82	4.7	10
27	Combined genetic analysis of juvenile idiopathic arthritis clinical subtypes identifies novel risk loci, target genes and key regulatory mechanisms. <i>Annals of the Rheumatic Diseases</i> , 2020 ,	2.4	10

26	Genetics of RA susceptibility, what comes next?. <i>RMD Open</i> , 2015 , 1, e000028	5.9	9
25	Genetic susceptibility to rheumatoid arthritis and its implications for novel drug discovery. <i>Expert Opinion on Drug Discovery</i> , 2016 , 11, 805-13	6.2	8
24	Characterisation of the genomic architecture of human chromosome 17q and evaluation of different methods for haplotype block definition. <i>BMC Genetics</i> , 2005 , 6, 21	2.6	8
23	Two novel polymorphisms in the human transforming growth factor beta 2 gene. <i>Genes and Immunity</i> , 2001 , 2, 295-6	4.4	8
22	The predictive value of serum S100A9 and response to etanercept is not confirmed in a large UK rheumatoid arthritis cohort. <i>Rheumatology</i> , 2017 , 56, 1019-1024	3.9	7
21	Investigation of an interleukin-6 receptor gene polymorphism (rs2228145) as a predictor of cardiovascular mortality in inflammatory polyarthritis: results from the Norfolk Arthritis Register. <i>Annals of the Rheumatic Diseases</i> , 2014 , 73, 787-8	2.4	7
20	Common genetic variants associated with disease from genome-wide association studies are mutually exclusive in prostate cancer and rheumatoid arthritis. <i>BJU International</i> , 2013 , 111, 1148-55	5.6	7
19	Examining the overlap between genome-wide rare variant association signals and linkage peaks in rheumatoid arthritis. <i>Arthritis and Rheumatism</i> , 2011 , 63, 1522-6		7
18	ASSIMILATOR: a new tool to inform selection of associated genetic variants for functional studies. <i>Bioinformatics</i> , 2011 , 27, 144-6	7.2	7
17	The skin microbiome in psoriatic arthritis: methodology development and pilot data. <i>Lancet, The</i> , 2015 , 385 Suppl 1, S27	4.0	6
16	ImmunoChip Analyses of Epistasis in Rheumatoid Arthritis Confirm Multiple Interactions within MHC and Suggest Novel Non-MHC Epistatic Signals. <i>Journal of Rheumatology</i> , 2016 , 43, 839-45	4.1	6
15	Functional genomics atlas of synovial fibroblasts defining rheumatoid arthritis heritability. <i>Genome Biology</i> , 2021 , 22, 247	18.3	6
14	Investigating CD11c expression as a potential genomic biomarker of response to TNF inhibitor biologics in whole blood rheumatoid arthritis samples. <i>Arthritis Research and Therapy</i> , 2015 , 17, 359	5.7	5
13	Take your PICS: moving from GWAS to immune function. <i>Immunity</i> , 2014 , 41, 883-5	32.3	5
12	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. <i>Journal of Investigative Dermatology</i> , 2008 , 128, 2904-5	4.3	5
11	Uncovering genetic mechanisms of hypertension through multi-omic analysis of the kidney. <i>Nature Genetics</i> , 2021 , 53, 630-637	36.3	5
10	No evidence for association of the KLF12 gene with rheumatoid arthritis in a large UK cohort. <i>Annals of the Rheumatic Diseases</i> , 2010 , 69, 1407-8	2.4	4
9	Exploring the overlap between rheumatoid arthritis susceptibility loci and long non-coding RNA annotations. <i>PLoS ONE</i> , 2020 , 15, e0223939	3.7	1

- 8 Functional genomics atlas of synovial fibroblasts defining rheumatoid arthritis heritability 1
- 7 Genetics of Rheumatic Diseases **2017**, 327-343
- 6 Case Study on Rheumatoid Arthritis **2011**, 307-323
- 5 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 2.4
- 4 Exploring the overlap between rheumatoid arthritis susceptibility loci and long non-coding RNA annotations **2020**, 15, e0223939
- 3 Exploring the overlap between rheumatoid arthritis susceptibility loci and long non-coding RNA annotations **2020**, 15, e0223939
- 2 Exploring the overlap between rheumatoid arthritis susceptibility loci and long non-coding RNA annotations **2020**, 15, e0223939
- 1 Exploring the overlap between rheumatoid arthritis susceptibility loci and long non-coding RNA annotations **2020**, 15, e0223939