

STAT4 and the Risk of Rheumatoid Arthritis and

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Citation Report

#	ARTICLE	IF	CITATIONS
1	Rheumatoid Arthritis, Systemic Lupus Erythematosus, and <i>STAT4</i> . New England Journal of Medicine, 2007, 357, 2517-2518.	27.0	7
2	<i>TRAF1</i> and <i>C5</i> as a Risk Locus for Rheumatoid Arthritis – A Genomewide Study. New England Journal of Medicine, 2007, 357, 1199-1209.	27.0	729
3	Recent advances in the genetics of RA susceptibility. Rheumatology, 2007, 47, 399-402.	1.9	138
4	Mechanisms of Disease: genetics of rheumatoid arthritis – ethnic differences in disease-associated genes. Nature Clinical Practice Rheumatology, 2007, 3, 644-650.	3.2	33
6	Association of <i>STAT4</i> with Rheumatoid Arthritis in the Korean Population. Molecular Medicine, 2007, 13, 455-460.	4.4	151
8	Contribution of genetic studies in rodent models of autoimmune arthritis to understanding and treatment of rheumatoid arthritis. Genes and Immunity, 2007, 8, 523-531.	4.1	9
9	HSPD1 is not a major susceptibility gene for rheumatoid arthritis in the French Caucasian population. Journal of Human Genetics, 2007, 52, 1036-1039.	2.3	3
10	<i>STAT4</i> : Genetics, mechanisms, and implications for autoimmunity. Current Allergy and Asthma Reports, 2008, 8, 398-403.	5.3	158
11	The burden of rheumatoid arthritis and access to treatment: a medical overview. European Journal of Health Economics, 2008, 8, 39-47.	2.8	20
12	<i>IL23R</i> and <i>IL12B</i> polymorphisms in spanish IBD patients: No evidence of interaction. Inflammatory Bowel Diseases, 2008, 14, 1192-1196.	1.9	38
13	Human leukocyte antigen polymorphisms in italian primary biliary cirrhosis: A multicenter study of 664 patients and 1992 healthy controls. Hepatology, 2008, 48, 1906-1912.	7.3	120
14	The inflammatory disease – associated variants in <i>IL12B</i> and <i>IL23R</i> are not associated with rheumatoid arthritis. Arthritis and Rheumatism, 2008, 58, 1877-1881.	6.7	41
15	Association of <i>STAT4</i> with susceptibility to rheumatoid arthritis and systemic lupus erythematosus in the Japanese population. Arthritis and Rheumatism, 2008, 58, 1940-1946.	6.7	139
16	Association of <i>STAT4</i> with rheumatoid arthritis: A replication study in three European populations. Arthritis and Rheumatism, 2008, 58, 1974-1980.	6.7	93
17	A broad analysis of <i>IL1</i> polymorphism and rheumatoid arthritis. Arthritis and Rheumatism, 2008, 58, 1947-1957.	6.7	31
18	Genome-wide association study of rheumatoid arthritis in the Spanish population: <i>KLF12</i> as a risk locus for rheumatoid arthritis susceptibility. Arthritis and Rheumatism, 2008, 58, 2275-2286.	6.7	100
19	Association of the <i>STAT4</i> gene with increased susceptibility for some immune-mediated diseases. Arthritis and Rheumatism, 2008, 58, 2598-2602.	6.7	118
20	Replication of the tumor necrosis factor receptor-associated factor 1/complement component 5 region as a susceptibility locus for rheumatoid arthritis in a European family-based study. Arthritis and Rheumatism, 2008, 58, 2670-2674.	6.7	32

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21	Vitamin E in the primary prevention of rheumatoid arthritis: The women's health study. Arthritis and Rheumatism, 2008, 59, 1589-1595.	6.7	53
22	STAT4 Polymorphism Is Associated with Early-Onset Type 1 Diabetes, but Not with Late-Onset Type 1 Diabetes. Annals of the New York Academy of Sciences, 2008, 1150, 93-98.	3.8	38
23	Th17 cells in human disease. Immunological Reviews, 2008, 223, 87-113.	6.0	960
24	How the study of children with rheumatic diseases identified interferon- γ and interleukin-1 as novel therapeutic targets. Immunological Reviews, 2008, 223, 39-59.	6.0	68
25	Molecular aspects of rheumatoid arthritis: role of transcription factors. FEBS Journal, 2008, 275, 4463-4470.	4.7	76
26	Variant form of STAT4 is associated with primary Sjögren's syndrome. Genes and Immunity, 2008, 9, 267-270.	4.1	165
27	STAT4 but not TRAF1/C5 variants influence the risk of developing rheumatoid arthritis and systemic lupus erythematosus in Colombians. Genes and Immunity, 2008, 9, 379-382.	4.1	86
28	The role and therapeutic implications of fibroblast-like synoviocytes in inflammation and cartilage erosion in rheumatoid arthritis. Immunological Reviews, 2008, 223, 252-270.	6.0	294
29	Genetic variants near TNFAIP3 on 6q23 are associated with systemic lupus erythematosus. Nature Genetics, 2008, 40, 1059-1061.	21.4	534
30	Functional SNPs in CD244 increase the risk of rheumatoid arthritis in a Japanese population. Nature Genetics, 2008, 40, 1224-1229.	21.4	106
31	Common variants at CD40 and other loci confer risk of rheumatoid arthritis. Nature Genetics, 2008, 40, 1216-1223.	21.4	476
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33	The developing mosaic of autoimmune disease risk. Nature Genetics, 2008, 40, 131-132.	21.4	13
34	Hypoxic reprogramming. Nature Genetics, 2008, 40, 132-134.	21.4	19
35	The genetics and immunopathogenesis of inflammatory bowel disease. Nature Reviews Immunology, 2008, 8, 458-466.	22.7	819
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37	Autoimmunity since the 1957 clonal selection theory: a little acorn to a large oak. Immunology and Cell Biology, 2008, 86, 67-71.	2.3	21
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40	Immune-mediated pathways in chronic inflammatory arthritis. <i>Best Practice and Research in Clinical Rheumatology</i> , 2008, 22, 221-238.	3.3	39
41	CCL genes in multiple sclerosis and systemic lupus erythematosus. <i>Journal of Neuroimmunology</i> , 2008, 200, 145-152.	2.3	21
42	Specificity of the STAT4 Genetic Association for Severe Disease Manifestations of Systemic Lupus Erythematosus. <i>PLoS Genetics</i> , 2008, 4, e1000084.	3.5	180
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44	Does gamma-aminobutyric acid (GABA) influence the development of chronic inflammation in rheumatoid arthritis?. <i>Journal of Neuroinflammation</i> , 2008, 5, 1.	7.2	99
45	Role of STAT4 polymorphisms in systemic lupus erythematosus in a Japanese population: a case-control association study of the STAT1-STAT4 region. <i>Arthritis Research and Therapy</i> , 2008, 10, R113.	3.5	88
46	High avidity autoreactive T cells with a low signalling capacity through the T-cell receptor: central to rheumatoid arthritis pathogenesis?. <i>Arthritis Research and Therapy</i> , 2008, 10, 210.	3.5	25
47	Circulating immune complexes contain citrullinated fibrinogen in rheumatoid arthritis. <i>Arthritis Research and Therapy</i> , 2008, 10, R94.	3.5	157
48	Genetic polymorphisms in PTPN22, PADI-4, and CTLA-4 and risk for rheumatoid arthritis in two longitudinal cohort studies: evidence of gene-environment interactions with heavy cigarette smoking. <i>Arthritis Research and Therapy</i> , 2008, 10, R52.	3.5	81
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52	Association of a TRAF1 and a STAT4 gene polymorphism with increased risk for rheumatoid arthritis in a genetically homogeneous population. <i>Human Immunology</i> , 2008, 69, 567-571.	2.4	75
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58	Genes and Sjögren's Syndrome. Rheumatic Disease Clinics of North America, 2008, 34, 847-868.	1.9	50
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60	Collaboration, Genetic Associations, and Lupus Erythematosus. New England Journal of Medicine, 2008, 358, 956-961.	27.0	86
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71	Primer: SNP-associated studies and what they can teach us. Nature Clinical Practice Rheumatology, 2008, 4, 210-217.	3.2	11
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74	SNP-guided microRNA maps (MirMaps) of 16 common human disorders identify a clinically accessible therapy reversing transcriptional aberrations of nuclear import and inflammasome pathways. Cell Cycle, 2008, 7, 3564-3576.	2.6	60
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82	STAT4 and the Risk of Rheumatoid Arthritis and Systemic Lupus Erythematosus. Yearbook of Dermatology and Dermatologic Surgery, 2008, 2008, 222-223.	0.0	0
83	TRAF1-C5 as a Risk Locus for Rheumatoid ArthritisA Genomewide Study. Yearbook of Dermatology and Dermatologic Surgery, 2008, 2008, 220-222.	0.0	0
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87	ç¬Œ2âŹé–Çç€Œfªã, ãfžãfã•HLA. Major Histocompatibility Complex, 2008, 15, 235-248.	0.1	0
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94	Systemic Lupus Erythematosus Features in Rheumatoid Arthritis and Their Effect on Overall Mortality. Journal of Rheumatology, 2009, 36, 50-57.	2.0	39

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96	Recent advances and opportunities in research on lupus: environmental influences and mechanisms of disease. Ciencia E Saude Coletiva, 2009, 14, 1865-1876.	0.5	8
97	Deconvolution of Blood Microarray Data Identifies Cellular Activation Patterns in Systemic Lupus Erythematosus. PLoS ONE, 2009, 4, e6098.	2.5	388
98	Association of STAT4 polymorphism with systemic sclerosis in a Japanese population. Annals of the Rheumatic Diseases, 2009, 68, 1375-1376.	0.9	64
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100	Cigarette Smoking, <i>STAT4</i> and <i>TNFRSF1B</i> Polymorphisms, and Systemic Lupus Erythematosus in a Japanese Population. Journal of Rheumatology, 2009, 36, 2195-2203.	2.0	27
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105	STAT4 single nucleotide polymorphism, rs7574865 G/T, as a risk for antiphospholipid syndrome. Annals of the Rheumatic Diseases, 2009, 68, 1366-1367.	0.9	19
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115	Cytokines in Systemic Lupus Erythematosus. <i>Current Molecular Medicine</i> , 2009, 9, 242-254.	1.3	85
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117	Data for Genetic Analysis Workshop 16 Problem 1, association analysis of rheumatoid arthritis data. <i>BMC Proceedings</i> , 2009, 3, S2.	1.6	48
118	Replication of recently identified associated single-nucleotide polymorphisms from six autoimmune diseases in Genetic Analysis Workshop 16 rheumatoid arthritis data. <i>BMC Proceedings</i> , 2009, 3, S31.	1.6	7
119	Incorporating biological knowledge in the search for gene – gene interaction in genome-wide association studies. <i>BMC Proceedings</i> , 2009, 3, S81.	1.6	3
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125	A new gene-based association test for genome-wide association studies. <i>BMC Proceedings</i> , 2009, 3, S130.	1.6	19
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127	Prediction of Candidate Primary Immunodeficiency Disease Genes Using a Support Vector Machine Learning Approach. <i>DNA Research</i> , 2009, 16, 345-351.	3.4	26
128	A haplotype in STAT4 gene associated with rheumatoid arthritis in Caucasians is not associated in the Han Chinese population, but with the presence of rheumatoid factor. <i>Rheumatology</i> , 2009, 48, 1363-1368.	1.9	14
129	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.	0.9	138
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131	A combinatorial approach for detecting gene-gene interaction using multiple traits of Genetic Analysis Workshop 16 rheumatoid arthritis data. BMC Proceedings, 2009, 3, S43.	1.6	4
132	IL2RA Genetic Heterogeneity in Multiple Sclerosis and Type 1 Diabetes Susceptibility and Soluble Interleukin-2 Receptor Production. PLoS Genetics, 2009, 5, e1000322.	3.5	210
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134	Genome-wide association analyses of North American Rheumatoid Arthritis Consortium and Framingham Heart Study data utilizing genome-wide linkage results. BMC Proceedings, 2009, 3, S103.	1.6	12
135	Gene hunting of the Genetic Analysis Workshop 16 rheumatoid arthritis data using rough set theory. BMC Proceedings, 2009, 3, S126.	1.6	9
136	Rheumatoid arthritis-associated gene-gene interaction network for rheumatoid arthritis candidate genes. BMC Proceedings, 2009, 3, S75.	1.6	17
137	Association of rheumatoid factor and anti-cyclic citrullinated peptide positivity, but not carriage of shared epitope or <i>PTPN22</i> susceptibility variants, with anti-tumour necrosis factor response in rheumatoid arthritis. Annals of the Rheumatic Diseases, 2009, 68, 69-74.	0.9	240
138	Investigating the viability of genetic screening/testing for RA susceptibility using combinations of five confirmed risk loci. Rheumatology, 2009, 48, 1369-1374.	1.9	20
139	Homozygosity for DNASE2 single nucleotide polymorphisms in the 5â€²-regulatory region is associated with rheumatoid arthritis. Annals of the Rheumatic Diseases, 2009, 68, 1498-1503.	0.9	20
140	Temporal Induction Pattern of STAT4 Target Genes Defines Potential for Th1 Lineage-Specific Programming. Journal of Immunology, 2009, 183, 3839-3847.	0.8	64
141	Current concepts in the pathogenesis of early rheumatoid arthritis. Best Practice and Research in Clinical Rheumatology, 2009, 23, 37-48.	3.3	51
142	The preclinical stages of RA: lessons from human studies and animal models. Best Practice and Research in Clinical Rheumatology, 2009, 23, 49-58.	3.3	28
143	The search for lupus biomarkers. Best Practice and Research in Clinical Rheumatology, 2009, 23, 507-523.	3.3	80
144	Rheumatic diseases: Environment and genetics. Joint Bone Spine, 2009, 76, 602-607.	1.6	35
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149	A clinical perspective of rheumatoid arthritis. European Journal of Immunology, 2009, 39, 2044-2048.	2.9	4

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151	The chromosome 7q region association with rheumatoid arthritis in females in a british population is not replicated in a North American caseâ€“control series. <i>Arthritis and Rheumatism</i> , 2009, 60, 47-52.	6.7	5
152	Study of functional variants of the <i>BANK1</i> gene in rheumatoid arthritis. <i>Arthritis and Rheumatism</i> , 2009, 60, 372-379.	6.7	50
153	Genetic risk factors for rheumatoid arthritis differ in caucasian and Korean populations. <i>Arthritis and Rheumatism</i> , 2009, 60, 364-371.	6.7	109
154	Familial associations of rheumatoid arthritis with autoimmune diseases and related conditions. <i>Arthritis and Rheumatism</i> , 2009, 60, 661-668.	6.7	188
155	Biomarkers of inflammation and development of rheumatoid arthritis in women from two prospective cohort studies. <i>Arthritis and Rheumatism</i> , 2009, 60, 641-652.	6.7	118
156	Specific association of type 1 diabetes mellitus with antiâ€“cyclic citrullinated peptideâ€“positive rheumatoid arthritis. <i>Arthritis and Rheumatism</i> , 2009, 60, 653-660.	6.7	76
157	Quantitative heritability of antiâ€“citrullinated protein antibodyâ€“positive and antiâ€“citrullinated protein antibodyâ€“negative rheumatoid arthritis. <i>Arthritis and Rheumatism</i> , 2009, 60, 916-923.	6.7	200
158	Highâ€“density genotyping of STAT4 reveals multiple haplotypic associations with systemic lupus erythematosus in different racial groups. <i>Arthritis and Rheumatism</i> , 2009, 60, 1085-1095.	6.7	82
159	The <i>PRL</i> â€“1149 G/T polymorphism and rheumatoid arthritis susceptibility. <i>Arthritis and Rheumatism</i> , 2009, 60, 1250-1254.	6.7	23
160	Confirmation of <i>STAT4</i> , <i>IL2/IL21</i> , and <i>CTLA4</i> polymorphisms in rheumatoid arthritis. <i>Arthritis and Rheumatism</i> , 2009, 60, 1255-1260.	6.7	84
161	Geneâ€“environment interaction between the DRB1 shared epitope and smoking in the risk of antiâ€“citrullinated protein antibodyâ€“positive rheumatoid arthritis: All alleles are important. <i>Arthritis and Rheumatism</i> , 2009, 60, 1597-1603.	6.7	129
162	Variants in <i>TNFAIP3</i> , <i>STAT4</i> , and <i>C12orf30</i> loci associated with multiple autoimmune diseases are also associated with juvenile idiopathic arthritis. <i>Arthritis and Rheumatism</i> , 2009, 60, 2124-2130.	6.7	75
163	Genetic susceptibility to rheumatoid arthritis: An emerging picture. <i>Arthritis and Rheumatism</i> , 2009, 61, 1441-1446.	6.7	79
164	Regulation of the interferonâ€“ γ production induced by RNAâ€“containing immune complexes in plasmacytoid dendritic cells. <i>Arthritis and Rheumatism</i> , 2009, 60, 2418-2427.	6.7	121
165	<i>STAT4</i> is a genetic risk factor for systemic sclerosis having additive effects with <i>IRF5</i> on disease susceptibility and related pulmonary fibrosis. <i>Arthritis and Rheumatism</i> , 2009, 60, 2472-2479.	6.7	139
166	Association of <i>STAT4</i> and <i>BLK</i> , but not <i>BANK1</i> or <i>IRF5</i> , with primary antiphospholipid syndrome. <i>Arthritis and Rheumatism</i> , 2009, 60, 2468-2471.	6.7	66
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