Ana Dopazo

List of Publications by Year in descending order

Source: https://exaly.com/author-pdf/4746222/publications.pdf

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186209 2,899 61 28 citations papers

52 h-index g-index 64 64 64 5600 all docs docs citations times ranked citing authors

175177

#	Article	IF	CITATIONS
1	Trained immunity induction by the inactivated mucosal vaccine MV130 protects against experimental viral respiratory infections. Cell Reports, 2022, 38, 110184.	2.9	34
2	Immune synapse instructs epigenomic and transcriptomic functional reprogramming in dendritic cells. Science Advances, 2021, 7, .	4.7	10
3	Neutrophil subtypes shape HIV-specific CD8 T-cell responses after vaccinia virus infection. Npj Vaccines, 2021, 6, 52.	2.9	6
4	Identification of a peripheral blood gene signature predicting aortic valve calcification. Physiological Genomics, 2020, 52, 563-574.	1.0	11
5	Harmine and Piperlongumine Revert TRIB2-Mediated Drug Resistance. Cancers, 2020, 12, 3689.	1.7	14
6	Discovery of New Targets to Control Metastasis in Pancreatic Cancer by Single-cell Transcriptomics Analysis of Circulating Tumor Cells. Molecular Cancer Therapeutics, 2020, 19, 1751-1760.	1.9	31
7	Oxidized Low-Density Lipoprotein Receptor in Lymphocytes Prevents Atherosclerosis and Predicts Subclinical Disease. Circulation, 2019, 139, 243-255.	1.6	36
8	In vivo phosphoproteomics reveals kinase activity profiles that predict treatment outcome in triple-negative breast cancer. Nature Communications, 2018, 9, 3501.	5.8	45
9	Differential leucocyte RNA expression in the coronary arteries compared to systemic circulation discriminates between patients with and those without coronary artery disease. ClÃnica E Investigación En Arteriosclerosis, 2017, 29, 60-66.	0.4	2
10	5-gene differential expression predicts stability of human intestinal allografts. Experimental and Molecular Pathology, 2017, 103, 163-171.	0.9	3
11	DNA Methylation of miR-7 is a Mechanism Involved in Platinum Response through <i>MAFG</i> Overexpression in Cancer Cells. Theranostics, 2017, 7, 4118-4134.	4.6	52
12	AG-NGS: A powerful and user-friendly computing application for the semi-automated preparation of next-generation sequencing libraries using open liquid handling platforms. BioTechniques, 2017, 62, xvi.	0.8	1
13	Methotrexate selectively targets human proinflammatory macrophages through a thymidylate synthase/p53 axis. Annals of the Rheumatic Diseases, 2016, 75, 2157-2165.	0.5	35
14	Technologies to Study Genetics and Molecular Pathways. , 2016, , 251-269.		0
15	Genomeâ€Wide Transcriptional and Functional Analysis of Endoglin Isoforms in the Human Promonocytic Cell Line U937. Journal of Cellular Physiology, 2015, 230, 947-958.	2.0	13
16	AG-NGS: A powerful and user-friendly computing application for the semi-automated preparation of next-generation sequencing libraries using open liquid handling platforms. BioTechniques, 2014, 56, 28-35.	0.8	9
17	A novel phosphatidylinositol 3-kinase (PI3K) inhibitor directs a potent FOXO-dependent, p53-independent cell cycle arrest phenotype characterized by the differential induction of a subset of FOXO-regulated genes. Breast Cancer Research, 2014, 16, 482.	2.2	41
18	Disease-modifying factors in hereditary angioedema: an RNA expression-based screening. Orphanet Journal of Rare Diseases, 2013, 8, 77.	1.2	12

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19	Human mesenchymal stem cell-replicative senescence and oxidative stress are closely linked to aneuploidy. Cell Death and Disease, 2013, 4, e691-e691.	2.7	192
20	Embryological-Origin–Dependent Differences in Homeobox Expression in Adult Aorta. Arteriosclerosis, Thrombosis, and Vascular Biology, 2013, 33, 1248-1256.	1.1	53
21	CSF3R T618I co-occurs with mutations of splicing and epigenetic genes and with a new PIM3 truncated fusion gene in chronic neutrophilic leukemia. Blood Cancer Journal, 2013, 3, e158-e158.	2.8	15
22	Aryl hydrocarbon receptor contributes to the MEK/ERK-dependent maintenance of the immature state of human dendritic cells. Blood, 2013, 121, e108-e117.	0.6	37
23	Chemokines induce axon outgrowth downstream of Hepatocyte Growth Factor and TCF/\hat{l}^2 -catenin signaling. Frontiers in Cellular Neuroscience, 2013, 7, 52.	1.8	23
24	Understanding cardiovascular disease: a journey through the genome (and what we found there). DMM Disease Models and Mechanisms, 2012, 5, 434-443.	1,2	40
25	Identification of Disease-Relevant Genes for Molecularly-Targeted Drug Discovery. Current Cancer Drug Targets, 2012, 12, 1-13.	0.8	13
26	Culture of human mesenchymal stem cells at low oxygen tension improves growth and genetic stability by activating glycolysis. Cell Death and Differentiation, 2012, 19, 743-755.	5.0	230
27	Cooperativity of Stress-Responsive Transcription Factors in Core Hypoxia-Inducible Factor Binding Regions. PLoS ONE, 2012, 7, e45708.	1.1	46
28	Spatial and Temporal Gene Expression Differences in Core and Periinfarct Areas in Experimental Stroke: A Microarray Analysis. PLoS ONE, 2012, 7, e52121.	1,1	59
29	Insights into Global Mechanisms and Disease by Gene Expression Profiling. Methods in Molecular Biology, 2011, 719, 269-298.	0.4	2
30	Automatic Genomics: a user-friendly program for the automatic designing and plate loading of medium-throughput qPCR experiments. BioTechniques, 2011, 50, 46-50.	0.8	2
31	Formin1 Mediates the Induction of Dendritogenesis and Synaptogenesis by Neurogenin3 in Mouse Hippocampal Neurons. PLoS ONE, 2011, 6, e21825.	1.1	26
32	miR-335 orchestrates cell proliferation, migration and differentiation in human mesenchymal stem cells. Cell Death and Differentiation, 2011, 18, 985-995.	5.0	265
33	Characterization of host genetic expression patterns in HIV-infected individuals with divergent disease progression. Virology, 2011, 411, 103-112.	1.1	22
34	Early peroxisome proliferator-activated receptor gamma regulated genes involved in expansion of pancreatic beta cell mass. BMC Medical Genomics, 2011, 4, 86.	0.7	15
35	Progress and challenges in developing a molecular diagnostic test for neurofibromatosis type 1. Expert Review of Molecular Diagnostics, 2011, 11, 671-673.	1.5	3
36	PGC- $1\hat{l}\pm$ Regulates Translocated in Liposarcoma Activity: Role in Oxidative Stress Gene Expression. Antioxidants and Redox Signaling, 2011, 15, 325-337.	2.5	24

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37	Processing of Agilent microRNA array data. BMC Research Notes, 2010, 3, 18.	0.6	77
38	Human TRIB2 is a repressor of FOXO that contributes to the malignant phenotype of melanoma cells. Oncogene, 2010, 29, 2973-2982.	2.6	85
39	Modifications in host cell cytoskeleton structure and function mediated by intracellular HIV-1 Tat protein are greatly dependent on the second coding exon. Nucleic Acids Research, 2010, 38, 3287-3307.	6.5	55
40	Large-scale gene expression in bone marrow mesenchymal stem cells: a putative role for COL10A1 in osteoarthritis. Annals of the Rheumatic Diseases, 2010, 69, 1880-1885.	0.5	38
41	The novel RUNX3/p33 isoform is induced upon monocyte-derived dendritic cell maturation and downregulates IL-8 expression. Immunobiology, 2010, 215, 812-820.	0.8	19
42	Cux1 and Cux2 Regulate Dendritic Branching, Spine Morphology, and Synapses of the Upper Layer Neurons of the Cortex. Neuron, 2010, 66, 523-535.	3.8	247
43	Modifications in host cell structure and functions mediated by Tat intracellular expression are greatly dependent on the second exon. Retrovirology, 2009, 6, .	0.9	0
44	Estradiol Activates Î ² -Catenin Dependent Transcription in Neurons. PLoS ONE, 2009, 4, e5153.	1.1	71
45	Transcriptional expression ofcis-acting andtrans-acting splicing mutations cause autosomal dominant retinitis pigmentosa. Human Mutation, 2008, 29, 869-878.	1.1	31
46	Codelink: an R package for analysis of GE healthcare gene expression bioarrays. Bioinformatics, 2007, 23, 1168-1169.	1.8	24
47	Position-dependent expression of GADD45α in rat brain tumours. Medical Oncology, 2007, 24, 436-444.	1.2	3
48	The Repression of E2F-1 Is Critical for the Activity of Minerval against Cancer. Journal of Pharmacology and Experimental Therapeutics, 2005, 315, 466-474.	1.3	38
49	Gene expression analysis of chromosomal regions with gain or loss of genetic material detected by comparative genomic hybridization. Genes Chromosomes and Cancer, 2004, 41, 353-365.	1.5	17
50	Growth and molecular profile of lung cancer cells expressing ectopic LKB1: down-regulation of the phosphatidylinositol 3'-phosphate kinase/PTEN pathway. Cancer Research, 2003, 63, 1382-8.	0.4	57
51	The molecular signature of mantle cell lymphoma reveals multiple signals favoring cell survival. Cancer Research, 2003, 63, 8226-32.	0.4	130
52	Identification of Genes Involved in Resistance to Interferon- \hat{l}_{\pm} in Cutaneous T-Cell Lymphoma. American Journal of Pathology, 2002, 161, 1825-1837.	1.9	106
53	Overview of the most prevalent hypothalamus-specific mRNAs, as identified by directional tag PCR subtraction Proceedings of the National Academy of Sciences of the United States of America, 1996, 93, 8733-8738.	3.3	255
54	Characterization of mutant alleles of the cell division protein FtsA, a regulator and structural component of the Escherichia coli septator. Biochimie, 1994, 76, 1071-1074.	1.3	4

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55	Primary structure of mouse secretogranin III and its absence from deficient mice. Journal of Molecular Neuroscience, 1993, 4, 225-233.	1.1	17
56	An amino-proximal domain required for the localization of FtsQ in the cytoplasmic membrane, and for its biological function in Escherichia coli. Molecular Microbiology, 1992, 6, 715-722.	1.2	24
57	On the chronology and topography of bacterial cell division. Research in Microbiology, 1991, 142, 253-257.	1.0	12
58	The native form of FtsA, a septal protein of Escherichia coli, is located in the cytoplasmic membrane. Journal of Bacteriology, 1990, 172, 5097-5102.	1.0	58
59	Structural inhibition and reactivation of Escherichia coli septation by elements of the SOS and TER pathways. Journal of Bacteriology, 1987, 169, 1772-1776.	1.0	13
60	Coupling between DNA replication and cell division mediated by the FtsA protein in Escherichia coli: a pathway independent of the SOS response, the "TER" pathway. Journal of Bacteriology, 1985, 164, 950-953.	1.0	27
61	Guest Commentary on Chapter 4: Integrative Approaches to Genotype-Phenotype Association Discovery. , 0, , 73-76.		0