Paula Sandrin-Garcia

List of Publications by Year in descending order

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Version: 2024-02-01

42 papers 851 citations

16 h-index 27 g-index

42 all docs 42 docs citations

42 times ranked 1479 citing authors

#	Article	IF	CITATIONS
1	Differential distribution of vitamin D receptor (<i>VDR</i>) gene variants and its expression in systemic lupus erythematosus. International Journal of Immunogenetics, 2022, 49, 181-192.	1.8	2
2	High burden of acute kidney injury in COVID-19 pandemic: systematic review and meta-analysis. Journal of Clinical Pathology, 2021, 74, 796-803.	2.0	26
3	Is there an Inflammation Role for MYD88 in Rheumatoid Arthritis?. Inflammation, 2021, 44, 1014-1022.	3.8	3
4	ADA2 deficiency (DADA2) associated with Evans syndrome and a severe <i>ADA2</i> genotype. Rheumatology, 2021, 60, e237-e239.	1.9	6
5	MYD88, IRAK3 and Rheumatoid Arthritis pathogenesis: Analysis of differential gene expression in CD14Â+Âmonocytes and the inflammatory cytokine levels. Immunobiology, 2021, 226, 152152.	1.9	3
6	The Role of NLRP3 Inflammasome in Lupus Nephritis. International Journal of Molecular Sciences, 2021, 22, 12476.	4.1	27
7	Are key cytokines genetic and serum levels variations related to rheumatoid arthritis clinical severity?. Gene, 2020, 722, 144098.	2.2	15
8	T-cell specific upregulation of Sema4A as risk factor for autoimmunity in systemic lupus erythematosus and rheumatoid arthritis. Autoimmunity, 2020, 53, 65-70.	2.6	4
9	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.	2.0	17
10	<i>>IL1β</i> , <i>IL18</i> , <i>NFKB1</i> and <i>IFNG</i> gene interactions are associated with severity of rheumatoid arthritis: A pilot study. Autoimmunity, 2020, 53, 95-101.	2.6	8
11	Differential expression of the inflammasome complex genes in systemic lupus erythematosus. Immunogenetics, 2020, 72, 217-224.	2.4	31
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.	3.5	22
13	Postmenopausal Osteoporosis reference genes for qPCR expression assays. Scientific Reports, 2019, 9, 16533.	3.3	10
14	<i>Mannose-Binding Lectin2</i> Gene Polymorphism and IgG4 in Membranous Nephropathy. Nephron, 2018, 139, 181-188.	1.8	7
15	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.	4.0	45
16	CTLA-4 gene polymorphisms are associated with obesity in Turner Syndrome. Genetics and Molecular Biology, 2018, 41, 727-734.	1.3	3
17	<i>CCR5Δ32</i> and the genetic susceptibility to rheumatoid arthritis in admixed populations: a multicentre study. Rheumatology, 2017, 56, kew398.	1.9	6
18	<i>PTPN22</i> 1858C > T polymorphism and susceptibility to systemic lupus erythematosus: a meta-analysis update. Autoimmunity, 2017, 50, 428-434.	2.6	12

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19	Ficolin Gene Polymorphisms in Systemic Lupus Erythematosus and Rheumatoid Arthritis. Annals of Human Genetics, 2016, 80, 1-6.	0.8	20
20	Vitamin D receptor polymorphisms and expression profile in rheumatoid arthritis brazilian patients. Molecular Biology Reports, 2016, 43, 41-51.	2.3	14
21	LIG4 and RAD52 DNA repair genes polymorphisms and systemic lupus erythematosus. Molecular Biology Reports, 2014, 41, 2249-2256.	2.3	6
22	FYB gene polymorphisms are associated with susceptibility for systemic lupus erythemathosus (SLE). Human Immunology, 2013, 74, 1009-1014.	2.4	8
23	Polymorphisms in STK17A gene are associated with systemic lupus erythematosus and its clinical manifestations. Gene, 2013, 527, 435-439.	2.2	20
24	Vitamin D receptor (VDR) gene polymorphisms and age onset in type 1 diabetes mellitus. Autoimmunity, 2013, 46, 382-387.	2.6	19
25	Polimorphisms in Inflammasome Genes Are Involved in the Predisposition to Systemic Lupus Erythematosus. Autoimmunity, 2012, 45, 271-278.	2.6	143
26	Mannose binding lectin gene (MBL2) functional polymorphisms are associated with systemic lupus erythematosus in southern Brazilians. Human Immunology, 2011, 72, 516-521.	2.4	22
27	Delayed effects of exposure to a moderate radiation dose on transcription profiles in human primary fibroblasts. Environmental and Molecular Mutagenesis, 2011, 52, 117-129.	2.2	9
28	Alterations in gene expression profiles correlated with cisplatin cytotoxicity in the glioma U343 cell line. Genetics and Molecular Biology, 2010, 33, 159-168.	1.3	17
29	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.	3.8	13
30	Gene Expression Profiles in Radiation Workers Occupationally Exposed to Ionizing Radiation. Journal of Radiation Research, 2009, 50, 61-71.	1.6	73
31	Comprehensive gene expression profiling in lungs of mice infected with <i>Mycobacterium tuberculosis</i> following DNAhsp65 immunotherapy. Journal of Gene Medicine, 2009, 11, 66-78.	2.8	22
32	Differential gene expression of peripheral blood mononuclear cells from rheumatoid arthritis patients may discriminate immunogenetic, pathogenic and treatment features. Immunology, 2009, 127, 365-372.	4.4	20
33	Gene Expression Profiles Stratified according to Type 1 Diabetes Mellitus Susceptibility Regions. Annals of the New York Academy of Sciences, 2008, 1150, 282-289.	3.8	13
34	Gene Expression Profiles in Human Lymphocytes Irradiated In Vitro with Low Doses of Gamma Rays. Radiation Research, 2007, 168, 650.	1.5	59
35	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.	1.3	7
36	Typical phenotypic spectrum of velocardiofacial syndrome occurs independently of deletion size in chromosome 22q11.2. Molecular and Cellular Biochemistry, 2007, 303, 9-17.	3.1	20

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37	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.	3.1	3
38	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.	3.8	6
39	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.	3.8	4
40	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.	5.8	5
41	Promiscuous Gene Expression in the Thymus: The Root of Central Tolerance. Clinical and Developmental Immunology, 2006, 13, 81-99.	3.3	28
42	Gene expression profiles in human cells submitted to genotoxic stress. Mutation Research - Reviews in Mutation Research, 2003, 544, 403-413.	5.5	53