

Anne M Hinks

List of Publications by Year in descending order

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60
papers

11,073
citations

100601

38
h-index

162838

57
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61
all docs

61
docs citations

61
times ranked

17382
citing authors

#	ARTICLE	IF	CITATIONS
1	<i>IL1RN</i> Variation Influences Both Disease Susceptibility and Response to Recombinant Human Interleukin-1 Receptor Antagonist Therapy in Systemic Juvenile Idiopathic Arthritis. <i>Arthritis and Rheumatology</i> , 2018, 70, 1319-1330.	2.9	40
2	Brief Report: The Genetic Profile of Rheumatoid Factor-Positive Polyarticular Juvenile Idiopathic Arthritis Resembles That of Adult Rheumatoid Arthritis. <i>Arthritis and Rheumatology</i> , 2018, 70, 957-962.	2.9	53
3	Genetic architecture distinguishes systemic juvenile idiopathic arthritis from other forms of juvenile idiopathic arthritis: clinical and therapeutic implications. <i>Annals of the Rheumatic Diseases</i> , 2017, 76, 906-913.	0.5	123
4	<i>HLA-DRB1*11</i> and variants of the MHC class II locus are strong risk factors for systemic juvenile idiopathic arthritis. <i>Proceedings of the National Academy of Sciences of the United States of America</i> , 2015, 112, 15970-15975.	3.3	139
5	The genetics of juvenile idiopathic arthritis: current understanding and future prospects. <i>Rheumatology</i> , 2014, 53, 592-599.	0.9	31
6	Autoinflammatory gene polymorphisms and susceptibility to UK juvenile idiopathic arthritis. <i>Pediatric Rheumatology</i> , 2013, 11, 14.	0.9	18
7	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
8	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
9	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
10	Association of the IL-10 Gene Family Locus on Chromosome 1 with Juvenile Idiopathic Arthritis (JIA). <i>PLoS ONE</i> , 2012, 7, e47673.	1.1	26
11	Genome-wide association analysis of juvenile idiopathic arthritis identifies a new susceptibility locus at chromosomal region 3q13. <i>Arthritis and Rheumatism</i> , 2012, 64, 2781-2791.	6.7	62
12	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
13	Association of the 5-aminoimidazole-4-carboxamide ribonucleotide transformylase gene with response to methotrexate in juvenile idiopathic arthritis. <i>Annals of the Rheumatic Diseases</i> , 2011, 70, 1395-1400.	0.5	62
14	The rheumatoid arthritis and juvenile idiopathic arthritis associated major (A) allele of rs2104286 is a loss of expression variant of IL2RA. <i>Annals of the Rheumatic Diseases</i> , 2011, 70, A6-A6.	0.5	0
15	Study of the common genetic background for rheumatoid arthritis and systemic lupus erythematosus. <i>Annals of the Rheumatic Diseases</i> , 2011, 70, 463-468.	0.5	130
16	Generation of novel pharmacogenomic candidates in response to methotrexate in juvenile idiopathic arthritis: correlation between gene expression and genotype. <i>Pharmacogenetics and Genomics</i> , 2010, 20, 665-676.	0.7	49
17	Confirmation of association of the REL locus with rheumatoid arthritis susceptibility in the UK population. <i>Annals of the Rheumatic Diseases</i> , 2010, 69, 1572-1573.	0.5	32
18	Association of the AFF3 gene and IL2/IL21 gene region with juvenile idiopathic arthritis. <i>Genes and Immunity</i> , 2010, 11, 194-198.	2.2	54

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19	Association of the CCR5 gene with juvenile idiopathic arthritis. <i>Genes and Immunity</i> , 2010, 11, 584-589.	2.2	24
20	Genome-wide association study of CNVs in 16,000 cases of eight common diseases and 3,000 shared controls. <i>Nature</i> , 2010, 464, 713-720.	13.7	737
21	Genome-wide association study meta-analysis identifies seven new rheumatoid arthritis risk loci. <i>Nature Genetics</i> , 2010, 42, 508-514.	9.4	1,132
22	Combined effects of three independent SNPs greatly increase the risk estimate for RA at 6q23. <i>Human Molecular Genetics</i> , 2010, 19, 4544-4544.	1.4	0
23	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
24	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
25	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
26	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
27	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
28	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
29	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
30	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
31	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
32	Investigating the viability of genetic screening/testing for RA susceptibility using combinations of five confirmed risk loci. <i>Rheumatology</i> , 2009, 48, 1369-1374.	0.9	20
33	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
34	Association of the IL2RA/CD25 gene with juvenile idiopathic arthritis. <i>Arthritis and Rheumatism</i> , 2009, 60, 251-257.	6.7	93
35	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
36	CD226 Gly307Ser association with multiple autoimmune diseases. <i>Genes and Immunity</i> , 2009, 10, 5-10.	2.2	227

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37	Genome-wide association study identifies eight loci associated with blood pressure. <i>Nature Genetics</i> , 2009, 41, 666-676.	9.4	1,104
38	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
39	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
40	Autoinflammatory genes and susceptibility to psoriatic juvenile idiopathic arthritis. <i>Arthritis and Rheumatism</i> , 2008, 58, 2142-2146.	6.7	64
41	Rheumatoid arthritis susceptibility loci at chromosomes 10p15, 12q13 and 22q13. <i>Nature Genetics</i> , 2008, 40, 1156-1159.	9.4	143
42	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
43	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
44	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
45	SNPs in the FOXP3 gene region show no association with Juvenile Idiopathic Arthritis in a UK Caucasian population. <i>Rheumatology</i> , 2007, 46, 1263-1265.	0.9	17
46	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
47	Rheumatoid arthritis association at 6q23. <i>Nature Genetics</i> , 2007, 39, 1431-1433.	9.4	361
48	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
49	Fine mapping of genes within the IDDM8 region in rheumatoid arthritis. <i>Arthritis Research and Therapy</i> , 2006, 8, R145.	1.6	7
50	PTPN22: a confirmed rheumatoid arthritis susceptibility gene?. <i>Future Rheumatology</i> , 2006, 1, 153-158.	0.2	0
51	Association between the PTPN22 gene and rheumatoid arthritis and juvenile idiopathic arthritis in a UK population: Further support that PTPN22 is an autoimmunity gene. <i>Arthritis and Rheumatism</i> , 2005, 52, 1694-1699.	6.7	266
52	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
53	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
54	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

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55	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
56	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
57	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
58	Evidence for a novel rheumatoid arthritis susceptibility locus on chromosome 6p. <i>Arthritis and Rheumatism</i> , 2004, 50, 3823-3830.	6.7	18
59	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
60	Linkage analysis of cross-sectional and longitudinally derived phenotypic measures to identify loci influencing blood pressure. <i>BMC Genetics</i> , 2003, 4, S26.	2.7	2