

Anne Barton

List of Publications by Year in descending order

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Version: 2024-02-01

342
papers

33,319
citations

14614

66
h-index

4203

174
g-index

359
all docs

359
docs citations

359
times ranked

35406
citing authors

#	ARTICLE	IF	CITATIONS
1	Towards Personalising the Use of Biologics in Rheumatoid Arthritis: A Discrete Choice Experiment. <i>Patient</i> , 2022, 15, 109-119.	1.1	1
2	Pre-defined gene co-expression modules in rheumatoid arthritis transition towards molecular health following anti-TNF therapy. <i>Rheumatology</i> , 2022, 61, 4935-4944.	0.9	3
3	HLA-DRB1 haplotypes predict cardiovascular mortality in inflammatory polyarthritis independent of CRP and anti-CCP status. <i>Arthritis Research and Therapy</i> , 2022, 24, 90.	1.6	3
4	P200 Combining protein quantitative trait and genetic risk score analysis to identify biomarkers of treatment response to TNFi in patients with rheumatoid arthritis. <i>Rheumatology</i> , 2022, 61, .	0.9	0
5	OA24 Predicting drug immunogenicity to tumour necrosis factor inhibitors in patients with rheumatoid arthritis. <i>Rheumatology</i> , 2022, 61, .	0.9	0
6	P190 Unsupervised automated clustering of mass cytometry data identifies unique CD4+ T cell subsets in rheumatoid arthritis. <i>Rheumatology</i> , 2022, 61, .	0.9	0
7	OA15 Drivers of change in four and two component disease activity scores after etanercept treatment, in a multi-centre cohort of patients with established rheumatoid arthritis. <i>Rheumatology</i> , 2022, 61, .	0.9	0
8	P189 A longitudinal study of psychological predictors of response to adalimumab in patients with rheumatoid arthritis. <i>Rheumatology</i> , 2022, 61, .	0.9	1
9	Comparative Genetic Analysis of Psoriatic Arthritis and Psoriasis for the Discovery of Genetic Risk Factors and Risk Prediction Modeling. <i>Arthritis and Rheumatology</i> , 2022, 74, 1535-1543.	2.9	15
10	OA28 Exploring the potential of polygenic risk scores for predicting coronary artery disease in patients with rheumatoid arthritis. <i>Rheumatology</i> , 2022, 61, .	0.9	0
11	OA16 Therapeutic certolizumab pegol drug levels to achieve good EULAR response in patients with rheumatoid arthritis: results from the Biologics in Rheumatoid Arthritis Genetics and Genomics Study Syndicate (BRAGSS) cohort. <i>Rheumatology</i> , 2022, 61, .	0.9	0
12	Translating research into clinical practice: quality improvement to halve non-adherence to methotrexate. <i>Rheumatology</i> , 2021, 60, 125-131.	0.9	6
13	Changes in the illness perceptions of patients with rheumatoid arthritis over the first year of methotrexate therapy. <i>Rheumatology</i> , 2021, 60, 2355-2365.	0.9	11
14	Machine learning in precision medicine: lessons to learn. <i>Nature Reviews Rheumatology</i> , 2021, 17, 5-6.	3.5	18
15	Do people with rheumatoid arthritis maintain their physical activity level at treatment onset over the first year of methotrexate therapy?. <i>Rheumatology</i> , 2021, 60, 4633-4642.	0.9	5
16	Prediction of response of methotrexate in patients with rheumatoid arthritis using serum lipidomics. <i>Scientific Reports</i> , 2021, 11, 7266.	1.6	21
17	Transcriptome-wide study of TNF-inhibitor therapy in rheumatoid arthritis reveals early signature of successful treatment. <i>Arthritis Research and Therapy</i> , 2021, 23, 80.	1.6	11
18	O08 Developing a DNA methylation signature for predicting rheumatoid arthritis using a machine learning pipeline. <i>Rheumatology</i> , 2021, 60, .	0.9	0

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19	Comprehensive analysis of the major histocompatibility complex in systemic sclerosis identifies differential HLA associations by clinical and serological subtypes. <i>Annals of the Rheumatic Diseases</i> , 2021, 80, 1040-1047.	0.5	24
20	Cell-specific epigenetic drivers of pathogenesis in rheumatoid arthritis. <i>Epigenomics</i> , 2021, 13, 549-560.	1.0	4
21	Robust optimization of SWATH-MS workflow for human blood serum proteome analysis using a quality by design approach. <i>Clinical Proteomics</i> , 2021, 18, 20.	1.1	2
22	Application of information theoretic feature selection and machine learning methods for the development of genetic risk prediction models. <i>Scientific Reports</i> , 2021, 11, 23335.	1.6	10
23	The predictors of and reasons for non-adherence in an observational cohort of patients with rheumatoid arthritis commencing methotrexate. <i>Rheumatology</i> , 2020, 59, 213-223.	0.9	22
24	Alopecia areata is characterized by dysregulation in systemic type 17 and type 2 cytokines, which may contribute to disease-associated psychological morbidity. <i>British Journal of Dermatology</i> , 2020, 182, 130-137.	1.4	52
25	Adding value to real-world data: the role of biomarkers. <i>Rheumatology</i> , 2020, 59, 31-38.	0.9	11
26	Differential DNA methylation correlates with response to methotrexate in rheumatoid arthritis. <i>Rheumatology</i> , 2020, 59, 1364-1371.	0.9	43
27	Long-term outcomes of patients who rate symptoms of rheumatoid arthritis as "satisfactory". <i>Rheumatology</i> , 2020, 59, 1853-1861.	0.9	8
28	Proteomic analysis to define predictors of treatment response to adalimumab or methotrexate in rheumatoid arthritis patients. <i>Pharmacogenomics Journal</i> , 2020, 20, 516-523.	0.9	6
29	The use of missing values in proteomic data-independent acquisition mass spectrometry to enable disease activity discrimination. <i>Bioinformatics</i> , 2020, 36, 2217-2223.	1.8	29
30	Therapeutic monitoring of TNF inhibitors for rheumatoid arthritis: evidence required following NICE's recommendations. <i>Rheumatology Advances in Practice</i> , 2020, 4, rkaa023.	0.3	0
31	O11's 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
32	Characterisation of CD4+ T-cell subtypes using single cell RNA sequencing and the impact of cell number and sequencing depth. <i>Scientific Reports</i> , 2020, 10, 19825.	1.6	17
33	Genetic feature engineering enables characterisation of shared risk factors in immune-mediated diseases. <i>Genome Medicine</i> , 2020, 12, 106.	3.6	12
34	Investigation of genetically regulated gene expression and response to treatment in rheumatoid arthritis highlights an association between <i>IL18RAP</i> expression and treatment response. <i>Annals of the Rheumatic Diseases</i> , 2020, 79, 1446-1452.	0.5	13
35	Using functional genomics to advance the understanding of psoriatic arthritis. <i>Rheumatology</i> , 2020, 59, 3137-3146.	0.9	8
36	Predicting treatment response to IL6R blockers in rheumatoid arthritis. <i>Rheumatology</i> , 2020, 59, 3603-3610.	0.9	15

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37	Latent Class Trajectory Modeling of 2-Component Disease Activity Score in 28 Joints Identifies Multiple Rheumatoid Arthritis Phenotypes of Response to Biologic Disease-Modifying Antirheumatic Drugs. <i>Arthritis and Rheumatology</i> , 2020, 72, 1632-1642.	2.9	9
38	Personalized medicine in rheumatic diseases: how close are we to being able to use genetic biomarkers to predict response to TNF inhibitors?. <i>Expert Review of Clinical Immunology</i> , 2020, 16, 389-396.	1.3	8
39	Predictors of presenteeism, absenteeism and job loss in patients commencing methotrexate or biologic therapy for rheumatoid arthritis. <i>Rheumatology</i> , 2020, 59, 2908-2919.	0.9	17
40	A restricted spectrum of missense KMT2D variants cause a multiple malformations disorder distinct from Kabuki syndrome. <i>Genetics in Medicine</i> , 2020, 22, 867-877.	1.1	41
41	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
42	Association of Pharmacological Biomarkers with Treatment Response and Longterm Disability in Patients with Psoriatic Arthritis: Results from OUTPASS. <i>Journal of Rheumatology</i> , 2020, 47, 1204-1208.	1.0	10
43	Pharmacogenetics of TNF inhibitor response in rheumatoid arthritis utilizing the two-component disease activity score. <i>Pharmacogenomics</i> , 2020, 21, 1151-1156.	0.6	3
44	THU0022...DIFFERENTIAL DNA METHYLATION AS A PREDICTOR OF TOCILIZUMAB RESPONSE IN RHEUMATOID ARTHRITIS PATIENTS. <i>Annals of the Rheumatic Diseases</i> , 2020, 79, 224.1-224.	0.5	0
45	THU0005...VARIABILITY OF DNA METHYLATION IS A DRIVER OF LYMPHOCYTE DYSREGULATION IN EARLY RHEUMATOID ARTHRITIS.. <i>Annals of the Rheumatic Diseases</i> , 2020, 79, 215.2-215.	0.5	0
46	Investigating the pharmacogenetics of anti-TNF response in patients with rheumatoid arthritis utilising the re-weighted disease activity score: results from the biologics in rheumatoid arthritis genetics and genomics study syndicate. <i>Rheumatology</i> , 2019, 58, .	0.9	0
47	Distinct HLA Associations with Rheumatoid Arthritis Subsets Defined by Serological Subphenotype. <i>American Journal of Human Genetics</i> , 2019, 105, 616-624.	2.6	27
48	Translational genomics and precision medicine: Moving from the lab to the clinic. <i>Science</i> , 2019, 365, 1409-1413.	6.0	133
49	Using the Immunophenotype to Predict Response to Biologic Drugs in Rheumatoid Arthritis. <i>Journal of Personalized Medicine</i> , 2019, 9, 46.	1.1	13
50	Development and validation of a methotrexate adherence assay. <i>Annals of the Rheumatic Diseases</i> , 2019, 78, 1192-1197.	0.5	13
51	What do we mean by treatment response and can we predict it?. <i>Rheumatology</i> , 2019, 58, .	0.9	0
52	Chromatin interactions reveal novel gene targets for drug repositioning in rheumatic diseases. <i>Annals of the Rheumatic Diseases</i> , 2019, 78, 1127-1134.	0.5	23
53	Can interleukin IL-17A levels predict response to biologic treatment in patients with rheumatoid arthritis?. <i>Rheumatology</i> , 2019, 58, .	0.9	0
54	Association of response to TNF inhibitors in rheumatoid arthritis with quantitative trait loci for CD40 and CD39. <i>Annals of the Rheumatic Diseases</i> , 2019, 78, 1055-1061.	0.5	25

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55	Validity of a two-component imaging-derived disease activity score for improved assessment of synovitis in early rheumatoid arthritis. <i>Rheumatology</i> , 2019, 58, 1400-1409.	0.9	39
56	Expression of STAT3-regulated genes in circulating CD4+ T cells discriminates rheumatoid arthritis independently of clinical parameters in early arthritis. <i>Rheumatology</i> , 2019, 58, 1250-1258.	0.9	14
57	SAT0062â€¦STRATIFIED MEDICINE FOR RHEUMATOID ARTHRITIS: PREDICTING RESPONSE TO BIOLOGIC THERAPY USING IMMUNE CELL SIGNATURES. , 2019, , .		0
58	P106/O25â€¦DNA methylation in lymphocyte subsets as a mediator of genetic risk in early rheumatoid arthritis. , 2019, , .		0
59	FR10004â€¦CHROMATIN INTERACTIONS IN NOVEL CELL TYPES REVEAL PARK7 AND ERRF1 AS PUTATIVE CAUSAL GENES IN THE SUSCEPTIBILITY TO PSORIATIC ARTHRITIS. , 2019, , .		0
60	Prediction of infection risk in rheumatoid arthritis patients treated with biologics: are we any closer to risk stratification?. <i>Current Opinion in Rheumatology</i> , 2019, 31, 285-292.	2.0	39
61	Profiling of Gene Expression Biomarkers as a Classifier of Methotrexate Nonresponse in Patients With Rheumatoid Arthritis. <i>Arthritis and Rheumatology</i> , 2019, 71, 678-684.	2.9	50
62	Prediction of response to methotrexate in rheumatoid arthritis. <i>Expert Review of Clinical Immunology</i> , 2018, 14, 419-429.	1.3	23
63	Rheumatoid arthritis. <i>Nature Reviews Disease Primers</i> , 2018, 4, 18001.	18.1	1,441
64	Genotypic variability-based genome-wide association study identifies non-additive loci HLA-C and IL12B for psoriasis. <i>Journal of Human Genetics</i> , 2018, 63, 289-296.	1.1	9
65	Genetics of immune-mediated inflammatory diseases. <i>Clinical and Experimental Immunology</i> , 2018, 193, 3-12.	1.1	66
66	O12â€¦Validity of a 2-component imaging-derived disease activity score (2C-DAS28) for improved assessment of synovitis in early rheumatoid arthritis. <i>Rheumatology</i> , 2018, 57, .	0.9	0
67	248â€¦Genome-wide association study of response to tumour necrosis factor inhibitor therapy in rheumatoid arthritis. <i>Rheumatology</i> , 2018, 57, .	0.9	0
68	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
69	227â€¦Predictors of presenteeism and absenteeism in patients commencing treatment with methotrexate monotherapy or biologic therapy for rheumatoid arthritis. <i>Rheumatology</i> , 2018, 57, .	0.9	0
70	e111â€¦Patterns of the patient acceptable symptom state over 12 months following the initiation of methotrexate therapy in patients with rheumatoid arthritis, and the association between these patterns and disability and disease activity. <i>Rheumatology</i> , 2018, 57, .	0.9	0
71	P124â€¦Altered CD4+ T cell DNA methylation in early rheumatoid arthritis. , 2018, , .		0
72	O89â€¦The association between poor prognostic factors at methotrexate initiation and disease activity and disability over one year: results from the Rheumatoid Arthritis Medication Study. <i>Rheumatology</i> , 2018, 57, .	0.9	0

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73	e12â€fClinical phenotypes of patients with rheumatoid arthritis who identify as in a patient acceptable symptom state at methotrexate initiation and a comparison of the outcome of these phenotypes over 12 months. <i>Rheumatology</i> , 2018, 57, .	0.9	0
74	Prediction of treatment response in rheumatoid arthritis patients using genomeâ€wide SNP data. <i>Genetic Epidemiology</i> , 2018, 42, 754-771.	0.6	15
75	Increased DNA methylation variability in rheumatoid arthritis-discordant monozygotic twins. <i>Genome Medicine</i> , 2018, 10, 64.	3.6	71
76	Assessing the Role of DNA Methylation-Derived Neutrophil-to-Lymphocyte Ratio in Rheumatoid Arthritis. <i>Journal of Immunology Research</i> , 2018, 2018, 1-10.	0.9	13
77	Genome-wide association study of response to tumour necrosis factor inhibitor therapy in rheumatoid arthritis. <i>Pharmacogenomics Journal</i> , 2018, 18, 657-664.	0.9	41
78	Genome-wide association study of response to methotrexate in early rheumatoid arthritis patients. <i>Pharmacogenomics Journal</i> , 2018, 18, 528-538.	0.9	42
79	Psoriasis and Psoriatic Arthritis. , 2018, , 239-250.		0
80	Prediction of primary non-response to methotrexate therapy using demographic, clinical and psychosocial variables: results from the UK Rheumatoid Arthritis Medication Study (RAMS). <i>Arthritis Research and Therapy</i> , 2018, 20, 147.	1.6	73
81	The prevalence of co-morbidities and their impact on physical activity in people with inflammatory rheumatic diseases compared with the general population: results from the UK Biobank. <i>Rheumatology</i> , 2018, 57, 2172-2182.	0.9	30
82	THU0001â€...Differential methylation as a predictor of methotrexate response in patients with rheumatoid arthritis. , 2018, , .		0
83	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
84	High frequency of antidrug antibodies and association of random drug levels with efficacy in certolizumab pegol-treated patients with rheumatoid arthritis: results from the BRAGGSS cohort. <i>Annals of the Rheumatic Diseases</i> , 2017, 76, 208-213.	0.5	49
85	Drug-specific risk and characteristics of lupus and vasculitis-like events in patients with rheumatoid arthritis treated with TNFi: results from BSRBR-RA. <i>RMD Open</i> , 2017, 3, e000314.	1.8	29
86	HLA-A 31:01 is not associated with the development of methotrexate pneumonitis in the UK population: results from a genome-wide association study. <i>Annals of the Rheumatic Diseases</i> , 2017, 76, e51-e51.	0.5	11
87	A rare coding allele INFIH1is protective for psoriatic arthritis. <i>Annals of the Rheumatic Diseases</i> , 2017, 76, 1321-1324.	0.5	22
88	Genetics of rheumatoid arthritis susceptibility, severity, and treatment response. <i>Seminars in Immunopathology</i> , 2017, 39, 395-408.	2.8	73
89	Cross-phenotype association mapping of the MHC identifies genetic variants that differentiate psoriatic arthritis from psoriasis. <i>Annals of the Rheumatic Diseases</i> , 2017, 76, 1774-1779.	0.5	51
90	DNA methylation as a marker of response in rheumatoid arthritis. <i>Pharmacogenomics</i> , 2017, 18, 1323-1332.	0.6	22

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91	Genotypic variability based association identifies novel non-additive loci DHCR7 and IRF4 in sero-negative rheumatoid arthritis. <i>Scientific Reports</i> , 2017, 7, 5261.	1.6	20
92	Precision Medicine in Rheumatoid Arthritis. <i>Rheumatic Disease Clinics of North America</i> , 2017, 43, 377-387.	0.8	42
93	Brief Report: The Role of Rare Protein-coding Variants in Anti-Tumor Necrosis Factor Treatment Response in Rheumatoid Arthritis. <i>Arthritis and Rheumatology</i> , 2017, 69, 735-741.	2.9	8
94	Investigation of the genetic overlap between rheumatoid arthritis and psoriatic arthritis in a Greek population. <i>Scandinavian Journal of Rheumatology</i> , 2017, 46, 180-186.	0.6	14
95	05.10...Comparison of cd4+ and b lymphocyte expression quantitative trait associations at ra risk loci in untreated early arthritis patients. , 2017, , .		0
96	Stratified medicine in rheumatoid arthritis—the MATURA programme. <i>Rheumatology</i> , 2017, 56, 1247-1250.	0.9	22
97	THU0003...CD4+ and B lymphocyte expression quantitative traits at rheumatoid arthritis risk loci in untreated early arthritis: implications for causal gene identification?. , 2017, , .		2
98	THU0001...Differential methylation as a potential biomarker of methotrexate response in patients with rheumatoid arthritis. , 2017, , .		0
99	THU0004...Cross phenotype association mapping of the mhc identifies genetic variants that differentiate psoriatic arthritis from psoriasis. , 2017, , .		0
100	OP0298...Chromatin interactions reveal novel gene targets for drug repositioning in rheumatic diseases. , 2017, , .		0
101	FRI0724...Predictors of presenteeism and absenteeism in patients commencing treatment with methotrexate monotherapy or biologic therapy for rheumatoid arthritis. , 2017, , .		0
102	Genetics of Rheumatic Diseases. , 2017, , 327-343.		0
103	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
104	<i>KIF3A</i> and <i>IL-4</i> are disease-specific biomarkers for psoriatic arthritis susceptibility. <i>Oncotarget</i> , 2017, 8, 95401-95411.	0.8	12
105	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
106	HLA-DRB1 Amino Acid Positions 11/13, 71, and 74 Are Associated With Inflammation Level, Disease Activity, and the Health Assessment Questionnaire Score in Patients With Inflammatory Polyarthritis. <i>Arthritis and Rheumatology</i> , 2016, 68, 2618-2628.	2.9	23
107	OP0235...Identification of Novel Cd4+ Lymphocyte Expression Quantitative Trait Loci in Untreated Early Arthritis Patients. <i>Annals of the Rheumatic Diseases</i> , 2016, 75, 147.1-147.	0.5	0
108	Psychological factors predict adherence to methotrexate in rheumatoid arthritis; findings from a systematic review of rates, predictors and associations with patient-reported and clinical outcomes. <i>RMD Open</i> , 2016, 2, e000171.	1.8	40

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109	SAT0009â€¦Investigation of Differential Methylation as A Potential Biomarker of Methotrexate Response in Patients with Rheumatoid Arthritis. <i>Annals of the Rheumatic Diseases</i> , 2016, 75, 667.1-667.	0.5	0
110	Cryopreservation of cells does not substantially alter the DNA methylome of CD3+CD4+ T cells. <i>Scandinavian Journal of Rheumatology</i> , 2016, 45, 329-330.	0.6	0
111	A microcosting study of immunogenicity and tumour necrosis factor alpha inhibitor drug level tests for therapeutic drug monitoring in clinical practice. <i>Rheumatology</i> , 2016, 55, 2131-2137.	0.9	14
112	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
113	Replication of Associations of Genetic Loci Outside the HLA Region With Susceptibility to Antiâ€œCyclic Citrullinated Peptideâ€œNegative Rheumatoid Arthritis. <i>Arthritis and Rheumatology</i> , 2016, 68, 1603-1613.	2.9	33
114	Association Between Genetic Variation in <i>FOXO3</i> and Reductions in Inflammation and Disease Activity in Inflammatory Polyarthritis. <i>Arthritis and Rheumatology</i> , 2016, 68, 2629-2636.	2.9	32
115	Detection of anti-drug antibodies using a bridging ELISA compared with radioimmunoassay in adalimumab-treated rheumatoid arthritis patients with random drug levels. <i>Rheumatology</i> , 2016, 55, 2050-2055.	0.9	14
116	Rheumatoid arthritis patient perceptions on the value of predictive testing for treatments: a qualitative study. <i>BMC Musculoskeletal Disorders</i> , 2016, 17, 460.	0.8	9
117	Crowdsourced assessment of common genetic contribution to predicting anti-TNF treatment response in rheumatoid arthritis. <i>Nature Communications</i> , 2016, 7, 12460.	5.8	73
118	AB0727â€¦Increased Rates of Hypertension in Patients with Psoriatic Arthritis Compared To Psoriasis Alone: Results from The UK Biobank: Table 1.. <i>Annals of the Rheumatic Diseases</i> , 2016, 75, 1153.1-1153.	0.5	1
119	A6.13â€¦Identification of novel expression quantitative trait loci in CD4+T cells of untreated early arthritis patients. <i>Annals of the Rheumatic Diseases</i> , 2016, 75, A52.2-A52.	0.5	0
120	AB0005â€¦Weighted Gene Co-Expression Network Analysis Reveals Link between Protein Kinase Signalling and Response To Methotrexate in New-Onset Rheumatoid Arthritis. <i>Annals of the Rheumatic Diseases</i> , 2016, 75, 898.2-898.	0.5	0
121	Replication of a distinct psoriatic arthritis risk variant at the L23R locus. <i>Annals of the Rheumatic Diseases</i> , 2016, 75, 1417-1418.	0.5	9
122	Previously reported <i>PDE3A</i> â€œSLCO1C1 genetic variant does not correlate with anti-TNF response in a large UK rheumatoid arthritis cohort. <i>Pharmacogenomics</i> , 2016, 17, 715-720.	0.6	9
123	Association of Toll-like receptor 4 (TLR4) with chronic plaque type psoriasis and psoriatic arthritis. <i>Archives of Dermatological Research</i> , 2016, 308, 201-205.	1.1	35
124	One SNP at a Time: Moving beyond GWAS in Psoriasis. <i>Journal of Investigative Dermatology</i> , 2016, 136, 567-573.	0.3	48
125	Identifying a novel locus for psoriatic arthritis. <i>Rheumatology</i> , 2016, 55, 25-32.	0.9	13
126	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

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127	Anticarbamylated protein antibodies are associated with long-term disability and increased disease activity in patients with early inflammatory arthritis: results from the Norfolk Arthritis Register. <i>Annals of the Rheumatic Diseases</i> , 2016, 75, 1139-1144.	0.5	41
128	Identifying Causal Genes at the Multiple Sclerosis Associated Region 6q23 Using Capture Hi-C. <i>PLoS ONE</i> , 2016, 11, e0166923.	1.1	28
129	Do lean markers relate to exacerbation rate in chronic obstructive pulmonary disease? Preliminary results from AERIS study. <i>Proceedings of the Nutrition Society</i> , 2015, 74, .	0.4	0
130	Rheumatoid arthritis response to treatment across IgG1 allotype “ anti-TNF incompatibility: a case-only study. <i>Arthritis Research and Therapy</i> , 2015, 17, 63.	1.6	9
131	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
132	SAT0345“Lifestyle, Clinical and Psychosocial Predictors of Good Response to Methotrexate Therapy in the Rheumatoid Arthritis Medication Study (RAMS). <i>Annals of the Rheumatic Diseases</i> , 2015, 74, 783.3-784.	0.5	1
133	O53.“PTPN22 is Associated with Susceptibility to Psoriatic Arthritis but not Psoriasis: Evidence for a Further PSA-Specific Risk Locus. <i>Rheumatology</i> , 2015, , .	0.9	1
134	O54.“The Importance of IL-6-STAT3 Mediated Activation of Circulating CD4⁺ T Cells in the Pathogenesis of Early Seronegative Rheumatoid Arthritis: A Validation Study. <i>Rheumatology</i> , 2015, , .	0.9	0
135	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
136	Pharmacogenetics of Treatment Response in Psoriatic Arthritis. <i>Current Rheumatology Reports</i> , 2015, 17, 44.	2.1	16
137	Reply. <i>Arthritis Care and Research</i> , 2015, 67, 452-453.	1.5	0
138	Capture Hi-C reveals novel candidate genes and complex long-range interactions with related autoimmune risk loci. <i>Nature Communications</i> , 2015, 6, 10069.	5.8	161
139	PTPN22 is associated with susceptibility to psoriatic arthritis but not psoriasis: evidence for a further PsA-specific risk locus. <i>Annals of the Rheumatic Diseases</i> , 2015, 74, 1882-1885.	0.5	64
140	Impact of inadequate adherence on response to subcutaneously administered anti-tumour necrosis factor drugs: results from the Biologics in Rheumatoid Arthritis Genetics and Genomics Study Syndicate cohort. <i>Rheumatology</i> , 2015, 54, 494-499.	0.9	90
141	The skin microbiome in psoriatic arthritis: methodology development and pilot data. <i>Lancet, The</i> , 2015, 385, S27.	6.3	7
142	O10.“Risk and Characteristics of Drug-Induced Lupus in Patients Exposed to Tumour Necrosis Factor-± Inhibitor Therapy: Results from the British Society for Rheumatology Biologics Register for Rheumatoid Arthritis. <i>Rheumatology</i> , 2015, , .	0.9	0
143	Association of HLA-DRB1 Haplotypes With Rheumatoid Arthritis Severity, Mortality, and Treatment Response. <i>JAMA - Journal of the American Medical Association</i> , 2015, 313, 1645.	3.8	119
144	O49.“Personalized Genetic Medicine: Amino Acid Positions 11, 71 and 74 in HLA-DRB1 Predict Disease Severity, Mortality and Treatment Response in Rheumatoid Arthritis“Multi-Centre Prospective Cohort Studies. <i>Rheumatology</i> , 2015, , .	0.9	0

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145	A HPLC-SRM-MS based method for the detection and quantification of methotrexate in urine at doses used in clinical practice for patients with rheumatological disease: a potential measure of adherence. <i>Analyst, The</i> , 2015, 140, 1981-1987.	1.7	23
146	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
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