Juan Sainz

List of Publications by Year
in descending order

89 papers	1,559 citations	304368 22 h-index	344852 36 g-index
95 all docs	95 docs citations	95 times ranked	2630 citing authors

#	Article	IF	Citations
1	A polygenic risk score for multiple myeloma risk prediction. European Journal of Human Genetics, 2022, 30, 474-479.	1.4	5
2	Type 2 Diabetes-Related Variants Influence the Risk of Developing Prostate Cancer: A Population-Based Case-Control Study and Meta-Analysis. Cancers, 2022, 14, 2376.	1.7	6
3	Validation and functional characterization of GWAS-identified variants for chronic lymphocytic leukemia: a CRuCIAL study. Blood Cancer Journal, 2022, 12, 79.	2.8	1
4	Common gene variants within 3â€²â€untranslated regions as modulators of multiple myeloma risk and survival. International Journal of Cancer, 2021, 148, 1887-1894.	2.3	3
5	Do myeloproliferative neoplasms and multiple myeloma share the same genetic susceptibility loci?. International Journal of Cancer, 2021, 148, 1616-1624.	2.3	7
6	Expression quantitative trait loci of genes predicting outcome are associated with survival of multiple myeloma patients. International Journal of Cancer, 2021, 149, 327-336.	2.3	3
7	Functional Genetic Variants in ATG10 Are Associated with Acute Myeloid Leukemia. Cancers, 2021, 13, 1344.	1.7	4
8	Polymorphisms within Autophagy-Related Genes Influence the Risk of Developing Colorectal Cancer: A Meta-Analysis of Four Large Cohorts. Cancers, 2021, 13, 1258.	1.7	3
9	Genetically determined telomere length and multiple myeloma risk and outcome. Blood Cancer Journal, 2021, 11, 74.	2.8	10
10	OP0017â€...VALIDATION OF GWAS-IDENTIFIED VARIANTS FOR ANTI-TNF DRUG RESPONSE IN RHEUMATOID ARTHRITIS: A META-ANALYSIS OF THREE LARGE COHORTS. Annals of the Rheumatic Diseases, 2021, 80, 9.2-10.	0.5	1
11	Polymorphisms within the TNFSF4 and MAPKAPK2 Loci Influence the Risk of Developing Invasive Aspergillosis: A Two-Stage Case Control Study in the Context of the aspBIOmics Consortium. Journal of Fungi (Basel, Switzerland), 2021, 7, 4.	1.5	5
12	Validation of GWAS-Identified Variants for Anti-TNF Drug Response in Rheumatoid Arthritis: A Meta-Analysis of Two Large Cohorts. Frontiers in Immunology, 2021, 12, 672255.	2.2	6
13	Host immune genetic variations influence the risk of developing acute myeloid leukaemia: results from the NuCLEAR consortium. Blood Cancer Journal, 2020, 10, 75.	2.8	2
14	Genetic polymorphisms associated with telomere length and risk of developing myeloproliferative neoplasms. Blood Cancer Journal, 2020, 10, 89.	2.8	20
15	NFKB2 polymorphisms associate with the risk of developing rheumatoid arthritis and response to TNF inhibitors: Results from the REPAIR consortium. Scientific Reports, 2020, 10, 4316.	1.6	14
16	Polymorphisms within the ARNT2 and CX3CR1 Genes Are Associated with the Risk of Developing Invasive Aspergillosis. Infection and Immunity, 2020, 88, .	1.0	8
17	Steroid hormone-related polymorphisms associate with the development of bone erosions in rheumatoid arthritis and help to predict disease progression: Results from the REPAIR consortium. Scientific Reports, 2019, 9, 14812.	1.6	7
18	Exome sequencing identifies germline variants in DIS3 in familial multiple myeloma. Leukemia, 2019, 33, 2324-2330.	3.3	33

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19	Genetic polymorphisms in genes of class switch recombination and multiple myeloma risk and survival: an IMMEnSE study. Leukemia and Lymphoma, 2019, 60, 1803-1811.	0.6	11
20	Polymorphisms at phase I-metabolizing enzyme and hormone receptor loci influence the response to anti-TNF therapy in rheumatoid arthritis patients. Pharmacogenomics Journal, 2019, 19, 83-96.	0.9	10
21	Inherited variation in the xenobiotic transporter pathway and survival of multiple myeloma patients. British Journal of Haematology, 2018, 183, 375-384.	1.2	11
22	THU0010â€...Polymorphisms in phase i-metabolising enzyme and hormone receptor genes influence the response to anti-tnf therapy. , 2018, , .		0
23	IL-10 overexpression predisposes to invasive aspergillosis by suppressing antifungal immunity. Journal of Allergy and Clinical Immunology, 2017, 140, 867-870.e9.	1.5	37
24	Identification of miRSNPs associated with the risk of multiple myeloma. International Journal of Cancer, 2017, 140, 526-534.	2.3	8
25	Common Genetic Polymorphisms within NFÎºB-Related Genes and the Risk of Developing Invasive Aspergillosis. Frontiers in Microbiology, 2016, 7, 1243.	1.5	13
26	Early mortality in multiple myeloma: the timeâ€dependent impact of comorbidity: A populationâ€based study in 621 realâ€life patients. American Journal of Hematology, 2016, 91, 700-704.	2.0	28
27	Polymorphisms in Host Immunity-Modulating Genes and Risk of Invasive Aspergillosis: Results from the AspBIOmics Consortium. Infection and Immunity, 2016, 84, 643-657.	1.0	35
28	A common variant within the HNF1B gene is associated with overall survival of multiple myeloma patients: Results from the IMMEnSE consortium and meta-analysis. Oncotarget, 2016, 7, 59029-59048.	0.8	16
29	Genetic variants within immune-modulating genes influence the risk of developing rheumatoid arthritis and anti-TNF drug response. Pharmacogenetics and Genomics, 2015, 25, 432-443.	0.7	14
30	THU0002â€...Estrogen-Related Polymorphisms and Risk of Rheumatoid Arthritis: A Multicenter Study. Annals of the Rheumatic Diseases, 2015, 74, 193.2-193.	0.5	0
31	Genetic variants within the TNFRSF1B gene and susceptibility to rheumatoid arthritis and response to anti-TNF drugs. Pharmacogenetics and Genomics, 2015, 25, 323-333.	0.7	17
32	ISS Versus R-ISS for Risk Stratification of Multiple Myeloma Patients undergoing Autologous Stem Cell Transplant. Journal of Leukemia (Los Angeles, Calif), 2015, 03, .	0.1	2
33	TRAF1/C5but NotPTPRCVariants Are Potential Predictors of Rheumatoid Arthritis Response to Anti-Tumor Necrosis Factor Therapy. BioMed Research International, 2015, 2015, 1-9.	0.9	15
34	Genome-wide association study identifies variants at 16p13 associated with survival in multiple myeloma patients. Nature Communications, 2015, 6, 7539.	5.8	38
35	Transplantation-Associated Thrombotic Microangiopathy in Patients Treated With Sirolimus and Cyclosporine as Salvage Therapy for Graft-Versus-Host Disease. Annals of Pharmacotherapy, 2015, 49, 986-994.	0.9	20
36	Trends in survival of multiple myeloma: A thirty-year population-based study in a single institution. Cancer Epidemiology, 2015, 39, 693-699.	0.8	24

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37	Type 2 diabetes-related variants influence the risk of developing multiple myeloma: results from the IMMEnSE consortium. Endocrine-Related Cancer, 2015, 22, 545-559.	1.6	11
38	The Evolving Role of Stem Cell Transplant in Multiple Myeloma: A Single Institution Study. Clinical Lymphoma, Myeloma and Leukemia, 2015, 15, e170-e171.	0.2	0
39	Risk of multiple myeloma is associated with polymorphisms within telomerase genes and telomere length. International Journal of Cancer, 2015, 136, E351-8.	2.3	30
40	Light Chain Multiple Myeloma: A Single Institution Series. Journal of Leukemia (Los Angeles, Calif), 2015, 03, .	0.1	2
41	Comparative Baseline Health-Related Quality of Life in Real-Life Patients with Monoclonal Gammopathies. Journal of Leukemia (Los Angeles, Calif), 2015, 03, .	0.1	1
42	Smoldering Multiple Myeloma: Changing the Management Paradigm or Just the Definition ?. Journal of Leukemia (Los Angeles, Calif), 2014, 02, .	0.1	0
43	GWAS-Identified Common Variants for Obesity Are Not Associated with the Risk of Developing Colorectal Cancer. Cancer Epidemiology Biomarkers and Prevention, 2014, 23, 1125-1128.	1.1	3
44	Repeat polymorphisms in ESR2 and ARand colorectal cancer risk and prognosis: results from a German population-based case-control study. BMC Cancer, 2014, 14, 817.	1.1	16
45	Genetic Variants and Multiple Myeloma Risk: IMMEnSE Validation of the Best Reported Associationsâ€"An Extensive Replication of the Associations from the Candidate Gene Era. Cancer Epidemiology Biomarkers and Prevention, 2014, 23, 670-674.	1.1	13
46	THU0474â€...Association of Il4, IL4R and IL8RB Gene Polymorphisms with the Risk of Developing Rheumatoid Arthritis. Annals of the Rheumatic Diseases, 2014, 73, 347.2-347.	0.5	0
47	Obesity and Multiple Myeloma: What Do the Data Tell Us?. Journal of Leukemia (Los Angeles, Calif), 2014, 02, .	0.1	2
48	Type 2 Diabetes and Multiple Myeloma: The Latest Insights. Journal of Leukemia (Los Angeles, Calif), 2014, 02, .	0.1	1
49	Abstract 5078: Genome wide association study identifies variants at 16p13 associated with survival in multiple myeloma patients. , 2014, , .		0
50	Type 2 Diabetes-Related Variants Influence on the Risk of Developing Multiple Myeloma: Results from the Immense Consortium. Blood, 2014, 124, 2044-2044.	0.6	0
51	The International Multiple Myeloma Research (IMMEnSE) Consortium: Genetics of Multiple Myeloma Risk and Prognosis. Blood, 2014, 124, 3421-3421.	0.6	0
52	Polymorphisms in regulators of xenobiotic transport and metabolism genes PXR and CAR do not affect multiple myeloma risk: a caseâ€"control study in the context of the IMMEnSE consortium. Journal of Human Genetics, 2013, 58, 155-159.	1.1	5
53	Impacto del tipo de hospital en la supervivencia de pacientes con mieloma mÃºltiple: estudio MICORE. Revista Clinica Espanola, 2013, 213, 330-335.	0.2	2
54	The impact of the type of hospital on survival of multiple myeloma patients: The MICORE study. Revista Clínica Espanõla, 2013, 213, 330-335.	0.3	0

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55	Genetic variants in Câ€type lectin genes are associated with colorectal cancer susceptibility and clinical outcome. International Journal of Cancer, 2013, 133, 2325-2333.	2.3	28
56	AB0009â€...Gender-associated differences of dectin-2, dc-sign and mcp-1 polymorphisms in the susceptibility to rheumatoid arthritis. Annals of the Rheumatic Diseases, 2013, 72, A788.2-A788.	0.5	0
57	FRI0183â€...Long-term survival of tumor necrosis factor-alpha inhibitor therapies in a spanish cohort of rheumatoid arthritis patients. Annals of the Rheumatic Diseases, 2013, 72, A434.1-A434.	0.5	0
58	Gender-Specific Effects of Genetic Variants within Th1 and Th17 Cell-Mediated Immune Response Genes on the Risk of Developing Rheumatoid Arthritis. PLoS ONE, 2013, 8, e72732.	1.1	20
59	Polymorphisms in xenobiotic transporters ABCB1, ABCG2, ABCC2, ABCC1, ABCC3 and multiple myeloma risk: a caseâ€"control study in the context of the International Multiple Myeloma rESEarch (IMMEnSE) consortium. Leukemia, 2012, 26, 1419-1422.	3.3	14
60	Impact of polymorphic variation at 7p15.3, 3p22.1 and 2p23.3 loci on risk of multiple myeloma. British Journal of Haematology, 2012, 158, 805-809.	1.2	19
61	Effect of Type 2 Diabetes Predisposing Genetic Variants on Colorectal Cancer Risk. Journal of Clinical Endocrinology and Metabolism, 2012, 97, E845-E851.	1.8	56
62	Comprehensive investigation of genetic variation in the 8q24 region and multiple myeloma risk in the <scp>IMME</scp>n<scp>SE</scp> consortium. British Journal of Haematology, 2012, 157, 331-338.	1.2	13
63	Dectin-1 and DC-SIGN Polymorphisms Associated with Invasive Pulmonary Aspergillosis Infection. PLoS ONE, 2012, 7, e32273.	1.1	126
64	A Comprehensive Investigation on Common Polymorphisms in the MDR1/ABCB1 Transporter Gene and Susceptibility to Colorectal Cancer. PLoS ONE, 2012, 7, e32784.	1.1	30
65	Genome-wide gene expression analysis in mouse embryonic stem cells. International Journal of Developmental Biology, 2011, 55, 995-1006.	0.3	12
66	P1-507 Polymorphisms in genes related to sex steroid transport and signalling modulate menopausal hormone therapy effect on risk of colorectal cancer. Journal of Epidemiology and Community Health, 2011, 65, A207-A207.	2.0	0
67	Genetics and molecular epidemiology of multiple myeloma: The rationale for the IMMEnSE consortium (Review). International Journal of Oncology, 2011, 40, 625-38.	1.4	14
68	Modification of menopausal hormone therapy-associated colorectal cancer risk by polymorphisms in sex steroid signaling, metabolism and transport related genes. Endocrine-Related Cancer, 2011, 18, 371-384.	1.6	23
69	Association of genetic polymorphisms in ESR2, HSD17B1, ABCB1, and SHBG genes with colorectal cancer risk. Endocrine-Related Cancer, 2011, 18, 265-276.	1.6	59
70	Polymorphisms in Regulators of Xenobiotic Transport and Metabolism Genes NR1I2 and NR1I3 and Multiple Myeloma Risk: A Case-Control Study in the Context of IMMEnSE Consortium. Blood, 2011, 118, 5014-5014.	0.6	0
71	TNFR1 mRNA Expression Level and TNFR1 Gene Polymorphisms are Predictive Markers for Susceptibility to Develop Invasive Pulmonary Aspergillosis. International Journal of Immunopathology and Pharmacology, 2010, 23, 423-436.	1.0	53
72	Abstract 2855: Estrogen transport, metabolism, binding and signalling related single nucleotide polymorphisms modulate menopausal hormone therapy effect on risk of colorectal cancer. , 2010, , .		0

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73	IL1 Gene Cluster Polymorphisms and Its Haplotypes may Predict the Risk to Develop Invasive Pulmonary Aspergillosis and Modulate C-reactive Protein Level. Journal of Clinical Immunology, 2008, 28, 473-485.	2.0	81
74	Genetic variants of IL6 gene promoter influence on C-reactive protein levels but are not associated with susceptibility to invasive pulmonary aspergillosis in haematological patients. Cytokine, 2008, 41, 268-278.	1.4	25
75	Variable Number of Tandem Repeats of TNF Receptor Type 2 Promoter as Genetic Biomarker of Susceptibility to Develop Invasive Pulmonary Aspergillosis. Human Immunology, 2007, 68, 41-50.	1.2	78
76	Interleukin-10 promoter polymorphism as risk factor to develop invasive pulmonary aspergillosis. Immunology Letters, 2007, 109, 76-82.	1.1	77
77	Antioxidant Enzymes and Effects of Tempol on the Development of Hypertension Induced by Nitric Oxide Inhibition. American Journal of Hypertension, 2005, 18, 871-877.	1.0	41
78	Role of sex, gonadectomy and sex hormones in the development of nitric oxide inhibition-induced hypertension. Experimental Physiology, 2004, 89, 155-162.	0.9	26
79	Gender difference in the role of endothelium-derived relaxing factors modulating renal vascular reactivity. European Journal of Pharmacology, 2004, 486, 281-288.	1.7	28
80	Effects of Nitric Oxide on Aldosterone Synthesis and Nitric Oxide Synthase Activity in Glomerulosa Cells from Bovine Adrenal Gland. Endocrine, 2004, 24, 061-072.	2.2	19
81	Protective effects of the angiotensin II type I (AT1) receptor blockade in low-renin deoxycorticosterone acetate (DOCA)-treated spontaneously hypertensive rats. Clinical Science, 2004, 106, 251-259.	1.8	15
82	ANTIOXIDANT ENZYMES AND EFFECTS OF TEMPOL, A RADICAL SCAVENGER, ON THE DEVELOPMENT OF NO INHIBITION-INDUCED HYPERTENSION. Journal of Hypertension, 2004, 22, S63.	0.3	0
83	Chronic Blockade of Neuronal Nitric Oxide Synthase Does Not Affect Long-Term Control of Blood Pressure in Normal, Saline-Drinking or Deoxycorticosterone-Treated Rats. Experimental Physiology, 2003, 88, 243-250.	0.9	10
84	Increased Pressor Sensitivity to Chronic Nitric Oxide Deficiency in Hyperthyroid Rats. Hypertension, 2003, 42, 220-225.	1.3	33
85	Role of endothelium-derived relaxing factors in the renal response to vasoactive agents in hypothyroid rats. American Journal of Physiology - Endocrinology and Metabolism, 2003, 285, E182-E188.	1.8	19
86	Nitric oxide synthase activity in hyperthyroid and hypothyroid rats. European Journal of Endocrinology, 2002, 147, 117-122.	1.9	84
87	Role of endothelium-derived relaxing factors in adrenomedullin-induced vasodilation in the rat kidney. European Journal of Pharmacology, 2002, 444, 97-102.	1.7	15
88	Contribution of endothelium-derived relaxing factors to P2Y-purinoceptor-induced vasodilation in the isolated rat kidney. General Pharmacology, 2000, 35, 129-133.	0.7	9
89	Does a Multiple Myeloma Polygenic Risk Score Predict Overall Survival of Myeloma Patients?. Cancer Epidemiology Biomarkers and Prevention, 0, , .	1.1	2