Stephano S Mello

List of Publications by Year
in descending order

46 papers	3,577 citations	361413 20 h-index	289244 40 g-index
46 all docs	46 docs citations	46 times ranked	7468 citing authors

#	Article	IF	Citations
1	The HIF target MAFF promotes tumor invasion and metastasis through IL11 and STAT3 signaling. Nature Communications, 2021, 12, 4308.	12.8	45
2	Zmat3 Is a Key Splicing Regulator in the p53 Tumor Suppression Program. Molecular Cell, 2020, 80, 452-469.e9.	9.7	44
3	The Long Noncoding RNA<i>NEAT1</i>Promotes Sarcoma Metastasis by Regulating RNA Splicing Pathways. Molecular Cancer Research, 2020, 18, 1534-1544.	3.4	16
4	p53 deficiency triggers dysregulation of diverse cellular processes in physiological oxygen. Journal of Cell Biology, 2020, 219, .	5.2	26
5	Counting the Minutes. ELife, 2020, 9, .	6.0	3
6	Deciphering p53 signaling in tumor suppression. Current Opinion in Cell Biology, 2018, 51, 65-72.	5.4	170
7	Neat-en-ing up our understanding of p53 pathways in tumor suppression. Cell Cycle, 2018, 17, 1527-1535.	2.6	9
8	Dynamin impacts homology-directed repair and breast cancer response to chemotherapy. Journal of Clinical Investigation, 2018, 128, 5307-5321.	8.2	20
9	Abstract IA07: Deconstructing p53 pathways in tumor suppression. , 2018, , .		0
10	A p53 Super-tumor Suppressor Reveals a Tumor Suppressive p53-Ptpn14-Yap Axis in Pancreatic Cancer. Cancer Cell, 2017, 32, 460-473.e6.	16.8	142
11	<i>Neat1</i> is a p53-inducible lincRNA essential for transformation suppression. Genes and Development, 2017, 31, 1095-1108.	5.9	179
12	Abstract 1628: MAFF, a new hypoxia target gene involving tumor invasion and metastasis. , 2016, , .		0
13	Abstract PR07: Deconstructing p53 transcriptional networks in pancreatic cancer suppression. , 2016, , .		0
14	Abstract A45: Neat1 is a p53-inducible lincRNA important for pancreatic cancer suppression. , 2016, , .		0
15	The p53 Target Gene <i>SIVA</i> Enables Nonâ€“Small Cell Lung Cancer Development. Cancer Discovery, 2015, 5, 622-635.	9.4	24
16	Combined inhibition of BET family proteins and histone deacetylases as a potential epigenetics-based therapy for pancreatic ductal adenocarcinoma. Nature Medicine, 2015, 21, 1163-1171.	30.7	349
17	Unravelling mechanisms of p53-mediated tumour suppression. Nature Reviews Cancer, 2014, 14, 359-370.	28.4	1,090
18	Ionizing radiation-induced gene expression changes in TP53 proficient and deficient glioblastoma cell lines. Mutation Research - Genetic Toxicology and Environmental Mutagenesis, 2013, 756, 46-55.	1.7	24

#	ARTICLE	IF	CITATIONS
19	Not all p53 gain-of-function mutants are created equal. Cell Death and Differentiation, 2013, 20, 855-857.	11.2	52
20	Global genomic profiling reveals an extensive p53-regulated autophagy program contributing to key p53 responses. Genes and Development, 2013, 27, 1016-1031.	5.9	353
21	Abstract IA4: Deconstructing p53 pathways in vivo. , 2013, , .		0
22	Abstract B12: Deciphering mechanisms of p53-mediated pancreatic cancer suppression.. , 2012, , .		0
23	Distinct p53 Transcriptional Programs Dictate Acute DNA-Damage Responses and Tumor Suppression. Cell, 2011, 145, 571-583.	28.9	443
24	The non-coding RNA BC1 is down-regulated in the hippocampus of Wistar Audiogenic Rat (WAR) strain after audiogenic kindling. Brain Research, 2011, 1367, 114-121.	2.2	22
25	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
26	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
27	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
28	Gene Expression Profiles in Radiation Workers Occupationally Exposed to Ionizing Radiation. Journal of Radiation Research, 2009, 50, 61-71.	1.6	73
29	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
30	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
31	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
32	Transcriptional changes in U343 MG-a glioblastoma cell line exposed to ionizing radiation. Human and Experimental Toxicology, 2008, 27, 919-929.	2.2	19
33	Gene Expression Profiles in Human Lymphocytes Irradiated In Vitro with Low Doses of Gamma Rays. Radiation Research, 2007, 168, 650.	1.5	59
34	Profiling Meta-Analysis Reveals Primarily Gene Coexpression Concordance between Systemic Lupus Erythematosus and Rheumatoid Arthritis. Annals of the New York Academy of Sciences, 2007, 1110, 33-46.	3.8	25
35	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
36	Hybridization signatures of gamma-irradiated murine fetal thymus organ culture (FTOC) reveal modulation of genes associated with T-cell receptor V(D)J recombination and DNA repair. Molecular Immunology, 2006, 43, 464-472.	2.2	7

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37	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
38	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
39	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
40	Changes in the gene expression profiling of the thymus in response to fibrosarcoma growth. Molecular and Cellular Biochemistry, 2005, 276, 81-88.	3.1	1
41	Genomic Instability:Signaling Pathways Orchestrating the Responsesto Ionizing Radiation and Cisplatin. Genome Dynamics and Stability, 2005, , 423-452.	1.1	1
42	Transcriptional Profiles of the Human Pathogenic Fungus Paracoccidioides brasiliensis in Mycelium and Yeast Cells. Journal of Biological Chemistry, 2005, 280, 24706-24714.	3.4	169
43	Hybridization signatures during thymus ontogeny reveals modulation of genes coding for T-cell signaling proteins. Molecular Immunology, 2005, 42, 1043-1048.	2.2	9
44	Immunosuppressive therapy modulates T lymphocyte gene expression in patients with systemic lupus erythematosus. Immunology, 2004, 113, 99-105.	4.4	27
45	Gene expression profiles in human cells submitted to genotoxic stress. Mutation Research - Reviews in Mutation Research, 2003, 544, 403-413.	5.5	53
46	Chromosomal rearrangements involving telomeric DNA sequences in Balb/3T3 cells transfected with the Ha-ras oncogene. Mutagenesis, 2002, 17, 67-72.	2.6	11