

Elena Sanchez

List of Publications by Year in descending order

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52
papers

3,546
citations

126858

33
h-index

161767

54
g-index

54
all docs

54
docs citations

54
times ranked

5657
citing authors

#	ARTICLE	IF	CITATIONS
1	Phenotypic and genotypic characterization of families with complex intellectual disability identified pathogenic genetic variations in known and novel disease genes. <i>Scientific Reports</i> , 2020, 10, 968.	1.6	8
2	Trans-ethnic kidney function association study reveals putative causal genes and effects on kidney-specific disease aetiologies. <i>Nature Communications</i> , 2019, 10, 29.	5.8	113
3	Pla2g6 Deficiency in Zebrafish Leads to Dopaminergic Cell Death, Axonal Degeneration, Increased β -Synuclein Expression, and Defects in Brain Functions and Pathways. <i>Molecular Neurobiology</i> , 2018, 55, 6734-6754.	1.9	17
4	Effects of Amerindian Genetic Ancestry on Clinical Variables and Therapy in Patients with Rheumatoid Arthritis. <i>Journal of Rheumatology</i> , 2017, 44, 1804-1812.	1.0	1
5	A Novel p.Glu298Lys Mutation in the ACMSD Gene in Sporadic Parkinson's Disease. <i>Journal of Parkinson's Disease</i> , 2017, 7, 459-463.	1.5	15
6	Identification of a Large DNAJB2 Deletion in a Family with Spinal Muscular Atrophy and Parkinsonism. <i>Human Mutation</i> , 2016, 37, 1180-1189.	1.1	36
7	Genomic Insights into the Ancestry and Demographic History of South America. <i>PLoS Genetics</i> , 2015, 11, e1005602.	1.5	198
8	<i>SORT1</i> Mutation Resulting in Sortilin Deficiency and p75 ^{NTR} Upregulation in a Family With Essential Tremor. <i>ASN Neuro</i> , 2015, 7, 175909141559829.	1.5	28
9	<i>SCN4A</i> pore mutation pathogenetically contributes to autosomal dominant essential tremor and may increase susceptibility to epilepsy. <i>Human Molecular Genetics</i> , 2015, 24, ddv410.	1.4	38
10	Intronic Variants in the NFKB1 Gene May Influence Hearing Forecast in Patients with Unilateral Sensorineural Hearing Loss in Meniere's Disease. <i>PLoS ONE</i> , 2014, 9, e112171.	1.1	37
11	Allelic heterogeneity in NCF2 associated with systemic lupus erythematosus (SLE) susceptibility across four ethnic populations. <i>Human Molecular Genetics</i> , 2014, 23, 1656-1668.	1.4	67
12	Fine mapping of Xq28: both <i>MECP2</i> and <i>IRAK1</i> contribute to risk for systemic lupus erythematosus in multiple ancestral groups. <i>Annals of the Rheumatic Diseases</i> , 2013, 72, 437-444.	0.5	97
13	Evidence of New Risk Genetic Factor to Systemic Lupus Erythematosus: The UBASH3A Gene. <i>PLoS ONE</i> , 2013, 8, e60646.	1.1	27
14	Genetic and physical interaction of the B-cell systemic lupus erythematosus-associated genes <i>BANK1</i> and <i>BLK</i> . <i>Annals of the Rheumatic Diseases</i> , 2012, 71, 136-142.	0.5	67
15	Analysis of autosomal genes reveals gene-sex interactions and higher total genetic risk in men with systemic lupus erythematosus. <i>Annals of the Rheumatic Diseases</i> , 2012, 71, 694-699.	0.5	87
16	Impact of genetic ancestry and sociodemographic status on the clinical expression of systemic lupus erythematosus in American Indian-European populations. <i>Arthritis and Rheumatism</i> , 2012, 64, 3687-3694.	6.7	70
17	Association study of <i>IRAK-M</i> and <i>SIGIRR</i> genes with SLE in a large European-descent population. <i>Lupus</i> , 2012, 21, 1166-1171.	0.8	11
18	Evidence for gene-gene epistatic interactions among susceptibility loci for systemic lupus erythematosus. <i>Arthritis and Rheumatism</i> , 2012, 64, 485-492.	6.7	53

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19	Fine-mapping and transethnic genotyping establish IL2/IL21 genetic association with lupus and localize this genetic effect to IL21. <i>Arthritis and Rheumatism</i> , 2011, 63, 1689-1697.	6.7	49
20	Identification of novel genetic susceptibility loci in African American lupus patients in a candidate gene association study. <i>Arthritis and Rheumatism</i> , 2011, 63, 3493-3501.	6.7	109
21	Early disease onset is predicted by a higher genetic risk for lupus and is associated with a more severe phenotype in lupus patients. <i>Annals of the Rheumatic Diseases</i> , 2011, 70, 151-156.	0.5	155
22	Phenotypic associations of genetic susceptibility loci in systemic lupus erythematosus. <i>Annals of the Rheumatic Diseases</i> , 2011, 70, 1752-1757.	0.5	110
23	A 3' untranslated region variant is associated with impaired expression of <i>CD226</i> in T and natural killer T cells and is associated with susceptibility to systemic lupus erythematosus. <i>Arthritis and Rheumatism</i> , 2010, 62, 3404-3414.	6.7	48
24	Genetically determined Amerindian ancestry correlates with increased frequency of risk alleles for systemic lupus erythematosus. <i>Arthritis and Rheumatism</i> , 2010, 62, 3722-3729.	6.7	70
25	Recent findings on genetics of systemic autoimmune diseases. <i>Current Opinion in Immunology</i> , 2010, 22, 698-705.	2.4	78
26	The TRAF1-C5 region on chromosome 9q33 is associated with multiple autoimmune diseases. <i>Annals of the Rheumatic Diseases</i> , 2010, 69, 696-699.	0.5	49
27	Promoter Insertion/Deletion in the <i>IRF5</i> Gene Is Highly Associated with Susceptibility to Systemic Lupus Erythematosus in Distinct Populations, But Exerts a Modest Effect on Gene Expression in Peripheral Blood Mononuclear Cells. <i>Journal of Rheumatology</i> , 2010, 37, 574-578.	1.0	32
28	Impact of interleukin-18 polymorphisms-607 and -137 on clinical characteristics of renal cell carcinoma patients. <i>Human Immunology</i> , 2010, 71, 309-313.	1.2	27
29	STAT4 associates with systemic lupus erythematosus through two independent effects that correlate with gene expression and act additively with IRF5 to increase risk. <i>Annals of the Rheumatic Diseases</i> , 2009, 68, 1746-1753.	0.5	138
30	Identification of a new putative functional IL18 gene variant through an association study in systemic lupus erythematosus. <i>Human Molecular Genetics</i> , 2009, 18, 3739-3748.	1.4	54
31	Replication of the TNFSF4 (OX40L) promoter region association with systemic lupus erythematosus. <i>Genes and Immunity</i> , 2009, 10, 248-253.	2.2	41
32	No evidence for genetic association of interferon regulatory factor 3 in systemic lupus erythematosus. <i>Lupus</i> , 2009, 18, 230-234.	0.8	13
33	Kallikrein genes are associated with lupus and glomerular basement membrane-specific antibody-induced nephritis in mice and humans. <i>Journal of Clinical Investigation</i> , 2009, 119, 911-923.	3.9	114
34	Functional variants in the B-cell gene BANK1 are associated with systemic lupus erythematosus. <i>Nature Genetics</i> , 2008, 40, 211-216.	9.4	436
35	Investigating the role of CD24 gene polymorphisms in rheumatoid arthritis. <i>Annals of the Rheumatic Diseases</i> , 2008, 67, 1197-1198.	0.5	18
36	MYO9B gene polymorphisms are associated with autoimmune diseases in Spanish population. <i>Human Immunology</i> , 2007, 68, 610-615.	1.2	33

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37	Association of a <i>CD24</i> gene polymorphism with susceptibility to systemic lupus erythematosus. <i>Arthritis and Rheumatism</i> , 2007, 56, 3080-3086.	6.7	47
38	Macrophage migration inhibitory factor gene influences the risk of developing tuberculosis in northwestern Colombian population. <i>Tissue Antigens</i> , 2007, 70, 28-33.	1.0	35
39	Analysis of interleukin-23 receptor (IL23R) gene polymorphisms in systemic lupus erythematosus. <i>Tissue Antigens</i> , 2007, 70, 233-237.	1.0	41
40	Association study of genetic variants of pro-inflammatory chemokine and cytokine genes in systemic lupus erythematosus. <i>BMC Medical Genetics</i> , 2006, 7, 48.	2.1	42
41	Evidence of association of macrophage migration inhibitory factor gene polymorphisms with systemic lupus erythematosus. <i>Genes and Immunity</i> , 2006, 7, 433-436.	2.2	91
42	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.	0.5	14
43	Study of a functional polymorphism in the p53 gene in systemic lupus erythematosus: lack of replication in a Spanish population. <i>Lupus</i> , 2006, 15, 658-661.	0.8	12
44	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.	0.5	59
45	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
46	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
47	Analysis of a GT Microsatellite in the Promoter of the foxp3/scurfin Gene in Autoimmune Diseases. <i>Human Immunology</i> , 2005, 66, 869-873.	1.2	25
48	Analysis of a Functional BTNL2 Polymorphism in Type 1 Diabetes, Rheumatoid Arthritis, and Systemic Lupus Erythematosus. <i>Human Immunology</i> , 2005, 66, 1235-1241.	1.2	70
49	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
50	Absence of COCH mutations in patients with Meniere disease. <i>European Journal of Human Genetics</i> , 2004, 12, 75-78.	1.4	37
51	Association of the CT60 marker of the CTLA4 gene with systemic lupus erythematosus. <i>Arthritis and Rheumatism</i> , 2004, 50, 2211-2215.	6.7	81
52	Inducible nitric oxide synthase promoter polymorphism in human brucellosis. <i>Microbes and Infection</i> , 2003, 5, 1165-1169.	1.0	17