

Javier Martin

List of Publications by Year in descending order

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Version: 2024-02-01

555
papers

29,448
citations

9234

74
h-index

10127

140
g-index

584
all docs

584
docs citations

584
times ranked

31091
citing authors

#	ARTICLE	IF	CITATIONS
1	Genetics of rheumatoid arthritis contributes to biology and drug discovery. <i>Nature</i> , 2014, 506, 376-381.	13.7	1,974
2	Genomewide Association Study of Severe Covid-19 with Respiratory Failure. <i>New England Journal of Medicine</i> , 2020, 383, 1522-1534.	13.9	1,548
3	Genetic mechanisms of critical illness in COVID-19. <i>Nature</i> , 2021, 591, 92-98.	13.7	1,014
4	Genetic association analyses implicate aberrant regulation of innate and adaptive immunity genes in the pathogenesis of systemic lupus erythematosus. <i>Nature Genetics</i> , 2015, 47, 1457-1464.	9.4	730
5	Identification of multiple risk variants for ankylosing spondylitis through high-density genotyping of immune-related loci. <i>Nature Genetics</i> , 2013, 45, 730-738.	9.4	699
6	A common haplotype of interferon regulatory factor 5 (IRF5) regulates splicing and expression and is associated with increased risk of systemic lupus erythematosus. <i>Nature Genetics</i> , 2006, 38, 550-555.	9.4	593
7	High-density genetic mapping identifies new susceptibility loci for rheumatoid arthritis. <i>Nature Genetics</i> , 2012, 44, 1336-1340.	9.4	558
8	Epidemiology of giant cell arteritis and polymyalgia rheumatica. <i>Arthritis and Rheumatism</i> , 2009, 61, 1454-1461.	6.7	460
9	Functional variants in the B-cell gene BANK1 are associated with systemic lupus erythematosus. <i>Nature Genetics</i> , 2008, 40, 211-216.	9.4	436
10	Genome-wide association study of systemic sclerosis identifies CD247 as a new susceptibility locus. <i>Nature Genetics</i> , 2010, 42, 426-429.	9.4	351
11	Transancestral mapping and genetic load in systemic lupus erythematosus. <i>Nature Communications</i> , 2017, 8, 16021.	5.8	314
12	HLA-DRB1 and persistent chronic inflammation contribute to cardiovascular events and cardiovascular mortality in patients with rheumatoid arthritis. <i>Arthritis and Rheumatism</i> , 2007, 57, 125-132.	6.7	312
13	Rheumatoid Arthritis: A Disease Associated with Accelerated Atherogenesis. <i>Seminars in Arthritis and Rheumatism</i> , 2005, 35, 8-17.	1.6	282
14	Association of a functional single-nucleotide polymorphism of PTPN22, encoding lymphoid protein phosphatase, with rheumatoid arthritis and systemic lupus erythematosus. <i>Arthritis and Rheumatism</i> , 2005, 52, 219-224.	6.7	275
15	Association of a functional variant downstream of TNFAIP3 with systemic lupus erythematosus. <i>Nature Genetics</i> , 2011, 43, 253-258.	9.4	242
16	Evaluation of the TREX1 gene in a large multi-ancestral lupus cohort. <i>Genes and Immunity</i> , 2011, 12, 270-279.	2.2	226
17	High prevalence of subclinical atherosclerosis in psoriatic arthritis patients without clinically evident cardiovascular disease or classic atherosclerosis risk factors. <i>Arthritis and Rheumatism</i> , 2007, 57, 1074-1080.	6.7	220
18	Major histocompatibility complex associations of ankylosing spondylitis are complex and involve further epistasis with ERAP1. <i>Nature Communications</i> , 2015, 6, 7146.	5.8	220

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19	Carotid Intima-Media Thickness Predicts the Development of Cardiovascular Events in Patients with Rheumatoid Arthritis. <i>Seminars in Arthritis and Rheumatism</i> , 2009, 38, 366-371.	1.6	211
20	Identification of Novel Genetic Markers Associated with Clinical Phenotypes of Systemic Sclerosis through a Genome-Wide Association Strategy. <i>PLoS Genetics</i> , 2011, 7, e1002178.	1.5	201
21	Endothelial dysfunction in psoriatic arthritis patients without clinically evident cardiovascular disease or classic atherosclerosis risk factors. <i>Arthritis and Rheumatism</i> , 2007, 57, 287-293.	6.7	185
22	ImmunoChip Analysis Identifies Multiple Susceptibility Loci for Systemic Sclerosis. <i>American Journal of Human Genetics</i> , 2014, 94, 47-61.	2.6	182
23	Giant Cell Arteritis in Northwestern Spain. <i>Medicine (United States)</i> , 2007, 86, 61-68.	0.4	168
24	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
25	The STAT4 gene influences the genetic predisposition to systemic sclerosis phenotype. <i>Human Molecular Genetics</i> , 2009, 18, 2071-2077.	1.4	163
26	Identification of IRF8, TMEM39A, and IKZF3-ZBP2 as Susceptibility Loci for Systemic Lupus Erythematosus in a Large-Scale Multiracial Replication Study. <i>American Journal of Human Genetics</i> , 2012, 90, 648-660.	2.6	161
27	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
28	Investigating the Causal Relationship of C-Reactive Protein with 32 Complex Somatic and Psychiatric Outcomes: A Large-Scale Cross-Consortium Mendelian Randomization Study. <i>PLoS Medicine</i> , 2016, 13, e1001976.	3.9	150
29	Giant Cell Arteritis. <i>Medicine (United States)</i> , 2005, 84, 277-290.	0.4	149
30	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
31	The IL23R Arg381Gln non-synonymous polymorphism confers susceptibility to ankylosing spondylitis. <i>Annals of the Rheumatic Diseases</i> , 2008, 67, 1451-1454.	0.5	142
32	Unraveling Multiple MHC Gene Associations with Systemic Lupus Erythematosus: Model Choice Indicates a Role for HLA Alleles and Non-HLA Genes in Europeans. <i>American Journal of Human Genetics</i> , 2012, 91, 778-793.	2.6	140
33	STAT4 associates with systemic lupus erythematosus through two independent effects that correlate with gene expression and act additively with IRF5 to increase risk. <i>Annals of the Rheumatic Diseases</i> , 2009, 68, 1746-1753.	0.5	138
34	Protection against anti-citrullinated protein antibody-positive rheumatoid arthritis is predominantly associated with HLA-DRB1*1301: A meta-analysis of HLA-DRB1 associations with anti-citrullinated protein antibody-positive and anti-citrullinated protein antibody-negative rheumatoid arthritis in four European populations. <i>Arthritis and Rheumatism</i> , 2010, 62, 1236-1245.	6.7	135
35	Involvement of Fcγ3 receptor IIIA genotypes in susceptibility to rheumatoid arthritis. <i>Arthritis and Rheumatism</i> , 2000, 43, 735.	6.7	130
36	Association of the C8orf13-BLK region with systemic sclerosis in North-American and European populations. <i>Journal of Autoimmunity</i> , 2010, 34, 155-162.	3.0	123

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37	Fine-mapping and functional studies highlight potential causal variants for rheumatoid arthritis and type 1 diabetes. <i>Nature Genetics</i> , 2018, 50, 1366-1374.	9.4	122
38	The High Prevalence of Subclinical Atherosclerosis in Patients With Ankylosing Spondylitis Without Clinically Evident Cardiovascular Disease. <i>Medicine (United States)</i> , 2009, 88, 358-365.	0.4	121
39	TYK2 Protein-Coding Variants Protect against Rheumatoid Arthritis and Autoimmunity, with No Evidence of Major Pleiotropic Effects on Non-Autoimmune Complex Traits. <i>PLoS ONE</i> , 2015, 10, e0122271.	1.1	120
40	PTPN22 C1858T polymorphism in Colombian patients with autoimmune diseases. <i>Genes and Immunity</i> , 2005, 6, 628-631.	2.2	117
41	TNFA \sim 308 (rs1800629) polymorphism is associated with a higher risk of cardiovascular disease in patients with rheumatoid arthritis. <i>Atherosclerosis</i> , 2011, 216, 125-130.	0.4	116
42	A genome-wide association study identifies a functional ERAP2 haplotype associated with birdshot chorioretinopathy. <i>Human Molecular Genetics</i> , 2014, 23, 6081-6087.	1.4	115
43	Polymorphism at the TNF loci in rheumatoid arthritis. <i>Tissue Antigens</i> , 1997, 49, 74-78.	1.0	114
44	Kallikrein genes are associated with lupus and glomerular basement membrane-specific antibody-induced nephritis in mice and humans. <i>Journal of Clinical Investigation</i> , 2009, 119, 911-923.	3.9	114
45	Polymorphisms of toll-like receptor 2 and 4 genes in rheumatoid arthritis and systemic lupus erythematosus. <i>Tissue Antigens</i> , 2004, 63, 54-57.	1.0	112
46	Phenotypic associations of genetic susceptibility loci in systemic lupus erythematosus. <i>Annals of the Rheumatic Diseases</i> , 2011, 70, 1752-1757.	0.5	110
47	Systemic Sclerosis in Northwestern Spain. <i>Medicine (United States)</i> , 2008, 87, 272-280.	0.4	109
48	Cardiovascular risk assessment in patients with rheumatoid arthritis: The relevance of clinical, genetic and serological markers. <i>Autoimmunity Reviews</i> , 2016, 15, 1013-1030.	2.5	107
49	Association of PDCD1 with susceptibility to systemic lupus erythematosus. <i>Arthritis and Rheumatism</i> , 2004, 50, 2590-2597.	6.7	106
50	A loss-of-function variant of PTPN22 is associated with reduced risk of systemic lupus erythematosus. <i>Human Molecular Genetics</i> , 2008, 18, 569-579.	1.4	106
51	Structural insertion/deletion variation in IRF5 is associated with a risk haplotype and defines the precise IRF5 isoforms expressed in systemic lupus erythematosus. <i>Arthritis and Rheumatism</i> , 2007, 56, 1234-1241.	6.7	105
52	A systemic sclerosis and systemic lupus erythematosus pan-meta-GWAS reveals new shared susceptibility loci. <i>Human Molecular Genetics</i> , 2013, 22, 4021-4029.	1.4	104
53	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
54	GWAS for systemic sclerosis identifies multiple risk loci and highlights fibrotic and vasculopathy pathways. <i>Nature Communications</i> , 2019, 10, 4955.	5.8	100

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55	Genetic associations of LYN with systemic lupus erythematosus. <i>Genes and Immunity</i> , 2009, 10, 397-403.	2.2	99
56	Identification of CSK as a systemic sclerosis genetic risk factor through Genome Wide Association Study follow-up. <i>Human Molecular Genetics</i> , 2012, 21, 2825-2835.	1.4	98
57	Endothelial Dysfunction, Carotid Intima-Media Thickness, and Accelerated Atherosclerosis in Rheumatoid Arthritis. <i>Seminars in Arthritis and Rheumatism</i> , 2008, 38, 67-70.	1.6	96
58	BANK1 functional variants are associated with susceptibility to diffuse systemic sclerosis in Caucasians. <i>Annals of the Rheumatic Diseases</i> , 2010, 69, 700-705.	0.5	96
59	Association of <i>STAT4</i> with rheumatoid arthritis: A replication study in three European populations. <i>Arthritis and Rheumatism</i> , 2008, 58, 1974-1980.	6.7	93
60	Therapeutic Effect of Vasoactive Intestinal Peptide on Experimental Autoimmune Encephalomyelitis. <i>American Journal of Pathology</i> , 2006, 168, 1179-1188.	1.9	91
61	Evidence of association of macrophage migration inhibitory factor gene polymorphisms with systemic lupus erythematosus. <i>Genes and Immunity</i> , 2006, 7, 433-436.	2.2	91
62	Analysis of autosomal genes reveals gene×sex interactions and higher total genetic risk in men with systemic lupus erythematosus. <i>Annals of the Rheumatic Diseases</i> , 2012, 71, 694-699.	0.5	87
63	IL-6 promoter polymorphisms in rheumatoid arthritis. <i>Genes and Immunity</i> , 2000, 1, 338-340.	2.2	86
64	Effect of anti-tumor necrosis factor $\hat{\pm}$ therapy on the progression of subclinical atherosclerosis in severe rheumatoid arthritis. <i>Arthritis and Rheumatism</i> , 2006, 55, 150-153.	6.7	86
65	STAT4 but not TRAF1/C5 variants influence the risk of developing rheumatoid arthritis and systemic lupus erythematosus in Colombians. <i>Genes and Immunity</i> , 2008, 9, 379-382.	2.2	86
66	<i>ERAP2</i> is associated with ankylosing spondylitis in <i>HLA-B27</i> -positive and <i>HLA-B27</i> -negative patients. <i>Annals of the Rheumatic Diseases</i> , 2015, 74, 1627-1629.	0.5	86
67	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
68	Genetic component of giant cell arteritis. <i>Rheumatology</i> , 2014, 53, 6-18.	0.9	83
69	3D Surgical Printing Cutting Guides for Open-Wedge High Tibial Osteotomy: Do It Yourself. <i>Journal of Knee Surgery</i> , 2016, 29, 690-695.	0.9	82
70	Association of the CT60 marker of the CTLA4 gene with systemic lupus erythematosus. <i>Arthritis and Rheumatism</i> , 2004, 50, 2211-2215.	6.7	81
71	Genome-wide meta-analysis reveals shared new <i>loci</i> in systemic seropositive rheumatic diseases. <i>Annals of the Rheumatic Diseases</i> , 2019, 78, 311-319.	0.5	81
72	Integrative Analysis Reveals a Molecular Stratification of Systemic Autoimmune Diseases. <i>Arthritis and Rheumatology</i> , 2021, 73, 1073-1085.	2.9	81

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73	Association of ATG16L1 and IRGM genes polymorphisms with inflammatory bowel disease: a meta-analysis approach. <i>Genes and Immunity</i> , 2009, 10, 356-364.	2.2	78
74	Genetic variants associated with antithyroid drug-induced agranulocytosis: a genome-wide association study in a European population. <i>Lancet Diabetes and Endocrinology</i> , 2016, 4, 507-516.	5.5	78
75	A Genome-wide Association Study Identifies Risk Alleles in Plasminogen and P4HA2 Associated with Giant Cell Arteritis. <i>American Journal of Human Genetics</i> , 2017, 100, 64-74.	2.6	78
76	Genetic basis of rheumatoid arthritis. <i>Biomedicine and Pharmacotherapy</i> , 2006, 60, 656-662.	2.5	77
77	A rare polymorphism in the gene for Toll-like receptor 2 is associated with systemic sclerosis phenotype and increases the production of inflammatory mediators. <i>Arthritis and Rheumatism</i> , 2012, 64, 264-271.	6.7	77
78	Metabolic Syndrome Is Associated with Increased Arterial Stiffness and Biomarkers of Subclinical Atherosclerosis in Patients with Systemic Lupus Erythematosus. <i>Journal of Rheumatology</i> , 2009, 36, 2204-2211.	1.0	75
79	Analysis of the influence of PTPN22 gene polymorphisms in systemic sclerosis. <i>Annals of the Rheumatic Diseases</i> , 2011, 70, 454-462.	0.5	75
80	Association of ferritin autoantibodies with giant cell arteritis/polymyalgia rheumatica. <i>Annals of the Rheumatic Diseases</i> , 2012, 71, 943-947.	0.5	75
81	A GWAS follow-up study reveals the association of the IL12RB2 gene with systemic sclerosis in Caucasian populations. <i>Human Molecular Genetics</i> , 2012, 21, 926-933.	1.4	74
82	The IRF5-TNPO3 association with systemic lupus erythematosus has two components that other autoimmune disorders variably share. <i>Human Molecular Genetics</i> , 2015, 24, 582-596.	1.4	74
83	Differential association of two PTPN22 coding variants with Crohn's disease and ulcerative colitis. <i>Inflammatory Bowel Diseases</i> , 2011, 17, 2287-2294.	0.9	73
84	Meta-analysis of Immunochip data of four autoimmune diseases reveals novel single-disease and cross-phenotype associations. <i>Genome Medicine</i> , 2018, 10, 97.	3.6	73
85	Identification of a Systemic Lupus Erythematosus Susceptibility Locus at 11p13 between PDHX and CD44 in a Multiethnic Study. <i>American Journal of Human Genetics</i> , 2011, 88, 83-91.	2.6	72
86	IRF5 polymorphism predicts prognosis in patients with systemic sclerosis. <i>Annals of the Rheumatic Diseases</i> , 2012, 71, 1197-1202.	0.5	72
87	NFKB1-94ATTG ins/del polymorphism (rs28362491) is associated with cardiovascular disease in patients with rheumatoid arthritis. <i>Atherosclerosis</i> , 2012, 224, 426-429.	0.4	72
88	HLA-DRA variants predict penicillin allergy in genome-wide fine-mapping genotyping. <i>Journal of Allergy and Clinical Immunology</i> , 2015, 135, 253-259.e10.	1.5	72
89	Genetics of immunoglobulin-A vasculitis (Henoch-Schönlein purpura): An updated review. <i>Autoimmunity Reviews</i> , 2018, 17, 301-315.	2.5	72
90	Chemokine receptor CCR5 polymorphisms and Chagas disease cardiomyopathy. <i>Tissue Antigens</i> , 2001, 58, 154-158.	1.0	71

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91	A1298C polymorphism in the MTHFR gene predisposes to cardiovascular risk in rheumatoid arthritis. <i>Arthritis Research and Therapy</i> , 2010, 12, R71.	1.6	71
92	Analysis of a Functional BTNL2 Polymorphism in Type 1 Diabetes, Rheumatoid Arthritis, and Systemic Lupus Erythematosus. <i>Human Immunology</i> , 2005, 66, 1235-1241.	1.2	70
93	Genetically determined Amerindian ancestry correlates with increased frequency of risk alleles for systemic lupus erythematosus. <i>Arthritis and Rheumatism</i> , 2010, 62, 3722-3729.	6.7	70
94	Impact of genetic ancestry and sociodemographic status on the clinical expression of systemic lupus erythematosus in American Indian and European populations. <i>Arthritis and Rheumatism</i> , 2012, 64, 3687-3694.	6.7	70
95	ABIN1 Dysfunction as a Genetic Basis for Lupus Nephritis. <i>Journal of the American Society of Nephrology: JASN</i> , 2013, 24, 1743-1754.	3.0	70
96	Association of two independent functional risk haplotypes in <i>TNIP1</i> with systemic lupus erythematosus. <i>Arthritis and Rheumatism</i> , 2012, 64, 3695-3705.	6.7	69
97	<i>HO1</i> promoter polymorphism associated with rheumatoid arthritis. <i>Arthritis and Rheumatism</i> , 2007, 56, 3953-3958.	6.7	68
98	Genetic and physical interaction of the B-cell systemic lupus erythematosus-associated genes <i>BANK1</i> and <i>BLK</i> . <i>Annals of the Rheumatic Diseases</i> , 2012, 71, 136-142.	0.5	67
99	Common variants at PVT1, ATG13 and AMBRA1, AHI1 and CLEC16A are associated with selective IgA deficiency. <i>Nature Genetics</i> , 2016, 48, 1425-1429.	9.4	67
100	Association of MicroRNA-618 Expression With Altered Frequency and Activation of Plasmacytoid Dendritic Cells in Patients With Systemic Sclerosis. <i>Arthritis and Rheumatology</i> , 2017, 69, 1891-1902.	2.9	67
101	A new molecular classification to drive precision treatment strategies in primary Sjögren's syndrome. <i>Nature Communications</i> , 2021, 12, 3523.	5.8	67
102	HLA class I loss and PD-L1 expression in lung cancer: impact on T-cell infiltration and immune escape. <i>Oncotarget</i> , 2018, 9, 4120-4133.	0.8	66
103	Prevalence of hepatitis C serum antibody in autoimmune diseases. <i>Journal of Autoimmunity</i> , 2009, 32, 261-266.	3.0	65
104	Cross-disorder analysis of schizophrenia and 19 immune-mediated diseases identifies shared genetic risk. <i>Human Molecular Genetics</i> , 2019, 28, 3498-3513.	1.4	65
105	Replication of an Association Between IL23R Gene Polymorphism With Inflammatory Bowel Disease. <i>Clinical Gastroenterology and Hepatology</i> , 2007, 5, 977-981.e2.	2.4	64
106	The <i>PTPN22</i> R263Q polymorphism is a risk factor for rheumatoid arthritis in Caucasian case-control samples. <i>Arthritis and Rheumatism</i> , 2011, 63, 365-372.	6.7	64
107	Transancestral mapping of the MHC region in systemic lupus erythematosus identifies new independent and interacting loci at <i>MSH5</i> , <i>HLA-DPB1</i> and <i>HLA-G</i> . <i>Annals of the Rheumatic Diseases</i> , 2012, 71, 777-784.	0.5	64
108	The telomeric part of the HLA region predisposes to rheumatoid arthritis independently of the class II loci. <i>Human Immunology</i> , 2001, 62, 75-84.	1.2	63

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109	A replication study confirms the association of <i>TNFSF4 (OX40L)</i> polymorphisms with systemic sclerosis in a large European cohort. <i>Annals of the Rheumatic Diseases</i> , 2011, 70, 638-641.	0.5	63
110	Novel identification of the <i>IRF7</i> region as an antacentromere autoantibody propensity locus in systemic sclerosis. <i>Annals of the Rheumatic Diseases</i> , 2012, 71, 114-119.	0.5	62
111	A genome-wide association study of rheumatoid arthritis without antibodies against citrullinated peptides. <i>Annals of the Rheumatic Diseases</i> , 2015, 74, e15-e15.	0.5	62
112	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
113	Inducible but not endothelial nitric oxide synthase polymorphism is associated with susceptibility to rheumatoid arthritis in northwest Spain. <i>British Journal of Rheumatology</i> , 2004, 43, 1182-1185.	2.5	61
114	Immunogenetics of systemic sclerosis: Defining heritability, functional variants and shared-autoimmunity pathways. <i>Journal of Autoimmunity</i> , 2015, 64, 53-65.	3.0	61
115	A functional variant of vascular endothelial growth factor is associated with severe ischemic complications in giant cell arteritis. <i>Journal of Rheumatology</i> , 2005, 32, 1737-41.	1.0	61
116	Variation in the <i>ICAM1</i> – <i>ICAM4</i> – <i>ICAM5</i> locus is associated with systemic lupus erythematosus susceptibility in multiple ancestries. <i>Annals of the Rheumatic Diseases</i> , 2012, 71, 1809-1814.	0.5	60
117	Changes in macrophage transcriptome associate with systemic sclerosis and mediate <i>GSDMA</i> contribution to disease risk. <i>Annals of the Rheumatic Diseases</i> , 2018, 77, 596-601.	0.5	60
118	Epistatic interaction between <i>FCRL3</i> and <i>NFAB1</i> genes in Spanish patients with rheumatoid arthritis. <i>Annals of the Rheumatic Diseases</i> , 2006, 65, 1188-1191.	0.5	59
119	Unraveling the genetic component of systemic sclerosis. <i>Human Genetics</i> , 2012, 131, 1023-1037.	1.8	59
120	Role of <i>MYH9</i> and <i>APOL1</i> in African and non-African populations with lupus nephritis. <i>Genes and Immunity</i> , 2012, 13, 232-238.	2.2	58
121	A combined large-scale meta-analysis identifies <i>COG6</i> as a novel shared risk locus for rheumatoid arthritis and systemic lupus erythematosus. <i>Annals of the Rheumatic Diseases</i> , 2017, 76, 286-294.	0.5	58
122	<i>CD40</i> : Novel Association with Crohn's Disease and Replication in Multiple Sclerosis Susceptibility. <i>PLoS ONE</i> , 2010, 5, e11520.	1.1	56
123	Confirmation of <i>TNIP1</i> but not <i>RHOB</i> and <i>PSORS1C1</i> as systemic sclerosis risk factors in a large independent replication study. <i>Annals of the Rheumatic Diseases</i> , 2013, 72, 602-607.	0.5	56
124	Isolation and characterization of the gene encoding histone H2A from <i>Trypanosoma cruzi</i> . <i>Molecular and Biochemical Parasitology</i> , 1994, 64, 1-10.	0.5	55
125	Genetic Influence of <i>PTPN22 R620W</i> Polymorphism in Tuberculosis. <i>Human Immunology</i> , 2005, 66, 1242-1247.	1.2	55
126	Identification of a new putative functional <i>IL18</i> gene variant through an association study in systemic lupus erythematosus. <i>Human Molecular Genetics</i> , 2009, 18, 3739-3748.	1.4	54

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127	Analysis of IRF5 gene functional polymorphisms in rheumatoid arthritis. <i>Arthritis and Rheumatism</i> , 2006, 54, 3815-3819.	6.7	53
128	Contribution of MHC class I region to genetic susceptibility for giant cell arteritis. <i>Rheumatology</i> , 2007, 46, 431-434.	0.9	53
129	Genetically Determined Partial Complement C4 Deficiency States Are Not Independent Risk Factors for SLE in UK and Spanish Populations. <i>American Journal of Human Genetics</i> , 2012, 90, 445-456.	2.6	53
130	Evidence for gene-gene epistatic interactions among susceptibility loci for systemic lupus erythematosus. <i>Arthritis and Rheumatism</i> , 2012, 64, 485-492.	6.7	53
131	Investigation of the IL23R gene in a Spanish rheumatoid arthritis cohort. <i>Human Immunology</i> , 2007, 68, 681-684.	1.2	52
132	Asymptomatic Hyperuricemia and Serum Uric Acid Concentration Correlate with Subclinical Atherosclerosis in Psoriatic Arthritis Patients Without Clinically Evident Cardiovascular Disease. Drs. Gonzalez-Gay, Gonzalez-Juanatey, and Llorca contributed equally to this study. <i>Seminars in Arthritis and Rheumatism</i> , 2009, 39, 157-162.	1.6	52
133	New insight on the Xq28 association with systemic sclerosis. <i>Annals of the Rheumatic Diseases</i> , 2013, 72, 2032-2038.	0.5	52
134	Analysis of the common genetic component of large-vessel vasculitides through a meta-immunochip strategy. <i>Scientific Reports</i> , 2017, 7, 43953.	1.6	52
135	The Erythrocyte Sedimentation Rate Is Associated with the Development of Visual Complications in Biopsy-Proven Giant Cell Arteritis. <i>Seminars in Arthritis and Rheumatism</i> , 2008, 38, 116-123.	1.6	51
136	Novel Association of the Interleukin 23 Region With Inflammatory Bowel Disease. <i>American Journal of Gastroenterology</i> , 2009, 104, 1968-1975.	0.2	51
137	Evaluation of TRAF6 in a large multiethnic lupus cohort. <i>Arthritis and Rheumatism</i> , 2012, 64, 1960-1969.	6.7	51
138	Identification of the PTPN22 functional variant R620W as susceptibility genetic factor for giant cell arteritis. <i>Annals of the Rheumatic Diseases</i> , 2013, 72, 1882-1886.	0.5	51
139	Macrophage migration inhibitory factor gene: Influence on rheumatoid arthritis susceptibility. <i>Human Immunology</i> , 2007, 68, 744-747.	1.2	50
140	Genetic association of vasoactive intestinal peptide receptor with rheumatoid arthritis: Altered expression and signal in immune cells. <i>Arthritis and Rheumatism</i> , 2008, 58, 1010-1019.	6.7	50
141	Study of functional variants of the BANK1 gene in rheumatoid arthritis. <i>Arthritis and Rheumatism</i> , 2009, 60, 372-379.	6.7	50
142	Specific association of a CLEC16A/KIAA0350 polymorphism with NOD2/CARD15 Crohn's disease patients. <i>European Journal of Human Genetics</i> , 2009, 17, 1304-1308.	1.4	50
143	Association of PTPN22 gene functional variants with development of pulmonary tuberculosis in Moroccan population. <i>Tissue Antigens</i> , 2009, 74, 228-232.	1.0	50
144	Influence of HLA DRB1 alleles in the susceptibility of rheumatoid arthritis and the regulation of antibodies against citrullinated proteins and rheumatoid factor. <i>Arthritis Research and Therapy</i> , 2010, 12, R62.	1.6	50

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