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Rheumatoid arthritis-associated DNA methylation sites in peripheral blood mononuclear cells

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#	Paper	IF	Citations
60	Epigenetic Changes in the Pathogenesis of Rheumatoid Arthritis. <i>Frontiers in Genetics</i> , 2019 , 10, 570	4.5	56
59	Autoimmune diseases in China. <i>Advances in Immunology</i> , 2019 , 144, 173-216	5.6	3
58	SOAT1 methylation is associated with coronary heart disease. <i>Lipids in Health and Disease</i> , 2019 , 18, 192	4.4	4
57	Epigenetically regulated co-expression network of genes significant for rheumatoid arthritis. <i>Epigenomics</i> , 2019 , 11, 1601-1612	4.4	5
56	SAMD9 is a (epi-) genetically regulated anti-inflammatory factor activated in RA patients. <i>Molecular and Cellular Biochemistry</i> , 2019 , 456, 135-144	4.2	7
55	Integrative analysis of DNA methylation in discordant twins unveils distinct architectures of systemic sclerosis subsets. <i>Clinical Epigenetics</i> , 2019 , 11, 58	7.7	16
54	DNA methylation and the hygiene hypothesis: connecting respiratory allergy and childhood acute lymphoblastic leukemia. <i>Epigenomics</i> , 2019 , 11, 1519-1537	4.4	2
53	Mendelian randomization analysis revealed potential causal factors for systemic lupus erythematosus. <i>Immunology</i> , 2020 , 159, 279-288	7.8	6
52	Decipher manifestations and Treg /Th17 imbalance in multi-staging rheumatoid arthritis and correlation with TSDR/RORC methylation. <i>Molecular Immunology</i> , 2020 , 127, 1-11	4.3	7
51	Response to 'Correspondence on 'Rheumatoid arthritis-associated DNA methylation sites in peripheral blood mononuclear cells'' by Wang and Niu. <i>Annals of the Rheumatic Diseases</i> , 2020 ,	2.4	
50	Correspondence on 'Rheumatoid arthritis-associated DNA methylation sites in peripheral blood mononuclear cells'. <i>Annals of the Rheumatic Diseases</i> , 2020 ,	2.4	0
49	Identification of differentially expressed and methylated genes associated with rheumatoid arthritis based on network. <i>Autoimmunity</i> , 2020 , 53, 303-313	3	6
48	Alteration of circulating microbiome and its associated regulation role in rheumatoid arthritis: Evidence from integration of multiomics data. <i>Clinical and Translational Medicine</i> , 2020 , 10, e229	5.7	3
47	Effects of Biological Therapies on Molecular Features of Rheumatoid Arthritis. <i>International Journal of Molecular Sciences</i> , 2020 , 21,	6.3	13
46	Epigenetics, pregnancy and autoimmune rheumatic diseases. <i>Autoimmunity Reviews</i> , 2020 , 19, 102685	13.6	7
45	Association between NPPA promoter methylation and hypertension: results from Gusu cohort and replication in an independent sample. <i>Clinical Epigenetics</i> , 2020 , 12, 133	7.7	5
44	Latin American Genes: The Great Forgotten in Rheumatoid Arthritis. <i>Journal of Personalized Medicine</i> , 2020 , 10,	3.6	1

43	Transcriptome-wide association study for persistent hepatitis B virus infection and related hepatocellular carcinoma. <i>Liver International</i> , 2020 , 40, 2117-2127	7.9	1
42	Type I Interferons in the Pathogenesis and Treatment of Autoimmune Diseases. <i>Clinical Reviews in Allergy and Immunology</i> , 2020 , 59, 248-272	12.3	28
41	Peptidyl Arginine Deiminase Type 4 Gene Promoter Hypo-Methylation in Rheumatoid Arthritis. <i>Journal of Clinical Medicine</i> , 2020 , 9,	5.1	3
40	Integrative lncRNA-mRNA co-expression network analysis identifies novel lncRNA E2F3-IT1 for rheumatoid arthritis. <i>Clinical and Translational Medicine</i> , 2021 , 11, e325	5.7	1
39	Integrative analysis identifies potential causal methylation-mRNA regulation chains for rheumatoid arthritis. <i>Molecular Immunology</i> , 2021 , 131, 89-96	4.3	2
38	The current status and challenges in the diagnosis and treatment of rheumatoid arthritis in China: An annual report of 2019. <i>Rheumatology and Immunology Research</i> , 2021 , 2, 49-56	0.2	2
37	Epigenome wide association study of response to methotrexate in early rheumatoid arthritis patients. <i>PLoS ONE</i> , 2021 , 16, e0247709	3.7	2
36	Evaluation of the Genetic Association and Methylation of Immune Response Pathway Genes with the Risk of Chronic Periodontitis in the Uighur Population. <i>Genetic Testing and Molecular Biomarkers</i> , 2021 , 25, 317-324	1.6	
35	Interaction between early-life pet exposure and methylation pattern of ADAM33 on allergic rhinitis among children aged 3-6 years in China. <i>Allergy, Asthma and Clinical Immunology</i> , 2021 , 17, 44	3.2	0
34	Association of methylation level and transcript level in TRAF5 gene with ankylosing spondylitis: a case-control study. <i>Genes and Immunity</i> , 2021 , 22, 101-107	4.4	2
33	Prediction of the Progression of Undifferentiated Arthritis to Rheumatoid Arthritis Using DNA Methylation Profiling. <i>Arthritis and Rheumatology</i> , 2021 , 73, 2229-2239	9.5	5
32	Association analysis of juvenile idiopathic arthritis genetic susceptibility factors in Estonian patients. <i>Clinical Rheumatology</i> , 2021 , 40, 4157-4165	3.9	1
31	Machine Learning Based Image Feature Recognition and Clinical Nursing of Children's Rheumatoid Arthritis-Related Lung Injury. <i>Scientific Programming</i> , 2021 , 2021, 1-8	1.4	
30	Methylation Promotes Inflammation and Activation of Fibroblast-Like Synoviocytes in Rheumatoid Arthritis. <i>Frontiers in Pharmacology</i> , 2021 , 12, 700373	5.6	2
29	Emerging epigenetic targets in rheumatoid arthritis. <i>Rheumatology International</i> , 2021 , 41, 2047-2067	3.6	2
28	Epigenetic and transcriptional control of interferon- γ . <i>Journal of Experimental Medicine</i> , 2021 , 218,	16.6	2
27	Ankylosing Spondylitis Patients Display Aberrant Gene DNA Methylation and Expression. <i>Immunological Investigations</i> , 2021 , 1-13	2.9	0
26	The role of oxidative stress in epigenetic changes underlying autoimmunity. <i>Antioxidants and Redox Signaling</i> , 2021 ,	8.4	0

25	Fetal malnutrition is associated with impairment of endogenous melatonin synthesis in pineal via hypermethylation of promoters of protein kinase C alpha and cAMP response element-binding. <i>Journal of Pineal Research</i> , 2021 , 71, e12764	10.4	1
24	The DNA methylation Profile of Undifferentiated Arthritis Patients Anticipates their Subsequent Differentiation to Rheumatoid Arthritis.		
23	Etiology and Risk Factors for Rheumatoid Arthritis: A State-of-the-Art Review.. <i>Frontiers in Medicine</i> , 2021 , 8, 689698	4.9	6
22	Differentially methylation of IFI44L gene promoter in Iranian patients with systemic lupus erythematosus and rheumatoid arthritis.. <i>Molecular Biology Reports</i> , 2022 , 49, 3065	2.8	1
21	Promoter DNA Methylation in GWAS-Identified Genes as Potential Functional Elements for Blood Pressure: An Observational and Mendelian Randomization Study.. <i>Frontiers in Genetics</i> , 2021 , 12, 791146	4.5	0
20	Serum Atrial Natriuretic Peptide, Promoter Methylation, and Cardiovascular Disease: A 10-year Follow-Up Study in Chinese Adults.. <i>Global Heart</i> , 2022 , 17, 27	2.9	0
19	piRNA-6426 increases DNMT3B-mediated SOAT1 methylation and improves heart failure.. <i>Aging</i> , 2022 , 14, 2678-2694	5.6	0
18	Association of DNA Methylation in Blood Pressure-Related Genes With Ischemic Stroke Risk and Prognosis.. <i>Frontiers in Cardiovascular Medicine</i> , 2022 , 9, 796245	5.4	0
17	Identification of novel rheumatoid arthritis-associated MiRNA-204-5p from plasma exosomes.. <i>Experimental and Molecular Medicine</i> , 2022 ,	12.8	0
16	Epigenetic Regulation in the Pathogenesis of Rheumatoid Arthritis.. <i>Frontiers in Immunology</i> , 2022 , 13, 859400	8.4	1
15	NPPA promoter hypomethylation predicts central obesity development: a prospective longitudinal study in Chinese adults. <i>Obesity Facts</i> , 2021 ,	5.1	0
14	Epigenetic Regulation of Immune and Inflammatory Responses in Rheumatoid Arthritis.. <i>Frontiers in Immunology</i> , 2022 , 13, 881191	8.4	0
13	[Methylation status and expression of gene promoter region in peripheral blood of patients with rheumatoid arthritis].. <i>Beijing Da Xue Xue Bao</i> , 2021 , 53, 1020-1025	0.2	
12	Gene Interaction Network Analysis Reveals IFI44L as a Drug Target in Rheumatoid Arthritis and Periodontitis.. <i>Molecules</i> , 2022 , 27,	4.8	0
11	DNA Methylation of the Natriuretic Peptide System Genes and Ischemic Stroke. <i>Neurology: Genetics</i> , 2022 , 8, e679	3.8	
10	Mitochondrial Dysfunction and Oxidative Stress in Rheumatoid Arthritis. <i>Antioxidants</i> , 2022 , 11, 1151	7.1	2
9	Interferon- β -mediated therapeutic resistance in early rheumatoid arthritis implicates epigenetic reprogramming. <i>Annals of the Rheumatic Diseases</i> , annrheumdis-2022-222370	2.4	0
8	DNA Methylation and mRNA Expression of B7-H3 Gene in Ankylosing Spondylitis: A Case-Control Study. <i>Immunological Investigations</i> , 1-10	2.9	0

7	Single-cell RNA sequencing analysis of T helper cell differentiation and heterogeneity. <i>Biochimica Et Biophysica Acta - Molecular Cell Research</i> , 2022 , 1869, 119321	4.9	
6	Geniposide restricts angiogenesis in experimental arthritis via inhibiting Dnmt1-mediated PTEN hypermethylation. <i>International Immunopharmacology</i> , 2022 , 111, 109087	5.8	1
5	Disease mechanisms in preclinical rheumatoid arthritis: A narrative review. 9,		2
4	Identification of methylation signatures and rules for predicting the severity of SARS-CoV-2 infection with machine learning methods. 13,		0
3	A novel long noncoding RNA, lnc-RNU12, influences the T-cell cycle via c-JUN and CCNL2 in rheumatoid arthritis.		0
2	Seropositivity-Dependent Association between LINE-1 Methylation and Response to Methotrexate Therapy in Early Rheumatoid Arthritis Patients. 2022 , 13, 2012		1
1	Differential CpG DNA methylation of peripheral B cells, CD4+ T cells, and salivary gland tissues in IgG4-related disease. 2023 , 25,		0