Paula Sandrin-Garcia

List of Publications by Year
in descending order

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42 papers	851 citations	516561 16 h-index	526166 27 g-index
42 all docs	42 docs citations	42 times ranked	1479 citing authors

#	Article	IF	Citations
1	Polimorphisms in Inflammasome Genes Are Involved in the Predisposition to Systemic Lupus Erythematosus. Autoimmunity, 2012, 45, 271-278.	1.2	143
2	Gene Expression Profiles in Radiation Workers Occupationally Exposed to Ionizing Radiation. Journal of Radiation Research, 2009, 50, 61-71.	0.8	73
3	Gene Expression Profiles in Human Lymphocytes Irradiated In Vitro with Low Doses of Gamma Rays. Radiation Research, 2007, 168, 650.	0.7	59
4	Gene expression profiles in human cells submitted to genotoxic stress. Mutation Research - Reviews in Mutation Research, 2003, 544, 403-413.	2.4	53
5	Polymorphisms and expression of inflammasome genes are associated with the development and severity of rheumatoid arthritis in Brazilian patients. Inflammation Research, 2018, 67, 255-264.	1.6	45
6	Differential expression of the inflammasome complex genes in systemic lupus erythematosus. Immunogenetics, 2020, 72, 217-224.	1.2	31
7	Promiscuous Gene Expression in the Thymus: The Root of Central Tolerance. Clinical and Developmental Immunology, 2006, 13, 81-99.	3.3	28
8	The Role of NLRP3 Inflammasome in Lupus Nephritis. International Journal of Molecular Sciences, 2021, 22, 12476.	1.8	27
9	High burden of acute kidney injury in COVID-19 pandemic: systematic review and meta-analysis. Journal of Clinical Pathology, 2021, 74, 796-803.	1.0	26
10	Comprehensive gene expression profiling in lungs of mice infected with Mycobacterium tuberculosis following DNAhsp65 immunotherapy. Journal of Gene Medicine, 2009, 11, 66-78.	1.4	22
11	Mannose binding lectin gene (MBL2) functional polymorphisms are associated with systemic lupus erythematosus in southern Brazilians. Human Immunology, 2011, 72, 516-521.	1.2	22
12	Higher interferon score and normal complement levels may identify a distinct clinical subset in children with systemic lupus erythematosus. Arthritis Research and Therapy, 2020, 22, 91.	1.6	22
13	Typical phenotypic spectrum of velocardiofacial syndrome occurs independently of deletion size in chromosome 22q11.2. Molecular and Cellular Biochemistry, 2007, 303, 9-17.	1.4	20
14	Differential gene expression of peripheral blood mononuclear cells from rheumatoid arthritis patients may discriminate immunogenetic, pathogenic and treatment features. Immunology, 2009, 127, 365-372.	2.0	20
15	Polymorphisms in STK17A gene are associated with systemic lupus erythematosus and its clinical manifestations. Gene, 2013, 527, 435-439.	1.0	20
16	Ficolin Gene Polymorphisms in Systemic Lupus Erythematosus and Rheumatoid Arthritis. Annals of Human Genetics, 2016, 80, 1-6.	0.3	20
17	Vitamin D receptor (VDR) gene polymorphisms and age onset in type 1 diabetes mellitus. Autoimmunity, 2013, 46, 382-387.	1.2	19
18	Alterations in gene expression profiles correlated with cisplatin cytotoxicity in the glioma U343 cell line. Genetics and Molecular Biology, 2010, 33, 159-168.	0.6	17

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19	Effect of a Single Apolipoprotein L1 Gene Nephropathy Variant on the Risk of Advanced Lupus Nephritis in Brazilians. Journal of Rheumatology, 2020, 47, 1209-1217.	1.0	17
20	Are key cytokines genetic and serum levels variations related to rheumatoid arthritis clinical severity?. Gene, 2020, 722, 144098.	1.0	15
21	Vitamin D receptor polymorphisms and expression profile in rheumatoid arthritis brazilian patients. Molecular Biology Reports, 2016, 43, 41-51.	1.0	14
22	Gene Expression Profiles Stratified according to Type 1 Diabetes Mellitus Susceptibility Regions. Annals of the New York Academy of Sciences, 2008, 1150, 282-289.	1.8	13
23	Shared and Unique Gene Expression in Systemic Lupus Erythematosus Depending on Disease Activity. Annals of the New York Academy of Sciences, 2009, 1173, 493-500.	1.8	13
24	PTPN22 1858Câ€‰>â€‰T polymorphism and susceptibility to systemic lupus erythematosus: a meta-analysis update. Autoimmunity, 2017, 50, 428-434.	1.2	12
25	Postmenopausal Osteoporosis reference genes for qPCR expression assays. Scientific Reports, 2019, 9, 16533.	1.6	10
26	Delayed effects of exposure to a moderate radiation dose on transcription profiles in human primary fibroblasts. Environmental and Molecular Mutagenesis, 2011, 52, 117-129.	0.9	9
27	FYB gene polymorphisms are associated with susceptibility for systemic lupus erythemathosus (SLE). Human Immunology, 2013, 74, 1009-1014.	1.2	8
28	IL1Î², IL18, NFKB1 and IFNG gene interactions are associated with severity of rheumatoid arthritis: A pilot study. Autoimmunity, 2020, 53, 95-101.	1.2	8
29	Fluorescence in situ hybridization (FISH) screening for the 22q11.2 deletion in patients with clinical features of velocardiofacial syndrome but without cardiac anomalies. Genetics and Molecular Biology, 2007, 30, 21-24.	0.6	7
30	<i>Mannose-Binding Lectin2</i> Gene Polymorphism and IgG4 in Membranous Nephropathy. Nephron, 2018, 139, 181-188.	0.9	7
31	Metabolism Genes Are among the Differentially Expressed Ones Observed in Lymphomononuclear Cells of Recently Diagnosed Type 1 Diabetes Mellitus Patients. Annals of the New York Academy of Sciences, 2006, 1079, 171-176.	1.8	6
32	LIG4 and RAD52 DNA repair genes polymorphisms and systemic lupus erythematosus. Molecular Biology Reports, 2014, 41, 2249-2256.	1.0	6
33	CCR5Î”32and the genetic susceptibility to rheumatoid arthritis in admixed populations: a multicentre study. Rheumatology, 2017, 56, kew398.	0.9	6
34	ADA2 deficiency (DADA2) associated with Evans syndrome and a severe ADA2 genotype. Rheumatology, 2021, 60, e237-e239.	0.9	6
35	Using cDNA microarrays to identify human CD19+ B cell gene products (ESTs) originated from systemic lupus erythematosus susceptibility loci. Autoimmunity Reviews, 2006, 5, 319-323.	2.5	5
36	Is HLA Class II Profile Relevant for the Study of Large-Scale Differentially Expressed Genes in Type 1 Diabetes Mellitus Patients?. Annals of the New York Academy of Sciences, 2006, 1079, 305-309.	1.8	4

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37	T-cell specific upregulation of Sema4A as risk factor for autoimmunity in systemic lupus erythematosus and rheumatoid arthritis. Autoimmunity, 2020, 53, 65-70.	1.2	4
38	cDNA microarray analysis of cyclosporin A (CsA)-treated human peripheral blood mononuclear cells reveal modulation of genes associated with apoptosis, cell-cycle regulation and DNA repair. Molecular and Cellular Biochemistry, 2007, 304, 235-241.	1.4	3
39	CTLA-4 gene polymorphisms are associated with obesity in Turner Syndrome. Genetics and Molecular Biology, 2018, 41, 727-734.	0.6	3
40	Is there an Inflammation Role for MYD88 in Rheumatoid Arthritis?. Inflammation, 2021, 44, 1014-1022.	1.7	3
41	MYD88, IRAK3 and Rheumatoid Arthritis pathogenesis: Analysis of differential gene expression in CD14Â+Âmonocytes and the inflammatory cytokine levels. Immunobiology, 2021, 226, 152152.	0.8	3
42	Differential distribution of vitamin D receptor (VDR) gene variants and its expression in systemic lupus erythematosus. International Journal of Immunogenetics, 2022, 49, 181-192.	0.8	2