

Gisela Orozco

List of Publications by Year in descending order

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Version: 2024-02-01

76
papers

2,946
citations

147801

31
h-index

175258

52
g-index

85
all docs

85
docs citations

85
times ranked

4649
citing authors

#	ARTICLE	IF	CITATIONS
1	Association of a functional single nucleotide polymorphism of <i>PTPN22</i> , encoding lymphoid protein phosphatase, with rheumatoid arthritis and systemic lupus erythematosus. <i>Arthritis and Rheumatism</i> , 2005, 52, 219-224.	6.7	275
2	Capture Hi-C reveals novel candidate genes and complex long-range interactions with related autoimmune risk loci. <i>Nature Communications</i> , 2015, 6, 10069.	12.8	161
3	The IL23R Arg381Gln non-synonymous polymorphism confers susceptibility to ankylosing spondylitis. <i>Annals of the Rheumatic Diseases</i> , 2008, 67, 1451-1454.	0.9	142
4	Study of the common genetic background for rheumatoid arthritis and systemic lupus erythematosus. <i>Annals of the Rheumatic Diseases</i> , 2011, 70, 463-468.	0.9	130
5	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
6	Confirmation of TNIP1 and IL23A as susceptibility loci for psoriatic arthritis. <i>Annals of the Rheumatic Diseases</i> , 2011, 70, 1641-1644.	0.9	103
7	GWAS for systemic sclerosis identifies multiple risk loci and highlights fibrotic and vasculopathy pathways. <i>Nature Communications</i> , 2019, 10, 4955.	12.8	100
8	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
9	Combined effects of three independent SNPs greatly increase the risk estimate for RA at 6q23. <i>Human Molecular Genetics</i> , 2009, 18, 2693-2699.	2.9	93
10	Capture Hi-C identifies a novel causal gene, IL20RA, in the pan-autoimmune genetic susceptibility region 6q23. <i>Genome Biology</i> , 2016, 17, 212.	8.8	85
11	Genetic basis of rheumatoid arthritis. <i>Biomedicine and Pharmacotherapy</i> , 2006, 60, 656-662.	5.6	77
12	Analysis of a Functional BTNL2 Polymorphism in Type 1 Diabetes, Rheumatoid Arthritis, and Systemic Lupus Erythematosus. <i>Human Immunology</i> , 2005, 66, 1235-1241.	2.4	70
13	Association of CD40 with rheumatoid arthritis confirmed in a large UK case-control study. <i>Annals of the Rheumatic Diseases</i> , 2010, 69, 813-816.	0.9	62
14	Epistatic interaction between FCRL3 and NFAB1 genes in Spanish patients with rheumatoid arthritis. <i>Annals of the Rheumatic Diseases</i> , 2006, 65, 1188-1191.	0.9	59
15	Synthetic associations in the context of genome-wide association scan signals. <i>Human Molecular Genetics</i> , 2010, 19, R137-R144.	2.9	53
16	Investigation of the IL23R gene in a Spanish rheumatoid arthritis cohort. <i>Human Immunology</i> , 2007, 68, 681-684.	2.4	52
17	Novel Association of the Interleukin 23 Interleukin 21 Region With Inflammatory Bowel Disease. <i>American Journal of Gastroenterology</i> , 2009, 104, 1968-1975.	0.4	51
18	Macrophage migration inhibitory factor gene: Influence on rheumatoid arthritis susceptibility. <i>Human Immunology</i> , 2007, 68, 744-747.	2.4	50

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19	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
20	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
21	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.	3.5	50
22	Brief Report: Identification of <i>BACH2</i> and <i>RAD51B</i> as Rheumatoid Arthritis Susceptibility Loci in a Meta-Analysis of Genome-Wide Data. <i>Arthritis and Rheumatism</i> , 2013, 65, 3058-3062.	6.7	43
23	C1858T Functional Variant of PTPN22 Gene Is Not Associated With Celiac Disease Genetic Predisposition. <i>Human Immunology</i> , 2005, 66, 848-852.	2.4	42
24	Novel Rheumatoid Arthritis Susceptibility Locus at 22q12 Identified in an Extended UK Genome-Wide Association Study. <i>Arthritis and Rheumatology</i> , 2014, 66, 24-30.	5.6	41
25	Auto-antibodies, HLA and PTPN22: susceptibility markers for rheumatoid arthritis. <i>Rheumatology</i> , 2007, 47, 138-141.	1.9	40
26	Protein tyrosine phosphatase gene (PTPN22) polymorphism in multiple sclerosis. <i>Journal of Neurology</i> , 2005, 252, 994-995.	3.6	38
27	Analysis of the functional NFKB1 promoter polymorphism in rheumatoid arthritis and systemic lupus erythematosus. <i>Tissue Antigens</i> , 2005, 65, 183-186.	1.0	35
28	Cytotoxic T-lymphocyte antigen-4-CT60 polymorphism in rheumatoid arthritis. <i>Tissue Antigens</i> , 2004, 64, 667-670.	1.0	34
29	Investigation of type 1 diabetes and coeliac disease susceptibility loci for association with juvenile idiopathic arthritis. <i>Annals of the Rheumatic Diseases</i> , 2010, 69, 2169-2172.	0.9	34
30	Update on the genetic risk factors for rheumatoid arthritis. <i>Expert Review of Clinical Immunology</i> , 2010, 6, 61-75.	3.0	34
31	The functional genetic variation in the PTPN22 gene has a negligible effect on the susceptibility to develop inflammatory bowel disease. <i>Tissue Antigens</i> , 2005, 66, 314-317.	1.0	32
32	Interleukin 12 (IL12B) and Interleukin 12 Receptor (IL12RB1) Gene Polymorphisms in Rheumatoid Arthritis. <i>Human Immunology</i> , 2005, 66, 710-714.	2.4	32
33	Combined genetic analysis of juvenile idiopathic arthritis clinical subtypes identifies novel risk loci, target genes and key regulatory mechanisms. <i>Annals of the Rheumatic Diseases</i> , 2021, 80, 321-328.	0.9	31
34	The genetics revolution in rheumatology: large scale genomic arrays and genetic mapping. <i>Nature Reviews Rheumatology</i> , 2017, 13, 421-432.	8.0	30
35	Patterns of constitutive and IFN- γ inducible expression of HLA class II molecules in human melanoma cell lines. <i>Immunogenetics</i> , 2007, 59, 123-133.	2.4	29
36	Rare variation at the TNFAIP3 locus and susceptibility to rheumatoid arthritis. <i>Human Genetics</i> , 2010, 128, 627-633.	3.8	29

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37	Identifying Causal Genes at the Multiple Sclerosis Associated Region 6q23 Using Capture Hi-C. PLoS ONE, 2016, 11, e0166923.	2.5	28
38	Asporin repeat polymorphism in rheumatoid arthritis. Annals of the Rheumatic Diseases, 2006, 66, 118-120.	0.9	27
39	Caspase 7 influences susceptibility to rheumatoid arthritis. Rheumatology, 2007, 46, 1243-1247.	1.9	27
40	Functional genomics atlas of synovial fibroblasts defining rheumatoid arthritis heritability. Genome Biology, 2021, 22, 247.	8.8	27
41	Analysis of a GT Microsatellite in the Promoter of the foxp3/scurfin Gene in Autoimmune Diseases. Human Immunology, 2005, 66, 869-873.	2.4	25
42	Comprehensive analysis of the major histocompatibility complex in systemic sclerosis identifies differential HLA associations by clinical and serological subtypes. Annals of the Rheumatic Diseases, 2021, 80, 1040-1047.	0.9	24
43	Lack of association between ankylosing spondylitis and a functional polymorphism of PTPN22 proposed as a general susceptibility marker for autoimmunity. Annals of the Rheumatic Diseases, 2006, 65, 687-688.	0.9	23
44	Chromatin interactions reveal novel gene targets for drug repositioning in rheumatic diseases. Annals of the Rheumatic Diseases, 2019, 78, 1127-1134.	0.9	23
45	Genetics of rheumatoid arthritis: GWAS and beyond. Open Access Rheumatology: Research and Reviews, 2011, 3, 31.	1.6	22
46	SLC22A4, RUNX1, and SUMO4 polymorphisms are not associated with rheumatoid arthritis: a case-control study in a Spanish population. Journal of Rheumatology, 2006, 33, 1235-9.	2.0	21
47	Sex-specific differences in effect size estimates at established complex trait loci. International Journal of Epidemiology, 2012, 41, 1376-1382.	1.9	19
48	Mapping DNA interaction landscapes in psoriasis susceptibility loci highlights KLF4 as a target gene in 9q31. BMC Biology, 2020, 18, 47.	3.8	19
49	Chromatin Looping Links Target Genes with Genetic Risk Loci for Dermatological Traits. Journal of Investigative Dermatology, 2021, 141, 1975-1984.	0.7	19
50	Inducible nitric oxide synthase promoter polymorphism in human brucellosis. Microbes and Infection, 2003, 5, 1165-1169.	1.9	17
51	Characterisation of CD4+ T-cell subtypes using single cell RNA sequencing and the impact of cell number and sequencing depth. Scientific Reports, 2020, 10, 19825.	3.3	17
52	Study of the role of a functional polymorphism of MHC2TA in rheumatoid arthritis in three ethnically different populations. Rheumatology, 2006, 45, 1442-1444.	1.9	15
53	HLA-DPB1-COL11A2 and three additional xMHC loci are independently associated with RA in a UK cohort. Genes and Immunity, 2011, 12, 169-175.	4.1	15
54	Study of the role of functional variants of SLC22A4, RUNX1 and SUMO4 in systemic lupus erythematosus. Annals of the Rheumatic Diseases, 2006, 65, 791-795.	0.9	14

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55	Identification of rheumatoid arthritis causal genes using functional genomics. <i>Scandinavian Journal of Immunology</i> , 2019, 89, e12753.	2.7	14
56	Lack of association of a functional single nucleotide polymorphism of PTPN22, encoding lymphoid protein phosphatase, with susceptibility to biopsy-proven giant cell arteritis. <i>Journal of Rheumatology</i> , 2005, 32, 1510-2.	2.0	12
57	HiChIP-Peaks: a HiChIP peak calling algorithm. <i>Bioinformatics</i> , 2020, 36, 3625-3631.	4.1	11
58	Transcriptome-wide study of TNF-inhibitor therapy in rheumatoid arthritis reveals early signature of successful treatment. <i>Arthritis Research and Therapy</i> , 2021, 23, 80.	3.5	11
59	Functional genomics in autoimmune diseases. <i>Human Molecular Genetics</i> , 2020, 29, R59-R65.	2.9	10
60	Common genetic variants associated with disease from genome-wide association studies are mutually exclusive in prostate cancer and rheumatoid arthritis. <i>BJU International</i> , 2013, 111, 1148-1155.	2.5	9
61	PTPN22C1858T polymorphism and human brucellosis. <i>Scandinavian Journal of Infectious Diseases</i> , 2009, 41, 109-112.	1.5	8
62	Analysis of SNP-SNP interactions and bone quantitative ultrasound parameter in early adulthood. <i>BMC Medical Genetics</i> , 2017, 18, 107.	2.1	8
63	Using functional genomics to advance the understanding of psoriatic arthritis. <i>Rheumatology</i> , 2020, 59, 3137-3146.	1.9	8
64	Fine mapping with epigenetic information and 3D structure. <i>Seminars in Immunopathology</i> , 2022, 44, 115-125.	6.1	8
65	Rheumatoid Arthritis-associated Polymorphisms at 6q23 Are Associated with Radiological Damage in Autoantibody-positive RA. <i>Journal of Rheumatology</i> , 2012, 39, 1781-1785.	2.0	7
66	Functional interrogation of autoimmune disease genetics using CRISPR/Cas9 technologies and massively parallel reporter assays. <i>Seminars in Immunopathology</i> , 2022, 44, 137-147.	6.1	5
67	Exploring the overlap between rheumatoid arthritis susceptibility loci and long non-coding RNA annotations. <i>PLoS ONE</i> , 2020, 15, e0223939.	2.5	2
68	Identification of new susceptibility markers for rheumatoid arthritis and systemic lupus erythematosus in the STAT4 gene. <i>Personalized Medicine</i> , 2008, 5, 169-174.	1.5	1
69	Common Genetic Component in Autoimmunity. <i>Rare Diseases of the Immune System</i> , 2019, , 221-236.	0.1	0
70	OP0191...EPIGENETIC PROFILING OF SYNOVIAL FIBROBLASTS REVEALS STRUCTURAL DNA DYNAMICS AT DISEASE IMPLICATED CHROMOSOME REGIONS. , 2019, , .		0
71	OP0192...DISSECTING THE LONG-RANGE GENE REGULATION OF RHEUMATOID ARTHRITIS RISK ENHANCERS AT THE 5Q11 LOCUS USING THE COMPLEMENTARY APPROACHES OF CRISPR AND CRISPRi. , 2019, , .		0
72	Title is missing!. , 2020, 15, e0223939.		0

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73	Title is missing!. , 2020, 15, e0223939.		0
74	Title is missing!. , 2020, 15, e0223939.		0
75	Title is missing!. , 2020, 15, e0223939.		0
76	OA30â€fIdentification of causal genes and mechanisms by which genetic variation mediates juvenile idiopathic arthritis susceptibility using functional genomics and CRISPR-Cas9. Rheumatology, 2022, 61, .	1.9	0