Sara Marsal

List of Publications by Year in descending order

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78 7,512 33 77 papers citations h-index g-index

82 82 82 82 19074

82 82 82 19074 all docs docs citations times ranked citing authors

#	Article	IF	CITATIONS
1	Genomewide Association Study of Severe Covid-19 with Respiratory Failure. New England Journal of Medicine, 2020, 383, 1522-1534.	27.0	1,548
2	Modeling Linkage Disequilibrium Increases Accuracy of Polygenic Risk Scores. American Journal of Human Genetics, 2015, 97, 576-592.	6.2	1,098
3	Genome-wide association study identifies eight risk loci and implicates metabo-psychiatric origins for anorexia nervosa. Nature Genetics, 2019, 51, 1207-1214.	21.4	641
4	Partitioning Heritability of Regulatory and Cell-Type-Specific Variants across 11 Common Diseases. American Journal of Human Genetics, 2014, 95, 535-552.	6.2	569
5	Analytical Methods in Untargeted Metabolomics: State of the Art in 2015. Frontiers in Bioengineering and Biotechnology, 2015, 3, 23.	4.1	495
6	Significant Locus and Metabolic Genetic Correlations Revealed in Genome-Wide Association Study of Anorexia Nervosa. American Journal of Psychiatry, 2017, 174, 850-858.	7.2	410
7	Genetic Structure of Europeans: A View from the North–East. PLoS ONE, 2009, 4, e5472.	2.5	279
8	Intravenous administration of expanded allogeneic adipose-derived mesenchymal stem cells in refractory rheumatoid arthritis (Cx611): results of a multicentre, dose escalation, randomised, single-blind, placebo-controlled phase lb/lla clinical trial. Annals of the Rheumatic Diseases, 2017, 76, 196-202.	0.9	194
9	Antiangiogenic effects of anti-tumor necrosis factor $\hat{l}\pm$ therapy with infliximab in psoriatic arthritis. Arthritis and Rheumatism, 2004, 50, 1636-1641.	6.7	137
10	Incidence of COVID-19 in a cohort of adult and paediatric patients with rheumatic diseases treated with targeted biologic and synthetic disease-modifying anti-rheumatic drugs. Seminars in Arthritis and Rheumatism, 2020, 50, 564-570.	3.4	129
11	Estimation of Genetic Correlation via Linkage Disequilibrium Score Regression and Genomic Restricted Maximum Likelihood. American Journal of Human Genetics, 2018, 102, 1185-1194.	6.2	119
12	Genome-wide association study meta-analysis identifies five new loci for systemic lupus erythematosus. Arthritis Research and Therapy, 2018, 20, 100.	3.5	102
13	Genomeâ€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
14	Urine metabolome profiling of immune-mediated inflammatory diseases. BMC Medicine, 2016, 14, 133.	5.5	97
15	An Eight-Gene Blood Expression Profile Predicts the Response to Infliximab in Rheumatoid Arthritis. PLoS ONE, 2009, 4, e7556.	2.5	94
16	Actualización 2014 del Documento de Consenso de la Sociedad Española de ReumatologÃa sobre el uso de terapias biológicas en la artritis reumatoide. ReumatologÃa ClÃnica, 2015, 11, 279-294.	0.5	89
17	Risk variants for psoriasis vulgaris in a large case–control collection and association with clinical subphenotypes. Human Molecular Genetics, 2012, 21, 4549-4557.	2.9	79
18	Epigenome-wide association study of rheumatoid arthritis identifies differentially methylated loci in B cells. Human Molecular Genetics, 2017, 26, 2803-2811.	2.9	67

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19	Synovial fibroblast hyperplasia in rheumatoid arthritis: Clinicopathologic correlations and partial reversal by anti–tumor necrosis factor therapy. Arthritis and Rheumatism, 2011, 63, 2575-2583.	6.7	66
20	Genetic characterization of northeastern Italian population isolates in the context of broader European genetic diversity. European Journal of Human Genetics, 2013, 21, 659-665.	2.8	64
21	GWAS replication study confirms the association of <i>PDE3A–SLCO1C1</i> with anti-TNF therapy response in rheumatoid arthritis. Pharmacogenomics, 2013, 14, 727-734.	1.3	61
22	Evidence for Genetic Overlap Between Schizophrenia and Age at First Birth in Women. JAMA Psychiatry, 2016, 73, 497.	11.0	51
23	Pro- and anti-inflammatory eicosanoids in psoriatic arthritis. Metabolomics, 2019, 15, 65.	3.0	49
24	AStream: an R package for annotating LC/MS metabolomic data. Bioinformatics, 2011, 27, 1339-1340.	4.1	46
25	Identification of Risk Loci for Crohn's Disease Phenotypes Using a Genome-Wide Association Study. Gastroenterology, 2015, 148, 794-805.	1.3	46
26	Genetic variation at the glycosaminoglycan metabolism pathway contributes to the risk of psoriatic arthritis but not psoriasis. Annals of the Rheumatic Diseases, 2019, 78, 355-364.	0.9	44
27	Association of HLA Class II Genes with Systemic Sclerosis in Spanish Patients. Journal of Rheumatology, 2009, 36, 2733-2736.	2.0	43
28	Associations Between Attention-Deficit/Hyperactivity Disorder and Various Eating Disorders: A Swedish Nationwide Population Study Using Multiple Genetically Informative Approaches. Biological Psychiatry, 2019, 86, 577-586.	1.3	43
29	A genome-wide association study on a southern European population identifies a new Crohn's disease susceptibility locus at <i>RBX1-EP300</i> . Gut, 2013, 62, 1440-1445.	12.1	42
30	A genome-wide association study identifies a novel locus at 6q22.1 associated with ulcerative colitis. Human Molecular Genetics, 2014, 23, 6927-6934.	2.9	39
31	Focus: A Robust Workflow for One-Dimensional NMR Spectral Analysis. Analytical Chemistry, 2014, 86, 1160-1169.	6.5	36
32	A functional variant of TLR10 modifies the activity of NFkB and may help predict a worse prognosis in patients with rheumatoid arthritis. Arthritis Research and Therapy, 2016, 18, 221.	3.5	35
33	Identification of a two-loci epistatic interaction associated with susceptibility to rheumatoid arthritis through reverse engineering and multifactor dimensionality reduction. Genomics, 2007, 90, 6-13.	2.9	34
34	Variation at FCGR2A and Functionally Related Genes Is Associated with the Response to Anti-TNF Therapy in Rheumatoid Arthritis. PLoS ONE, 2015, 10, e0122088.	2.5	33
35	Cardiovascular disease in immune-mediated inflammatory diseases. Medicine (United States), 2017, 96, e7308.	1.0	32
36	Shared genetic risk between eating disorder―and substance―use―elated phenotypes: Evidence from genomeâ€wide association studies. Addiction Biology, 2021, 26, e12880.	2.6	28

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37	Genome-Wide Pathway Analysis Identifies Genetic Pathways Associated with Psoriasis. Journal of Investigative Dermatology, 2016, 136, 593-602.	0.7	27
38	DAS-28-based EULAR response and HAQ improvement in rheumatoid arthritis patients switching between TNF antagonists. BMC Musculoskeletal Disorders, 2009, 10, 91.	1.9	26
39	Deletion of the late cornified envelope genes, <i>LCE3C</i> and <i>LCE3B</i> , is associated with rheumatoid arthritis. Arthritis and Rheumatism, 2010, 62, 1246-1251.	6.7	26
40	Genome-wide transcriptional analysis of T cell activation reveals differential gene expression associated with psoriasis. BMC Genomics, 2013, 14, 825.	2.8	25
41	A Combined Transcriptomic and Genomic Analysis Identifies a Gene Signature Associated With the Response to Anti-TNF Therapy in Rheumatoid Arthritis. Frontiers in Immunology, 2019, 10, 1459.	4.8	24
42	aRNApipe: a balanced, efficient and distributed pipeline for processing RNA-seq data in high-performance computing environments. Bioinformatics, 2017, 33, 1727-1729.	4.1	23
43	Identification of candidate genes for rituximab response in rheumatoid arthritis patients by microarray expression profiling in blood cells. Pharmacogenomics, 2009, 10, 1697-1708.	1.3	22
44	Novel Insights into the Regulatory Architecture of CD4+ T Cells in Rheumatoid Arthritis. PLoS ONE, 2014, 9, e100690.	2.5	22
45	Rheumatoid arthritis pharmacogenomics. Pharmacogenomics, 2010, 11, 617-619.	1.3	20
46	A deletion atADAMTS9-MAGIIlocus is associated with psoriatic arthritis risk. Annals of the Rheumatic Diseases, 2015, 74, 1875-1881.	0.9	18
47	Discoidin domain receptor 1 gene variants are associated with decreased white matter fractional anisotropy and decreased processing speed in schizophrenia. Journal of Psychiatric Research, 2019, 110, 74-82.	3.1	18
48	A cross-disease meta-GWAS identifies four new susceptibility loci shared between systemic sclerosis and Crohn's disease. Scientific Reports, 2020, 10, 1862.	3.3	18
49	CNstream: A method for the identification and genotyping of copy number polymorphisms using Illumina microarrays. BMC Bioinformatics, 2010, 11, 264.	2.6	15
50	A genome-wide association study identifies <i>SLC8A3</i> as a susceptibility locus for ACPA-positive rheumatoid arthritis. Rheumatology, 2016, 55, 1106-1111.	1.9	14
51	Genome-wide pathway analysis identifies VEGF pathway association with oral ulceration in systemic lupus erythematosus. Arthritis Research and Therapy, 2017, 19, 138.	3.5	14
52	Systemic Lupus Erythematosus: Genomics, Mechanisms, and Therapies. Clinical and Developmental Immunology, 2012, 2012, 1-2.	3.3	12
53	Biosimilar and interchangeable: Inseparable scientific concepts?. British Journal of Clinical Pharmacology, 2019, 85, 2460-2463.	2.4	12
54	Loss of imprinting of <i>IGF2</i> characterises high IGF2 mRNA-expressing type of fibroblast-like synoviocytes in rheumatoid arthritis. Annals of the Rheumatic Diseases, 2010, 69, 1239-1242.	0.9	11

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55	Variation at interleukin-6 receptor gene is associated to joint damage in rheumatoid arthritis. Arthritis Research and Therapy, 2015, 17, 242.	3.5	11
56	Targeting of the CD80/86 proinflammatory axis as a therapeutic strategy to prevent severe COVID-19. Scientific Reports, 2021, 11, 11462.	3.3	11
57	Asistencia reumatológica en el sector sanitario público de Cataluña: año 2012. ReumatologÃa ClÃnica, 2014, 10, 85-88.	0.5	10
58	The Genetic Architecture of Rheumatoid Arthritis: From Susceptibility to Clinical Subphenotype Associations. Current Topics in Medicinal Chemistry, 2013, 13, 720-731.	2.1	9
59	<i>PDE3A-SLCO1C1</i> locus is associated with response to anti-tumor necrosis factor therapy in psoriatic arthritis. Pharmacogenomics, 2014, 15, 1763-1769.	1.3	9
60	Longitudinal analysis of blood DNA methylation identifies mechanisms of response to tumor necrosis factor inhibitor therapy in rheumatoid arthritis. EBioMedicine, 2022, 80, 104053.	6.1	9
61	Retention of subcutaneous abatacept for the treatment of rheumatoid arthritis: real-world results from the ASCORE study: an international 2-year observational study. Clinical Rheumatology, 2022, , 1.	2.2	7
62	Metabolomics in rheumatic diseases. International Journal of Clinical Rheumatology, 2014, 9, 353-369.	0.3	6
63	Identification of <i>IRX1</i> as a Risk Locus for Rheumatoid Factor Positivity in Rheumatoid Arthritis in a Genomeâ€Wide Association Study. Arthritis and Rheumatology, 2016, 68, 1384-1391.	5.6	6
64	Food groups associated with immune-mediated inflammatory diseases: a Mendelian randomization and disease severity study. European Journal of Clinical Nutrition, 2021, 75, 1368-1382.	2.9	5
65	Genetic variation associated with cardiovascular risk in autoimmune diseases. PLoS ONE, 2017, 12, e0185889.	2.5	5
66	GStream: Improving SNP and CNV Coverage on Genome-Wide Association Studies. PLoS ONE, 2013, 8, e68822.	2.5	4
67	Pharmacogenomics of anti-TNF response in psoriasis, where are we?. Pharmacogenomics, 2016, 17, 323-326.	1.3	4
68	Genetic association between CD96 locus and immunogenicity to anti-TNF therapy in Crohn's disease. Pharmacogenomics Journal, 2019, 19, 547-555.	2.0	4
69	Interactions between rheumatoid arthritis antibodies are associated with the response to anti-tumor necrosis factor therapy. BMC Musculoskeletal Disorders, 2021, 22, 372.	1.9	4
70	Unveiling Caseâ€Control Relationships in Designing a Simple and Powerful Method for Detecting Geneâ€Gene Interactions. Genetic Epidemiology, 2012, 36, 710-716.	1.3	3
71	Clinical and therapeutic management of rheumatoid arthritis with biological disease-modifying antirheumatic drugs: RADAR study. Rheumatology International, 2019, 39, 2015-2024.	3.0	2
72	Populationâ€based identityâ€byâ€descent mapping combined with exome sequencing to detect rare risk variants for schizophrenia. American Journal of Medical Genetics Part B: Neuropsychiatric Genetics, 2019, 180, 223-231.	1.7	2

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73	Lack of association between the corticotropin-releasing hormone locus and rheumatoid arthritis. Arthritis and Rheumatism, 2004, 50, 2706-2708.	6.7	1
74	Functional rare variants influence the clinical response to anti-TNF therapy in Crohn's disease. Therapeutic Advances in Gastroenterology, 2019, 12, 175628481986784.	3.2	1
75	A questionnaire-based study on contraceptive practice in patients with rheumatic disease found no significant difference in age-matched healthy controls. Rheumatology International, 2020, 40, 1473-1480.	3.0	1
76	Synovial lymphoid neogenesis is associated with IL-23 expression and disease activity in rheumatoid arthritis. Journal of Translational Medicine, 2011, 9, .	4.4	0
77	Dietary habits in patients with fibromyalgia: a cross-sectional study. Clinical and Experimental Rheumatology, 2021, 39 Suppl 130, 170-173.	0.8	0
78	Trajectories in early rheumatoid arthritis related fatigue over 10 years: results from the ESPOIR cohort. Clinical and Experimental Rheumatology, 2021, , .	0.8	0