

Paula Sandrin-Garcia

List of Publications by Year in descending order

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papers

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516710

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1479
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#	ARTICLE	IF	CITATIONS
1	Polimorphisms in Inflammasome Genes Are Involved in the Predisposition to Systemic Lupus Erythematosus. <i>Autoimmunity</i> , 2012, 45, 271-278.	2.6	143
2	Gene Expression Profiles in Radiation Workers Occupationally Exposed to Ionizing Radiation. <i>Journal of Radiation Research</i> , 2009, 50, 61-71.	1.6	73
3	Gene Expression Profiles in Human Lymphocytes Irradiated In Vitro with Low Doses of Gamma Rays. <i>Radiation Research</i> , 2007, 168, 650.	1.5	59
4	Gene expression profiles in human cells submitted to genotoxic stress. <i>Mutation Research - Reviews in Mutation Research</i> , 2003, 544, 403-413.	5.5	53
5	Polymorphisms and expression of inflammasome genes are associated with the development and severity of rheumatoid arthritis in Brazilian patients. <i>Inflammation Research</i> , 2018, 67, 255-264.	4.0	45
6	Differential expression of the inflammasome complex genes in systemic lupus erythematosus. <i>Immunogenetics</i> , 2020, 72, 217-224.	2.4	31
7	Promiscuous Gene Expression in the Thymus: The Root of Central Tolerance. <i>Clinical and Developmental Immunology</i> , 2006, 13, 81-99.	3.3	28
8	The Role of NLRP3 Inflammasome in Lupus Nephritis. <i>International Journal of Molecular Sciences</i> , 2021, 22, 12476.	4.1	27
9	High burden of acute kidney injury in COVID-19 pandemic: systematic review and meta-analysis. <i>Journal of Clinical Pathology</i> , 2021, 74, 796-803.	2.0	26
10	Comprehensive gene expression profiling in lungs of mice infected with <i>Mycobacterium tuberculosis</i> following DNAhsp65 immunotherapy. <i>Journal of Gene Medicine</i> , 2009, 11, 66-78.	2.8	22
11	Mannose binding lectin gene (MBL2) functional polymorphisms are associated with systemic lupus erythematosus in southern Brazilians. <i>Human Immunology</i> , 2011, 72, 516-521.	2.4	22
12	Higher interferon score and normal complement levels may identify a distinct clinical subset in children with systemic lupus erythematosus. <i>Arthritis Research and Therapy</i> , 2020, 22, 91.	3.5	22
13	Typical phenotypic spectrum of velocardiofacial syndrome occurs independently of deletion size in chromosome 22q11.2. <i>Molecular and Cellular Biochemistry</i> , 2007, 303, 9-17.	3.1	20
14	Differential gene expression of peripheral blood mononuclear cells from rheumatoid arthritis patients may discriminate immunogenetic, pathogenic and treatment features. <i>Immunology</i> , 2009, 127, 365-372.	4.4	20
15	Polymorphisms in STK17A gene are associated with systemic lupus erythematosus and its clinical manifestations. <i>Gene</i> , 2013, 527, 435-439.	2.2	20
16	Ficolin Gene Polymorphisms in Systemic Lupus Erythematosus and Rheumatoid Arthritis. <i>Annals of Human Genetics</i> , 2016, 80, 1-6.	0.8	20
17	Vitamin D receptor (VDR) gene polymorphisms and age onset in type 1 diabetes mellitus. <i>Autoimmunity</i> , 2013, 46, 382-387.	2.6	19
18	Alterations in gene expression profiles correlated with cisplatin cytotoxicity in the glioma U343 cell line. <i>Genetics and Molecular Biology</i> , 2010, 33, 159-168.	1.3	17

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19	Effect of a Single Apolipoprotein L1 Gene Nephropathy Variant on the Risk of Advanced Lupus Nephritis in Brazilians. <i>Journal of Rheumatology</i> , 2020, 47, 1209-1217.	2.0	17
20	Are key cytokines genetic and serum levels variations related to rheumatoid arthritis clinical severity?. <i>Gene</i> , 2020, 722, 144098.	2.2	15
21	Vitamin D receptor polymorphisms and expression profile in rheumatoid arthritis brazilian patients. <i>Molecular Biology Reports</i> , 2016, 43, 41-51.	2.3	14
22	Gene Expression Profiles Stratified according to Type 1 Diabetes Mellitus Susceptibility Regions. <i>Annals of the New York Academy of Sciences</i> , 2008, 1150, 282-289.	3.8	13
23	Shared and Unique Gene Expression in Systemic Lupus Erythematosus Depending on Disease Activity. <i>Annals of the New York Academy of Sciences</i> , 2009, 1173, 493-500.	3.8	13
24	<i>PTPN22</i> 1858Câ€™%>â€™%T polymorphism and susceptibility to systemic lupus erythematosus: a meta-analysis update. <i>Autoimmunity</i> , 2017, 50, 428-434.	2.6	12
25	Postmenopausal Osteoporosis reference genes for qPCR expression assays. <i>Scientific Reports</i> , 2019, 9, 16533.	3.3	10
26	Delayed effects of exposure to a moderate radiation dose on transcription profiles in human primary fibroblasts. <i>Environmental and Molecular Mutagenesis</i> , 2011, 52, 117-129.	2.2	9
27	FYB gene polymorphisms are associated with susceptibility for systemic lupus erythematosus (SLE). <i>Human Immunology</i> , 2013, 74, 1009-1014.	2.4	8
28	<i>IL1Î²</i>, <i>IL18</i>, <i>NFKB1</i> and <i>IFNG</i> gene interactions are associated with severity of rheumatoid arthritis: A pilot study. <i>Autoimmunity</i> , 2020, 53, 95-101.	2.6	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. <i>Genetics and Molecular Biology</i> , 2007, 30, 21-24.	1.3	7
30	<i>Mannose-Binding Lectin2</i> Gene Polymorphism and IgG4 in Membranous Nephropathy. <i>Nephron</i> , 2018, 139, 181-188.	1.8	7
31	Metabolism Genes Are among the Differentially Expressed Ones Observed in Lymphomononuclear Cells of Recently Diagnosed Type 1 Diabetes Mellitus Patients. <i>Annals of the New York Academy of Sciences</i> , 2006, 1079, 171-176.	3.8	6
32	LIG4 and RAD52 DNA repair genes polymorphisms and systemic lupus erythematosus. <i>Molecular Biology Reports</i> , 2014, 41, 2249-2256.	2.3	6
33	<i>CCR5Î³32</i> and the genetic susceptibility to rheumatoid arthritis in admixed populations: a multicentre study. <i>Rheumatology</i> , 2017, 56, kew398.	1.9	6
34	ADA2 deficiency (DADA2) associated with Evans syndrome and a severe <i>ADA2</i> genotype. <i>Rheumatology</i> , 2021, 60, e237-e239.	1.9	6
35	Using cDNA microarrays to identify human CD19+ B cell gene products (ESTs) originated from systemic lupus erythematosus susceptibility loci. <i>Autoimmunity Reviews</i> , 2006, 5, 319-323.	5.8	5
36	Is HLA Class II Profile Relevant for the Study of Large-Scale Differentially Expressed Genes in Type 1 Diabetes Mellitus Patients?. <i>Annals of the New York Academy of Sciences</i> , 2006, 1079, 305-309.	3.8	4

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37	T-cell specific upregulation of Sema4A as risk factor for autoimmunity in systemic lupus erythematosus and rheumatoid arthritis. <i>Autoimmunity</i> , 2020, 53, 65-70.	2.6	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. <i>Molecular and Cellular Biochemistry</i> , 2007, 304, 235-241.	3.1	3
39	CTLA-4 gene polymorphisms are associated with obesity in Turner Syndrome. <i>Genetics and Molecular Biology</i> , 2018, 41, 727-734.	1.3	3
40	Is there an Inflammation Role for MYD88 in Rheumatoid Arthritis?. <i>Inflammation</i> , 2021, 44, 1014-1022.	3.8	3
41	MYD88, IRAK3 and Rheumatoid Arthritis pathogenesis: Analysis of differential gene expression in CD14 ⁺ monocytes and the inflammatory cytokine levels. <i>Immunobiology</i> , 2021, 226, 152152.	1.9	3
42	Differential distribution of vitamin D receptor (<i>VDR</i>) gene variants and its expression in systemic lupus erythematosus. <i>International Journal of Immunogenetics</i> , 2022, 49, 181-192.	1.8	2