

# Robert P Kimberly

## List of Publications by Year in Descending Order

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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.

193  
papers

15,052  
citations

66  
h-index

118  
g-index

202  
ext. papers

16,856  
ext. citations

9.5  
avg, IF

5.57  
L-index

#	Paper	IF	Citations
193	Exploring COVID-19 Vaccine Hesitancy Among Stakeholders in African American and Latinx Communities in the Deep South Through the Lens of the Health Belief Model. <i>American Journal of Health Promotion</i> , <b>2021</b> , 8901171211045038	2.5	0
192	Returning to Growth: One Academic Medical Center's Successful Five-Step Approach to Change Management. <i>Academic Medicine</i> , <b>2021</b> , 96, 1401-1407	3.9	1
191	Undiagnosed SARS-CoV-2 seropositivity during the first 6 months of the COVID-19 pandemic in the United States. <i>Science Translational Medicine</i> , <b>2021</b> , 13,	17.5	34
190	Fine-mapping, trans-ancestral and genomic analyses identify causal variants, cells, genes and drug targets for type 1 diabetes. <i>Nature Genetics</i> , <b>2021</b> , 53, 962-971	36.3	28
189	Re-engineering The Clinical Research Enterprise in Response to COVID-19: The Clinical Translational Science Award (CTSA) experience and proposed playbook for future pandemics. <i>Journal of Clinical and Translational Science</i> , <b>2021</b> , 5, e96	0.4	2
188	Mapping a Pandemic: SARS-CoV-2 Seropositivity in the United States <b>2021</b> ,		11
187	Complement genes contribute sex-biased vulnerability in diverse disorders. <i>Nature</i> , <b>2020</b> , 582, 577-581	50.4	71
186	Skewed allelic expression on X chromosome associated with aberrant expression of XIST on systemic lupus erythematosus lymphocytes. <i>Human Molecular Genetics</i> , <b>2020</b> , 29, 2523-2534	5.6	7
185	MRLR: unraveling high-resolution meiotic recombination by linked reads. <i>Bioinformatics</i> , <b>2020</b> , 36, 10-16	7.2	1
184	Population-Specific Patterns of Epigenetic Defects in the B Cell Lineage in Patients With Systemic Lupus Erythematosus. <i>Arthritis and Rheumatology</i> , <b>2020</b> , 72, 282-291	9.5	14
183	Genetic influences on susceptibility to rheumatoid arthritis in African-Americans. <i>Human Molecular Genetics</i> , <b>2019</b> , 28, 858-874	5.6	26
182	Type 1 Diabetes Risk in African-Ancestry Participants and Utility of an Ancestry-Specific Genetic Risk Score. <i>Diabetes Care</i> , <b>2019</b> , 42, 406-415	14.6	39
181	Individualized decision aid for diverse women with lupus nephritis (IDEA-WON): A randomized controlled trial. <i>PLoS Medicine</i> , <b>2019</b> , 16, e1002800	11.6	13
180	Association of response to TNF inhibitors in rheumatoid arthritis with quantitative trait loci for and CD39. <i>Annals of the Rheumatic Diseases</i> , <b>2019</b> , 78, 1055-1061	2.4	12
179	Genetic variants at the 16p13 locus confer risk for eosinophilic esophagitis. <i>Genes and Immunity</i> , <b>2019</b> , 20, 281-292	4.4	16
178	IFN $\gamma$ induces epigenetic programming of human T-bet B cells and promotes TLR7/8 and IL-21 induced differentiation. <i>ELife</i> , <b>2019</b> , 8,	8.9	65
177	Discovery and Validation of Circulating Biomarkers of Colorectal Adenoma by High-Depth Small RNA Sequencing. <i>Clinical Cancer Research</i> , <b>2018</b> , 24, 2092-2099	12.9	17

176	A plausibly causal functional lupus-associated risk variant in the STAT1-STAT4 locus. <i>Human Molecular Genetics</i> , <b>2018</b> , 27, 2392-2404	5.6	22
175	Genetic fine mapping of systemic lupus erythematosus MHC associations in Europeans and African Americans. <i>Human Molecular Genetics</i> , <b>2018</b> , 27, 3813-3824	5.6	23
174	Trans-Ethnic Mapping of Identifies Two Independent SLE-Risk Linkage Groups Enriched for Co-Transcriptional Splicing Marks. <i>International Journal of Molecular Sciences</i> , <b>2018</b> , 19,	6.3	9
173	Genome-wide association studies suggest that APOL1-environment interactions more likely trigger kidney disease in African Americans with nondiabetic nephropathy than strong APOL1-second gene interactions. <i>Kidney International</i> , <b>2018</b> , 94, 599-607	9.9	30
172	Transancestral mapping and genetic load in systemic lupus erythematosus. <i>Nature Communications</i> , <b>2017</b> , 8, 16021	17.4	171
171	Regulation of FcRIIb function by site-specific serine phosphorylation. <i>Journal of Leukocyte Biology</i> , <b>2017</b> , 101, 421-428	6.5	3
170	Preferential association of a functional variant in complement receptor 2 with antibodies to double-stranded DNA. <i>Annals of the Rheumatic Diseases</i> , <b>2016</b> , 75, 242-52	2.4	8
169	Decreased SMG7 expression associates with lupus-risk variants and elevated antinuclear antibody production. <i>Annals of the Rheumatic Diseases</i> , <b>2016</b> , 75, 2007-2013	2.4	12
168	X Chromosome Dose and Sex Bias in Autoimmune Diseases: Increased Prevalence of 47,XXX in Systemic Lupus Erythematosus and Sjögren's Syndrome. <i>Arthritis and Rheumatology</i> , <b>2016</b> , 68, 1290-1300	9.5	65
167	Identification of a Systemic Lupus Erythematosus Risk Locus Spanning ATG16L2, FCHSD2, and P2RY2 in Koreans. <i>Arthritis and Rheumatology</i> , <b>2016</b> , 68, 1197-1209	9.5	45
166	Estimation of Cell-Type Composition Including T and B Cell Subtypes for Whole Blood Methylation Microarray Data. <i>Frontiers in Genetics</i> , <b>2016</b> , 7, 23	4.5	14
165	Genome-Wide Association Study in an Amerindian Ancestry Population Reveals Novel Systemic Lupus Erythematosus Risk Loci and the Role of European Admixture. <i>Arthritis and Rheumatology</i> , <b>2016</b> , 68, 932-43	9.5	93
164	Clinical associations of anti-Smith antibodies in PROFILE: a multi-ethnic lupus cohort. <i>Clinical Rheumatology</i> , <b>2015</b> , 34, 1217-23	3.9	37
163	Lupus Risk Variant Increases pSTAT1 Binding and Decreases ETS1 Expression. <i>American Journal of Human Genetics</i> , <b>2015</b> , 96, 731-9	11	31
162	Genetic associations of leptin-related polymorphisms with systemic lupus erythematosus. <i>Clinical Immunology</i> , <b>2015</b> , 161, 157-62	9	8
161	Characterization of genetic loci that affect susceptibility to inflammatory bowel diseases in African Americans. <i>Gastroenterology</i> , <b>2015</b> , 149, 1575-1586	13.3	47
160	Blocking of targeted microRNAs from next-generation sequencing libraries. <i>Nucleic Acids Research</i> , <b>2015</b> , 43, e145	20.1	14
159	The IRF5-TNPO3 association with systemic lupus erythematosus has two components that other autoimmune disorders variably share. <i>Human Molecular Genetics</i> , <b>2015</b> , 24, 582-96	5.6	57

158	Lack of Association of the APOL1 G3 Haplotype in African Americans with ESRD. <i>Journal of the American Society of Nephrology: JASN</i> , <b>2015</b> , 26, 1021-5	12.7	4
157	Lupus risk variants in the PTK2B locus alter B-cell receptor internalization. <i>Frontiers in Genetics</i> , <b>2014</b> , 5, 450	4.5	22
156	Targeting the Fc receptor in autoimmune disease. <i>Expert Opinion on Therapeutic Targets</i> , <b>2014</b> , 18, 335-50	5.4	43
155	A polymorphism in TLR2 is associated with arterial thrombosis in a multiethnic population of patients with systemic lupus erythematosus. <i>Arthritis and Rheumatology</i> , <b>2014</b> , 66, 1882-7	9.5	17
154	Lupus nephritis susceptibility loci in women with systemic lupus erythematosus. <i>Journal of the American Society of Nephrology: JASN</i> , <b>2014</b> , 25, 2859-70	12.7	83
153	Fcγ receptor IIIa single-nucleotide polymorphisms and haplotypes affect human IgG binding and are associated with lupus nephritis in African Americans. <i>Arthritis and Rheumatology</i> , <b>2014</b> , 66, 1291-9	9.5	23
152	Human FcR polymorphism and disease. <i>Current Topics in Microbiology and Immunology</i> , <b>2014</b> , 382, 275-303	3.3	27
151	End-stage renal disease in African Americans with lupus nephritis is associated with APOL1. <i>Arthritis and Rheumatology</i> , <b>2014</b> , 66, 390-6	9.5	174
150	Two functional lupus-associated BLK promoter variants control cell-type- and developmental-stage-specific transcription. <i>American Journal of Human Genetics</i> , <b>2014</b> , 94, 586-98	11	49
149	Allelic-dependent expression of an activating Fc receptor on B cells enhances humoral immune responses. <i>Science Translational Medicine</i> , <b>2013</b> , 5, 216ra175	17.5	41
148	Brief Report: Single-nucleotide polymorphisms in VKORC1 are risk factors for systemic lupus erythematosus in Asians. <i>Arthritis and Rheumatism</i> , <b>2013</b> , 65, 211-5		8
147	Variable association of reactive intermediate genes with systemic lupus erythematosus in populations with different African ancestry. <i>Journal of Rheumatology</i> , <b>2013</b> , 40, 842-9	4.1	12
146	Preferential binding to Elk-1 by SLE-associated IL10 risk allele upregulates IL10 expression. <i>PLoS Genetics</i> , <b>2013</b> , 9, e1003870	6	33
145	Genome-wide association study and gene expression analysis identifies CD84 as a predictor of response to etanercept therapy in rheumatoid arthritis. <i>PLoS Genetics</i> , <b>2013</b> , 9, e1003394	6	127
144	Genome-wide DNA methylation analysis of systemic lupus erythematosus reveals persistent hypomethylation of interferon genes and compositional changes to CD4+ T-cell populations. <i>PLoS Genetics</i> , <b>2013</b> , 9, e1003678	6	241
143	Admixture mapping in lupus identifies multiple functional variants within IFIH1 associated with apoptosis, inflammation, and autoantibody production. <i>PLoS Genetics</i> , <b>2013</b> , 9, e1003222	6	87
142	Trans-ancestral studies fine map the SLE-susceptibility locus TNFSF4. <i>PLoS Genetics</i> , <b>2013</b> , 9, e1003554	6	41
141	MicroRNA-3148 modulates allelic expression of toll-like receptor 7 variant associated with systemic lupus erythematosus. <i>PLoS Genetics</i> , <b>2013</b> , 9, e1003336	6	89

140	Fine mapping of Xq28: both MECP2 and IRAK1 contribute to risk for systemic lupus erythematosus in multiple ancestral groups. <i>Annals of the Rheumatic Diseases</i> , <b>2013</b> , 72, 437-44	2.4	80
139	ABIN1 dysfunction as a genetic basis for lupus nephritis. <i>Journal of the American Society of Nephrology: JASN</i> , <b>2013</b> , 24, 1743-54	12.7	59
138	Abnormalities in Immune Complex Clearance and Fcγ Receptor Function <b>2013</b> , 127-140		1
137	Genomic copy number variants: evidence for association with antibody response to anthrax vaccine adsorbed. <i>PLoS ONE</i> , <b>2013</b> , 8, e64813	3.7	5
136	PTPN22 association in systemic lupus erythematosus (SLE) with respect to individual ancestry and clinical sub-phenotypes. <i>PLoS ONE</i> , <b>2013</b> , 8, e69404	3.7	48
135	Evidence for gene-gene epistatic interactions among susceptibility loci for systemic lupus erythematosus. <i>Arthritis and Rheumatism</i> , <b>2012</b> , 64, 485-92		47
134	Impact of genetic ancestry and sociodemographic status on the clinical expression of systemic lupus erythematosus in American Indian-European populations. <i>Arthritis and Rheumatism</i> , <b>2012</b> , 64, 3687-94		61
133	Association of two independent functional risk haplotypes in TNIP1 with systemic lupus erythematosus. <i>Arthritis and Rheumatism</i> , <b>2012</b> , 64, 3695-705		64
132	Variation in the ICAM1-ICAM4-ICAM5 locus is associated with systemic lupus erythematosus susceptibility in multiple ancestries. <i>Annals of the Rheumatic Diseases</i> , <b>2012</b> , 71, 1809-14	2.4	51
131	A genome-wide association study of host genetic determinants of the antibody response to Anthrax Vaccine Adsorbed. <i>Vaccine</i> , <b>2012</b> , 30, 4778-84	4.1	21
130	Genome-wide association study of African and European Americans implicates multiple shared and ethnic specific loci in sarcoidosis susceptibility. <i>PLoS ONE</i> , <b>2012</b> , 7, e43907	3.7	76
129	Treatment of arthritis by macrophage depletion and immunomodulation: testing an apoptosis-mediated therapy in a humanized death receptor mouse model. <i>Arthritis and Rheumatism</i> , <b>2012</b> , 64, 1098-109		45
128	Association of single-nucleotide polymorphisms in CCR6, TAGAP, and TNFAIP3 with rheumatoid arthritis in African Americans. <i>Arthritis and Rheumatism</i> , <b>2012</b> , 64, 1355-8		15
127	Evaluation of TRAF6 in a large multi-ancestral lupus cohort. <i>Arthritis and Rheumatism</i> , <b>2012</b> , 64, 1960-9		45
126	Genetic risk factors for thrombosis in systemic lupus erythematosus. <i>Journal of Rheumatology</i> , <b>2012</b> , 39, 1603-10	4.1	19
125	Identification of IRF8, TMEM39A, and IKZF3-ZBP2 as susceptibility loci for systemic lupus erythematosus in a large-scale multi-racial replication study. <i>American Journal of Human Genetics</i> , <b>2012</b> , 90, 648-60	11	134
124	The unique cytoplasmic domain of human FcγRIIIa regulates receptor-mediated function. <i>Journal of Immunology</i> , <b>2012</b> , 189, 4284-94	5.3	14
123	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> , <b>2012</b> , 71, 694-9	2.4	63

122	Serine phosphorylation of Fc $\gamma$ RI cytoplasmic domain directs lipid raft localization and interaction with protein 4.1G. <i>Journal of Leukocyte Biology</i> , <b>2012</b> , 91, 97-103	6.5	7
121	CSK regulatory polymorphism is associated with systemic lupus erythematosus and influences B-cell signaling and activation. <i>Nature Genetics</i> , <b>2012</b> , 44, 1227-30	36.3	87
120	Lupus-associated causal mutation in neutrophil cytosolic factor 2 (NCF2) brings unique insights to the structure and function of NADPH oxidase. <i>Proceedings of the National Academy of Sciences of the United States of America</i> , <b>2012</b> , 109, E59-67	11.5	124
119	IRF5 haplotypes demonstrate diverse serological associations which predict serum interferon alpha activity and explain the majority of the genetic association with systemic lupus erythematosus. <i>Annals of the Rheumatic Diseases</i> , <b>2012</b> , 71, 463-8	2.4	109
118	Association of a functional variant downstream of TNFAIP3 with systemic lupus erythematosus. <i>Nature Genetics</i> , <b>2011</b> , 43, 253-8	36.3	208
117	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> , <b>2011</b> , 88, 83-91	11	69
116	FAS mRNA editing in Human Systemic Lupus Erythematosus. <i>Human Mutation</i> , <b>2011</b> , 32, 1268-77	4.7	9
115	Fine-mapping and transethnic genotyping establish IL2/IL21 genetic association with lupus and localize this genetic effect to IL21. <i>Arthritis and Rheumatism</i> , <b>2011</b> , 63, 1689-97		41
114	Genetic analyses of interferon pathway-related genes reveal multiple new loci associated with systemic lupus erythematosus. <i>Arthritis and Rheumatism</i> , <b>2011</b> , 63, 2049-57		39
113	Association of PPP2CA polymorphisms with systemic lupus erythematosus susceptibility in multiple ethnic groups. <i>Arthritis and Rheumatism</i> , <b>2011</b> , 63, 2755-63		31
112	Identification of novel genetic susceptibility loci in African American lupus patients in a candidate gene association study. <i>Arthritis and Rheumatism</i> , <b>2011</b> , 63, 3493-501		86
111	IgA and IgG antineutrophil cytoplasmic antibody engagement of Fc receptor genetic variants influences granulomatosis with polyangiitis. <i>Proceedings of the National Academy of Sciences of the United States of America</i> , <b>2011</b> , 108, 20736-41	11.5	58
110	A comprehensive analysis of shared loci between systemic lupus erythematosus (SLE) and sixteen autoimmune diseases reveals limited genetic overlap. <i>PLoS Genetics</i> , <b>2011</b> , 7, e1002406	6	126
109	The functional polymorphism 844 A>G in Fc $\gamma$ RI (CD89) does not contribute to systemic sclerosis or rheumatoid arthritis susceptibility. <i>Journal of Rheumatology</i> , <b>2011</b> , 38, 446-9	4.1	4
108	Phenotypic associations of genetic susceptibility loci in systemic lupus erythematosus. <i>Annals of the Rheumatic Diseases</i> , <b>2011</b> , 70, 1752-7	2.4	89
107	Association of genetic variants in complement factor H and factor H-related genes with systemic lupus erythematosus susceptibility. <i>PLoS Genetics</i> , <b>2011</b> , 7, e1002079	6	145
106	Human FasL gene is a target of $\beta$ -catenin/T-cell factor pathway and complex FasL haplotypes alter promoter functions. <i>PLoS ONE</i> , <b>2011</b> , 6, e26143	3.7	10
105	Genetic variants of STAT4 associated with rheumatoid arthritis in persons of Asian and European ancestry do not replicate in African Americans. <i>Annals of the Rheumatic Diseases</i> , <b>2010</b> , 69, 625-6	2.4	19



104	The role of genetic variation near interferon-kappa in systemic lupus erythematosus. <i>Journal of Biomedicine and Biotechnology</i> , <b>2010</b> , 2010,		37
103	ITGAM coding variant (rs1143679) influences the risk of renal disease, discoid rash and immunological manifestations in patients with systemic lupus erythematosus with European ancestry. <i>Annals of the Rheumatic Diseases</i> , <b>2010</b> , 69, 1329-32	2.4	60
102	The non-muscle Myosin heavy chain 9 gene (MYH9) is not associated with lupus nephritis in African Americans. <i>American Journal of Nephrology</i> , <b>2010</b> , 32, 66-72	4.6	15
101	Genetic factors predisposing to systemic lupus erythematosus and lupus nephritis. <i>Seminars in Nephrology</i> , <b>2010</b> , 30, 164-76	4.8	66
100	Pathways: Strategies for susceptibility genes in SLE. <i>Autoimmunity Reviews</i> , <b>2010</b> , 9, 473-6	13.6	26
99	Wegener's granulomatosis: a model of auto-antibodies in mucosal autoimmunity. <i>Clinical Immunology</i> , <b>2010</b> , 134, 104-12	9	5
98	Most common single-nucleotide polymorphisms associated with rheumatoid arthritis in persons of European ancestry confer risk of rheumatoid arthritis in African Americans. <i>Arthritis and Rheumatism</i> , <b>2010</b> , 62, 3547-53		47
97	An African ancestry-specific allele of CTLA4 confers protection against rheumatoid arthritis in African Americans. <i>PLoS Genetics</i> , <b>2009</b> , 5, e1000424	6	9
96	Evaluation of imputation-based association in and around the integrin-alpha-M (ITGAM) gene and replication of robust association between a non-synonymous functional variant within ITGAM and systemic lupus erythematosus (SLE). <i>Human Molecular Genetics</i> , <b>2009</b> , 18, 1171-80	5.6	88
95	Variants within MECP2, a key transcription regulator, are associated with increased susceptibility to lupus and differential gene expression in patients with systemic lupus erythematosus. <i>Arthritis and Rheumatism</i> , <b>2009</b> , 60, 1076-84		69
94	A polymorphism within IL21R confers risk for systemic lupus erythematosus. <i>Arthritis and Rheumatism</i> , <b>2009</b> , 60, 2402-7		97
93	The FCRL3-169CT promoter single-nucleotide polymorphism, which is associated with systemic lupus erythematosus in a Japanese population, predicts expression of receptor protein on CD19+ B cells. <i>Arthritis and Rheumatism</i> , <b>2009</b> , 60, 3510-2		20
92	A large-scale replication study identifies TNIP1, PRDM1, JAZF1, UHRF1BP1 and IL10 as risk loci for systemic lupus erythematosus. <i>Nature Genetics</i> , <b>2009</b> , 41, 1228-33	36.3	626
91	Kallikrein genes are associated with lupus and glomerular basement membrane-specific antibody-induced nephritis in mice and humans. <i>Journal of Clinical Investigation</i> , <b>2009</b> , 119, 911-23	15.9	100
90	A nonsynonymous functional variant in integrin-alpha(M) (encoded by ITGAM) is associated with systemic lupus erythematosus. <i>Nature Genetics</i> , <b>2008</b> , 40, 152-4	36.3	247
89	Genome-wide association scan in women with systemic lupus erythematosus identifies susceptibility variants in ITGAM, PTK, KIAA1542 and other loci. <i>Nature Genetics</i> , <b>2008</b> , 40, 204-10	36.3	1021
88	Genetic variation in the CRP promoter: association with systemic lupus erythematosus. <i>Human Molecular Genetics</i> , <b>2008</b> , 17, 1147-55	5.6	52
87	Functional expression of IgA receptor FcalphaRI on human platelets. <i>Journal of Leukocyte Biology</i> , <b>2008</b> , 84, 1492-500	6.5	48

86	Immune opsonins modulate BLyS/BAFF release in a receptor-specific fashion. <i>Journal of Immunology</i> , <b>2008</b> , 181, 1012-8	5.3	23
85	Klinefelter's syndrome (47,XXY) in male systemic lupus erythematosus patients: support for the notion of a gene-dose effect from the X chromosome. <i>Arthritis and Rheumatism</i> , <b>2008</b> , 58, 2511-7		221
84	Features associated with, and the impact of, hemolytic anemia in patients with systemic lupus erythematosus: LX, results from a multiethnic cohort. <i>Arthritis and Rheumatism</i> , <b>2008</b> , 59, 1332-40		27
83	IgA1-secreting cell lines from patients with IgA nephropathy produce aberrantly glycosylated IgA1. <i>Journal of Clinical Investigation</i> , <b>2008</b> , 118, 629-39	15.9	178
82	Common variants within MECP2 confer risk of systemic lupus erythematosus. <i>PLoS ONE</i> , <b>2008</b> , 3, e1727	3.7	112
81	Genetics and Disease <b>2008</b> , 108-113		
80	A genome wide association study of systemic lupus erythematosus (SLE) by SLEGEN, the International SLE Genetics Consortium.. <i>FASEB Journal</i> , <b>2008</b> , 22, 850.1	0.9	
79	Variation in the relative copy number of the TLR7 gene in patients with systemic lupus erythematosus and healthy control subjects. <i>Arthritis and Rheumatism</i> , <b>2007</b> , 56, 3375-8		52
78	Biologic therapies in rheumatology: lessons learned, future directions. <i>Nature Reviews Drug Discovery</i> , <b>2007</b> , 6, 75-92	64.1	141
77	Curcumin reverses breast tumor exosomes mediated immune suppression of NK cell tumor cytotoxicity. <i>Biochimica Et Biophysica Acta - Molecular Cell Research</i> , <b>2007</b> , 1773, 1116-23	4.9	105
76	Fc receptor genes and the systemic lupus erythematosus diathesis. <i>Autoimmunity</i> , <b>2007</b> , 40, 567-81	3	54
75	Fc $\alpha$ RI (CD89) alleles determine the proinflammatory potential of serum IgA. <i>Journal of Immunology</i> , <b>2007</b> , 178, 3973-82	5.3	48
74	Expression profile of Fc $\gamma$ RIIb on leukocytes and its dysregulation in systemic lupus erythematosus. <i>Journal of Immunology</i> , <b>2007</b> , 178, 3272-80	5.3	144
73	Association of a transmembrane polymorphism of Fc $\gamma$ receptor IIb (FCGR2B) with systemic lupus erythematosus in Taiwanese patients. <i>Arthritis and Rheumatism</i> , <b>2006</b> , 54, 3908-17		59
72	Time to renal disease and end-stage renal disease in PROFILE: a multiethnic lupus cohort. <i>PLoS Medicine</i> , <b>2006</b> , 3, e396	11.6	75
71	Regional admixture mapping and structured association testing: conceptual unification and an extensible general linear model. <i>PLoS Genetics</i> , <b>2006</b> , 2, e137	6	53
70	Dectin-2 is a pattern recognition receptor for fungi that couples with the Fc receptor gamma chain to induce innate immune responses. <i>Journal of Biological Chemistry</i> , <b>2006</b> , 281, 38854-66	5.4	325
69	Inducible resistance of tumor cells to tumor necrosis factor-related apoptosis-inducing ligand receptor 2-mediated apoptosis by generation of a blockade at the death domain function. <i>Cancer Research</i> , <b>2006</b> , 66, 8520-8	10.1	54



68	Murine mammary carcinoma exosomes promote tumor growth by suppression of NK cell function. <i>Journal of Immunology</i> , <b>2006</b> , 176, 1375-85	5.3	380
67	No evidence for an association between the -871 T/C promoter polymorphism in the B-cell-activating factor gene and primary Sjögren's syndrome. <i>Arthritis Research and Therapy</i> , <b>2006</b> , 8, R30	5.7	31
66	JAB1 determines the response of rheumatoid arthritis synovial fibroblasts to tumor necrosis factor-alpha. <i>American Journal of Pathology</i> , <b>2006</b> , 169, 889-902	5.8	22
65	SLE: challenges and candidates in human disease. <i>Trends in Immunology</i> , <b>2005</b> , 26, 580-6	14.4	50
64	Cleavage of p53-vimentin complex enhances tumor necrosis factor-related apoptosis-inducing ligand-mediated apoptosis of rheumatoid arthritis synovial fibroblasts. <i>American Journal of Pathology</i> , <b>2005</b> , 167, 705-19	5.8	48
63	Genetics of susceptibility and severity in systemic lupus erythematosus. <i>Current Opinion in Rheumatology</i> , <b>2005</b> , 17, 529-37	5.3	53
62	Expression and occupancy of BAFF-R on B cells in systemic lupus erythematosus. <i>Arthritis and Rheumatism</i> , <b>2005</b> , 52, 3943-54		85
61	A promoter haplotype of the immunoreceptor tyrosine-based inhibitory motif-bearing FcgammaRIIb alters receptor expression and associates with autoimmunity. I. Regulatory FCGR2B polymorphisms and their association with systemic lupus erythematosus. <i>Journal of Immunology</i> , <b>2004</b> , 172, 7192-9	5.3	141
60	A promoter haplotype of the immunoreceptor tyrosine-based inhibitory motif-bearing FcgammaRIIb alters receptor expression and associates with autoimmunity. II. Differential binding of GATA4 and Yin-Yang1 transcription factors and correlated receptor expression and function. <i>Journal of Immunology</i> , <b>2004</b> , 172, 7192-9	5.3	92
59	Differential gene expression modulated by the cytoplasmic domain of Fc gamma RIa (CD64) alpha-chain. <i>Journal of Immunology</i> , <b>2004</b> , 173, 6211-9	5.3	8
58	An exposed domain in the severe acute respiratory syndrome coronavirus spike protein induces neutralizing antibodies. <i>Journal of Virology</i> , <b>2004</b> , 78, 7217-26	6.6	71
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1	Complement component 4 genes contribute sex-specific vulnerability in diverse illnesses		2