

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	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
2	Fine mapping with epigenetic information and 3D structure. <i>Seminars in Immunopathology</i> , 2022, 44, 115-125.	6.1	8
3	OA30 Identification of causal genes and mechanisms by which genetic variation mediates juvenile idiopathic arthritis susceptibility using functional genomics and CRISPR-Cas9. <i>Rheumatology</i> , 2022, 61, .	1.9	0
4	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
5	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
6	Comprehensive analysis of the major histocompatibility complex in systemic sclerosis identifies differential HLA associations by clinical and serological subtypes. <i>Annals of the Rheumatic Diseases</i> , 2021, 80, 1040-1047.	0.9	24
7	Chromatin Looping Links Target Genes with Genetic Risk Loci for Dermatological Traits. <i>Journal of Investigative Dermatology</i> , 2021, 141, 1975-1984.	0.7	19
8	Functional genomics atlas of synovial fibroblasts defining rheumatoid arthritis heritability. <i>Genome Biology</i> , 2021, 22, 247.	8.8	27
9	Characterisation of CD4+ T-cell subtypes using single cell RNA sequencing and the impact of cell number and sequencing depth. <i>Scientific Reports</i> , 2020, 10, 19825.	3.3	17
10	Using functional genomics to advance the understanding of psoriatic arthritis. <i>Rheumatology</i> , 2020, 59, 3137-3146.	1.9	8
11	Functional genomics in autoimmune diseases. <i>Human Molecular Genetics</i> , 2020, 29, R59-R65.	2.9	10
12	Mapping DNA interaction landscapes in psoriasis susceptibility loci highlights KLF4 as a target gene in 9q31. <i>BMC Biology</i> , 2020, 18, 47.	3.8	19
13	Exploring the overlap between rheumatoid arthritis susceptibility loci and long non-coding RNA annotations. <i>PLoS ONE</i> , 2020, 15, e0223939.	2.5	2
14	HiChIP-Peaks: a HiChIP peak calling algorithm. <i>Bioinformatics</i> , 2020, 36, 3625-3631.	4.1	11
15	Title is missing!. , 2020, 15, e0223939.		0
16	Title is missing!. , 2020, 15, e0223939.		0
17	Title is missing!. , 2020, 15, e0223939.		0
18	Title is missing!. , 2020, 15, e0223939.		0

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19	CWAS for systemic sclerosis identifies multiple risk loci and highlights fibrotic and vasculopathy pathways. <i>Nature Communications</i> , 2019, 10, 4955.	12.8	100
20	Identification of rheumatoid arthritis causal genes using functional genomics. <i>Scandinavian Journal of Immunology</i> , 2019, 89, e12753.	2.7	14
21	Common Genetic Component in Autoimmunity. <i>Rare Diseases of the Immune System</i> , 2019, , 221-236.	0.1	0
22	Chromatin interactions reveal novel gene targets for drug repositioning in rheumatic diseases. <i>Annals of the Rheumatic Diseases</i> , 2019, 78, 1127-1134.	0.9	23
23	OP0191â€¦EPIGENETIC PROFILING OF SYNOVIAL FIBROBLASTS REVEALS STRUCTURAL DNA DYNAMICS AT DISEASE IMPLICATED CHROMOSOME REGIONS. , 2019, , .		0
24	OP0192â€¦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
25	The genetics revolution in rheumatology: large scale genomic arrays and genetic mapping. <i>Nature Reviews Rheumatology</i> , 2017, 13, 421-432.	8.0	30
26	Analysis of SNP-SNP interactions and bone quantitative ultrasound parameter in early adulthood. <i>BMC Medical Genetics</i> , 2017, 18, 107.	2.1	8
27	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
28	Identifying Causal Genes at the Multiple Sclerosis Associated Region 6q23 Using Capture Hi-C. <i>PLoS ONE</i> , 2016, 11, e0166923.	2.5	28
29	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
30	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
31	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
32	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
33	Sex-specific differences in effect size estimates at established complex trait loci. <i>International Journal of Epidemiology</i> , 2012, 41, 1376-1382.	1.9	19
34	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
35	Genetics of rheumatoid arthritis: GWAS and beyond. <i>Open Access Rheumatology: Research and Reviews</i> , 2011, 3, 31.	1.6	22
36	HLA-DPB1-COL11A2 and three additional xMHC loci are independently associated with RA in a UK cohort. <i>Genes and Immunity</i> , 2011, 12, 169-175.	4.1	15

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37	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
38	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
39	Rare variation at the TNFAIP3 locus and susceptibility to rheumatoid arthritis. <i>Human Genetics</i> , 2010, 128, 627-633.	3.8	29
40	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
41	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
42	Synthetic associations in the context of genome-wide association scan signals. <i>Human Molecular Genetics</i> , 2010, 19, R137-R144.	2.9	53
43	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
44	Update on the genetic risk factors for rheumatoid arthritis. <i>Expert Review of Clinical Immunology</i> , 2010, 6, 61-75.	3.0	34
45	Novel Association of the Interleukin 2 Interleukin 21 Region With Inflammatory Bowel Disease. <i>American Journal of Gastroenterology</i> , 2009, 104, 1968-1975.	0.4	51
46	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
47	Study of functional variants of the BANK1 gene in rheumatoid arthritis. <i>Arthritis and Rheumatism</i> , 2009, 60, 372-379.	6.7	50
48	PTPN22C1858T polymorphism and human brucellosis. <i>Scandinavian Journal of Infectious Diseases</i> , 2009, 41, 109-112.	1.5	8
49	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
50	Association of STAT4 with rheumatoid arthritis: A replication study in three European populations. <i>Arthritis and Rheumatism</i> , 2008, 58, 1974-1980.	6.7	93
51	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
52	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
53	Caspase 7 influences susceptibility to rheumatoid arthritis. <i>Rheumatology</i> , 2007, 46, 1243-1247.	1.9	27
54	Auto-antibodies, HLA and PTPN22: susceptibility markers for rheumatoid arthritis. <i>Rheumatology</i> , 2007, 47, 138-141.	1.9	40

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55	Investigation of the IL23R gene in a Spanish rheumatoid arthritis cohort. <i>Human Immunology</i> , 2007, 68, 681-684.	2.4	52
56	Macrophage migration inhibitory factor gene: Influence on rheumatoid arthritis susceptibility. <i>Human Immunology</i> , 2007, 68, 744-747.	2.4	50
57	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
58	Genetic basis of rheumatoid arthritis. <i>Biomedicine and Pharmacotherapy</i> , 2006, 60, 656-662.	5.6	77
59	Asporin repeat polymorphism in rheumatoid arthritis. <i>Annals of the Rheumatic Diseases</i> , 2006, 66, 118-120.	0.9	27
60	Study of the role of functional variants of SLC22A4, RUNX1 and SUMO4 in systemic lupus erythematosus. <i>Annals of the Rheumatic Diseases</i> , 2006, 65, 791-795.	0.9	14
61	Study of the role of a functional polymorphism of MHC2TA in rheumatoid arthritis in three ethnically different populations. <i>Rheumatology</i> , 2006, 45, 1442-1444.	1.9	15
62	Epistatic interaction between FCRL3 and NF κ B1 genes in Spanish patients with rheumatoid arthritis. <i>Annals of the Rheumatic Diseases</i> , 2006, 65, 1188-1191.	0.9	59
63	Lack of association between ankylosing spondylitis and a functional polymorphism of PTPN22 proposed as a general susceptibility marker for autoimmunity. <i>Annals of the Rheumatic Diseases</i> , 2006, 65, 687-688.	0.9	23
64	SLC22A4, RUNX1, and SUMO4 polymorphisms are not associated with rheumatoid arthritis: a case-control study in a Spanish population. <i>Journal of Rheumatology</i> , 2006, 33, 1235-9.	2.0	21
65	Analysis of the functional NF κ B1 promoter polymorphism in rheumatoid arthritis and systemic lupus erythematosus. <i>Tissue Antigens</i> , 2005, 65, 183-186.	1.0	35
66	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
67	Association of a functional single nucleotide polymorphism of PTPN22, encoding lymphoid protein phosphatase, with rheumatoid arthritis and systemic lupus erythematosus. <i>Arthritis and Rheumatism</i> , 2005, 52, 219-224.	6.7	275
68	Protein tyrosine phosphatase gene (PTPN22) polymorphism in multiple sclerosis. <i>Journal of Neurology</i> , 2005, 252, 994-995.	3.6	38
69	Interleukin 12 (IL12B) and Interleukin 12 Receptor (IL12RB1) Gene Polymorphisms in Rheumatoid Arthritis. <i>Human Immunology</i> , 2005, 66, 710-714.	2.4	32
70	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
71	Analysis of a GT Microsatellite in the Promoter of the foxp3/scurfin Gene in Autoimmune Diseases. <i>Human Immunology</i> , 2005, 66, 869-873.	2.4	25
72	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

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73	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
74	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
75	Cytotoxic T-lymphocyte antigen-4-CT60 polymorphism in rheumatoid arthritis. <i>Tissue Antigens</i> , 2004, 64, 667-670.	1.0	34
76	Inducible nitric oxide synthase promoter polymorphism in human brucellosis. <i>Microbes and Infection</i> , 2003, 5, 1165-1169.	1.9	17