

Stephen Eyre

List of Publications by Year in descending order

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Version: 2024-02-01

140
papers

26,285
citations

34016

52
h-index

15683

125
g-index

144
all docs

144
docs citations

144
times ranked

31210
citing authors

#	ARTICLE	IF	CITATIONS
1	Association with HLA-DR ¹ position 37 distinguishes juvenile dermatomyositis from adult-onset myositis. <i>Human Molecular Genetics</i> , 2022, 31, 2471-2481.	1.4	9
2	No evidence that genetic predictors of susceptibility predict changes in core outcomes in JIA. <i>Rheumatology</i> , 2022, , .	0.9	0
3	Monogenic disorders as mimics of juvenile idiopathic arthritis. <i>Pediatric Rheumatology</i> , 2022, 20, .	0.9	4
4	Persistent inflammatory and non-inflammatory mechanisms in refractory rheumatoid arthritis. <i>Nature Reviews Rheumatology</i> , 2021, 17, 17-33.	3.5	118
5	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> , 2021, 80, 321-328.	0.5	31
6	Genetic risk factors associated with increased risk of uveitis in patients with juvenile idiopathic arthritis. <i>Rheumatology</i> , 2021, 60, .	0.9	0
7	Uncovering genetic mechanisms of hypertension through multi-omic analysis of the kidney. <i>Nature Genetics</i> , 2021, 53, 630-637.	9.4	37
8	Functional genomics atlas of synovial fibroblasts defining rheumatoid arthritis heritability. <i>Genome Biology</i> , 2021, 22, 247.	3.8	27
9	Lymphocyte DNA methylation mediates genetic risk at RA risk loci that are shared with other immune mediated diseases. <i>Rheumatology</i> , 2020, 59, .	0.9	0
10	Analysis of chromatin organization and gene expression in T cells identifies functional genes for rheumatoid arthritis. <i>Nature Communications</i> , 2020, 11, 4402.	5.8	37
11	Exploring the overlap between rheumatoid arthritis susceptibility loci and long non-coding RNA annotations. <i>PLoS ONE</i> , 2020, 15, e0223939.	1.1	2
12	Lymphocyte DNA methylation mediates genetic risk at shared immune-mediated disease loci. <i>Journal of Allergy and Clinical Immunology</i> , 2020, 145, 1438-1451.	1.5	20
13	Title is missing!. , 2020, 15, e0223939.		0
14	Title is missing!. , 2020, 15, e0223939.		0
15	Title is missing!. , 2020, 15, e0223939.		0
16	Title is missing!. , 2020, 15, e0223939.		0
17	SAT0055â€¦JOINT SPECIFIC TNF RESPONSE OF SYNOVIAL FIBROBLASTS IN RHEUMATOID ARTHRITIS. , 2019, , .		0
18	Genetics of rheumatoid arthritis: 2018 status. <i>Annals of the Rheumatic Diseases</i> , 2019, 78, 446-453.	0.5	141

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19	CD4+ and B Lymphocyte Expression Quantitative Traits at Rheumatoid Arthritis Risk Loci in Patients With Untreated Early Arthritis. <i>Arthritis and Rheumatology</i> , 2018, 70, 361-370.	2.9	37
20	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.	1.3	133
21	The genetics revolution in rheumatology: large scale genomic arrays and genetic mapping. <i>Nature Reviews Rheumatology</i> , 2017, 13, 421-432.	3.5	30
22	05.10â€¦Comparison of cd4+ and b lymphocyte expression quantitative trait associations at ra risk loci in untreated early arthritis patients. , 2017, , .		0
23	Genetics of Rheumatic Diseases. , 2017, , 327-343.		0
24	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, kew387.	0.9	10
25	Capture Hi-C identifies a novel causal gene, IL20RA, in the pan-autoimmune genetic susceptibility region 6q23. <i>Genome Biology</i> , 2016, 17, 212.	3.8	85
26	Major histocompatibility complex harbors widespread genotypic variability of non-additive risk of rheumatoid arthritis including epistasis. <i>Scientific Reports</i> , 2016, 6, 25014.	1.6	17
27	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, 803-810.	9.4	62
28	Differential Methylation as a Biomarker of Response to Etanercept in Patients With Rheumatoid Arthritis. <i>Arthritis and Rheumatology</i> , 2016, 68, 1353-1360.	2.9	59
29	Replication of Associations of Genetic Loci Outside the HLA Region With Susceptibility to Antiâ€“Cyclic Citrullinated Peptideâ€“Negative Rheumatoid Arthritis. <i>Arthritis and Rheumatology</i> , 2016, 68, 1603-1613.	2.9	33
30	Genetic susceptibility to rheumatoid arthritis and its implications for novel drug discovery. <i>Expert Opinion on Drug Discovery</i> , 2016, 11, 805-813.	2.5	9
31	Brief Report: <i>IRF4</i> Newly Identified as a Common Susceptibility Locus for Systemic Sclerosis and Rheumatoid Arthritis in a Crossâ€“Disease Metaâ€“Analysis of Genomeâ€“Wide Association Studies. <i>Arthritis and Rheumatology</i> , 2016, 68, 2338-2344.	2.9	46
32	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, 839-845.	1.0	8
33	One SNP at a Time: Moving beyond GWAS in Psoriasis. <i>Journal of Investigative Dermatology</i> , 2016, 136, 567-573.	0.3	48
34	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, 317-320.	0.5	19
35	Identifying Causal Genes at the Multiple Sclerosis Associated Region 6q23 Using Capture Hi-C. <i>PLoS ONE</i> , 2016, 11, e0166923.	1.1	28
36	257.â€“Incorporating Genotypic Variability Mapping Enhances Discovery of Risk Loci for Rheumatoid Arthritis. <i>Rheumatology</i> , 2015, , .	0.9	0

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37	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.	1.6	6
38	Statistical colocalization of genetic risk variants for related autoimmune diseases in the context of common controls. <i>Nature Genetics</i> , 2015, 47, 839-846.	9.4	128
39	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-e13.	0.5	100
40	The skin microbiome in psoriatic arthritis: methodology development and pilot data. <i>Lancet, The</i> , 2015, 385, S27.	6.3	7
41	Genetics of RA susceptibility, what comes next?. <i>RMD Open</i> , 2015, 1, e000028-e000028.	1.8	9
42	A weighted genetic risk score using all known susceptibility variants to estimate rheumatoid arthritis risk. <i>Annals of the Rheumatic Diseases</i> , 2015, 74, 170-176.	0.5	55
43	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, 565-580.	2.6	144
44	Widespread non-additive and interaction effects within HLA loci modulate the risk of autoimmune diseases. <i>Nature Genetics</i> , 2015, 47, 1085-1090.	9.4	164
45	Take Your PICS: Moving from GWAS to Immune Function. <i>Immunity</i> , 2014, 41, 883-885.	6.6	5
46	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.	2.9	41
47	Genetics of rheumatoid arthritis contributes to biology and drug discovery. <i>Nature</i> , 2014, 506, 376-381.	13.7	1,974
48	Investigation of an interleukin-6 receptor gene polymorphism (rs2228145) as a predictor of cardiovascular mortality in inflammatory polyarthritis: results from the Norfolk Arthritis Register: Table A1. <i>Annals of the Rheumatic Diseases</i> , 2014, 73, 787-788.	0.5	8
49	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, 522-532.	2.6	156
50	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, 777-784.	0.9	33
51	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.	1.2	20
52	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, 325-329.	2.2	18
53	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, 1148-1155.	1.3	9
54	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.	2.6	83

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55	Dense genotyping of immune-related disease regions identifies 14 new susceptibility loci for juvenile idiopathic arthritis. <i>Nature Genetics</i> , 2013, 45, 664-669.	9.4	337
56	Human Genetics in Rheumatoid Arthritis Guides a High-Throughput Drug Screen of the CD40 Signaling Pathway. <i>PLoS Genetics</i> , 2013, 9, e1003487.	1.5	52
57	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.	1.5	55
58	Brief Report: Identification of <i>BACH2</i> and <i>RAD51B</i> as Rheumatoid Arthritis Susceptibility Loci in a Meta-Analysis of Genome-Wide Data. <i>Arthritis and Rheumatism</i> , 2013, 65, 3058-3062.	6.7	43
59	The potential use of expression profiling: implications for predicting treatment response in rheumatoid arthritis. <i>Annals of the Rheumatic Diseases</i> , 2013, 72, 1118-1124.	0.5	38
60	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.	1.1	27
61	Genetic markers of rheumatoid arthritis susceptibility in anti-citrullinated peptide antibody negative patients. <i>Annals of the Rheumatic Diseases</i> , 2012, 71, 1984-1990.	0.5	93
62	Informed Conditioning on Clinical Covariates Increases Power in Case-Control Association Studies. <i>PLoS Genetics</i> , 2012, 8, e1003032.	1.5	78
63	Investigation of rheumatoid arthritis susceptibility loci in juvenile idiopathic arthritis confirms high degree of overlap. <i>Annals of the Rheumatic Diseases</i> , 2012, 71, 1117-1121.	0.5	40
64	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, 638-641.	0.5	354
65	Bayesian refinement of association signals for 14 loci in 3 common diseases. <i>Nature Genetics</i> , 2012, 44, 1294-1301.	9.4	469
66	A genetic marker at the <i>OLIG3/TNFAIP3</i> locus associates with methotrexate continuation in early inflammatory polyarthritis: results from the Norfolk Arthritis Register. <i>Pharmacogenomics Journal</i> , 2012, 12, 128-133.	0.9	14
67	High-density genetic mapping identifies new susceptibility loci for rheumatoid arthritis. <i>Nature Genetics</i> , 2012, 44, 1336-1340.	9.4	558
68	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.	1.6	18
69	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, 120-128.	2.2	50
70	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.	0.9	32
71	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.	1.6	60
72	Genetics of rheumatoid arthritis: GWAS and beyond. <i>Open Access Rheumatology: Research and Reviews</i> , 2011, 3, 31.	0.8	22

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73	Case Study on Rheumatoid Arthritis. , 2011, , 307-323.		0
74	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.	1.4	28
75	HLA-DPB1-COL11A2 and three additional xMHC loci are independently associated with RA in a UK cohort. Genes and Immunity, 2011, 12, 169-175.	2.2	15
76	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
77	Evidence to support <i>IL-13</i> as a risk locus for psoriatic arthritis but not psoriasis vulgaris. Annals of the Rheumatic Diseases, 2011, 70, 1016-1019.	0.5	68
78	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.	0.5	55
79	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.	0.5	0
80	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
81	ASSIMILATOR: a new tool to inform selection of associated genetic variants for functional studies. Bioinformatics, 2011, 27, 144-146.	1.8	8
82	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
83	Rare variation at the TNFAIP3 locus and susceptibility to rheumatoid arthritis. Human Genetics, 2010, 128, 627-633.	1.8	29
84	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.	1.5	13
85	Association of the AFF3 gene and IL2/IL21 gene region with juvenile idiopathic arthritis. Genes and Immunity, 2010, 11, 194-198.	2.2	54
86	Association of the CCR5 gene with juvenile idiopathic arthritis. Genes and Immunity, 2010, 11, 584-589.	2.2	24
87	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
88	Meta-analysis and imputation refines the association of 15q25 with smoking quantity. Nature Genetics, 2010, 42, 436-440.	9.4	581
89	Genome-wide association study meta-analysis identifies seven new rheumatoid arthritis risk loci. Nature Genetics, 2010, 42, 508-514.	9.4	1,132
90	Combined effects of three independent SNPs greatly increase the risk estimate for RA at 6q23. Human Molecular Genetics, 2010, 19, 4544-4544.	1.4	0

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91	PADI4 genotype is not associated with rheumatoid arthritis in a large UK Caucasian population. <i>Annals of the Rheumatic Diseases</i> , 2010, 69, 666-670.	0.5	73
92	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, 1407-1408.	0.5	9
93	Overlap of disease susceptibility loci for rheumatoid arthritis and juvenile idiopathic arthritis. <i>Annals of the Rheumatic Diseases</i> , 2010, 69, 1049-1053.	0.5	61
94	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.	1.4	0
95	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, 2169-2172.	0.5	34
96	Association of CD40 with rheumatoid arthritis confirmed in a large UK case-control study. <i>Annals of the Rheumatic Diseases</i> , 2010, 69, 813-816.	0.5	62
97	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.	1.6	92
98	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, 2518-2522.	1.4	78
99	Combined effects of three independent SNPs greatly increase the risk estimate for RA at 6q23. <i>Human Molecular Genetics</i> , 2009, 18, 2693-2699.	1.4	93
100	Identification of a novel susceptibility locus for juvenile idiopathic arthritis by genome-wide association analysis. <i>Arthritis and Rheumatism</i> , 2009, 60, 258-263.	6.7	72
101	Association of the IL2RA/CD25 gene with juvenile idiopathic arthritis. <i>Arthritis and Rheumatism</i> , 2009, 60, 251-257.	6.7	93
102	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, 2565-2576.	6.7	86
103	Outcomes of methotrexate therapy for psoriasis and relationship to genetic polymorphisms. <i>British Journal of Dermatology</i> , 2009, 160, 438-441.	1.4	64
104	Genome-wide association study identifies eight loci associated with blood pressure. <i>Nature Genetics</i> , 2009, 41, 666-676.	9.4	1,104
105	Genetic variants at CD28, PRDM1 and CD2/CD58 are associated with rheumatoid arthritis risk. <i>Nature Genetics</i> , 2009, 41, 1313-1318.	9.4	306
106	A re-evaluation of three putative functional single nucleotide polymorphisms in rheumatoid arthritis. <i>Annals of the Rheumatic Diseases</i> , 2009, 68, 1373-1375.	0.5	13
107	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, 2904-2905.	0.3	6
108	Genetic Variation in Efflux Transporters Influences Outcome to Methotrexate Therapy in Patients with Psoriasis. <i>Journal of Investigative Dermatology</i> , 2008, 128, 1925-1929.	0.3	109

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109	Rheumatoid arthritis susceptibility loci at chromosomes 10p15, 12q13 and 22q13. <i>Nature Genetics</i> , 2008, 40, 1156-1159.	9.4	143
110	Polymorphisms in the IL-12 β 2 and IL-23R Genes Are Associated with Psoriasis of Early Onset in a UK Cohort. <i>Journal of Investigative Dermatology</i> , 2008, 128, 1325-1327.	0.3	74
111	Polymorphisms in the PTPN22 region are associated with psoriasis of early onset. <i>British Journal of Dermatology</i> , 2008, 158, 962-968.	1.4	41
112	Association of the FCRL3 gene with rheumatoid arthritis: a further example of population specificity?. <i>Arthritis Research and Therapy</i> , 2008, 10, 405.	1.6	0
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, 2274-2279.	1.4	131
114	Investigation of association between the TRAF family genes and RA susceptibility. <i>Annals of the Rheumatic Diseases</i> , 2007, 66, 1322-1326.	0.5	41
115	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, 683-686.	0.5	30
116	Replication of Genome-Wide Association Signals in UK Samples Reveals Risk Loci for Type 2 Diabetes. <i>Science</i> , 2007, 316, 1336-1341.	6.0	2,040
117	Association scan of 14,500 nonsynonymous SNPs in four diseases identifies autoimmunity variants. <i>Nature Genetics</i> , 2007, 39, 1329-1337.	9.4	1,298
118	Rheumatoid arthritis association at 6q23. <i>Nature Genetics</i> , 2007, 39, 1431-1433.	9.4	361
119	Genome-wide association study of 14,000 cases of seven common diseases and 3,000 shared controls. <i>Nature</i> , 2007, 447, 661-678.	13.7	8,895
120	Localization of type 1 diabetes susceptibility to the MHC class I genes HLA-B and HLA-A. <i>Nature</i> , 2007, 450, 887-892.	13.7	493
121	Association of the FCRL3 gene with rheumatoid arthritis: a further example of population specificity?. <i>Arthritis Research and Therapy</i> , 2006, 8, R117.	1.6	36
122	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, 3417-3422.	6.7	21
123	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, 752-758.	6.7	33
124	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.	2.7	9
125	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, 1390-1393.	0.9	16
126	Investigation of polymorphisms in the PADI4 gene in determining severity of inflammatory polyarthritis. <i>Annals of the Rheumatic Diseases</i> , 2005, 64, 1311-1315.	0.5	32

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127	Polymorphisms in the tumour necrosis factor gene are not associated with severity of inflammatory polyarthritis. <i>Annals of the Rheumatic Diseases</i> , 2004, 63, 280-284.	0.5	30
128	Association of protein kinase C alpha (PRKCA) gene with multiple sclerosis in a UK population. <i>Brain</i> , 2004, 127, 1717-1722.	3.7	36
129	Polymorphisms of the equine major histocompatibility complex class II DRA locus. <i>Tissue Antigens</i> , 2004, 64, 173-179.	1.0	17
130	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, 729-735.	6.7	39
131	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, 748-752.	6.7	50
132	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, 1117-1121.	6.7	186
133	Evidence for a novel rheumatoid arthritis susceptibility locus on chromosome 6p. <i>Arthritis and Rheumatism</i> , 2004, 50, 3823-3830.	6.7	18
134	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.	2.6	209
135	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, 632-639.	6.7	184
136	Two novel polymorphisms in the human transforming growth factor beta 2 gene. <i>Genes and Immunity</i> , 2001, 2, 295-296.	2.2	8
137	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, 1901-1906.	1.4	52
138	Linkage of a marker in intron D of the estrogen synthase locus to rheumatoid arthritis. <i>Arthritis and Rheumatism</i> , 1999, 42, 1617-1620.	6.7	15
139	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, 904-908.	2.6	49
140	Risk-taking Behavior in Adolescents: The Paradigm. <i>Annals of the New York Academy of Sciences</i> , 1997, 817, 1-35.	1.8	69