Bruna De Felice

List of Publications by Year in descending order

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430754 454834 38 918 18 30 citations h-index g-index papers 40 40 40 1801 docs citations times ranked citing authors all docs

#	Article	IF	CITATIONS
1	A miRNA signature in leukocytes from sporadic amyotrophic lateral sclerosis. Gene, 2012, 508, 35-40.	1.0	126
2	miR-338-3p is over-expressed in blood, CFS, serum and spinal cord from sporadic amyotrophic lateral sclerosis patients. Neurogenetics, 2014, 15, 243-253.	0.7	99
3	Genome–wide microRNA expression profiling in placentas from pregnant women exposed to BPA. BMC Medical Genomics, 2015, 8, 56.	0.7	65
4	Differential apoptosis markers in human keloids and hypertrophic scars fibroblasts. Molecular and Cellular Biochemistry, 2009, 327, 191-201.	1.4	60
5	Quercetin Increases MUC2 and MUC5AC Gene Expression and Secretion in Intestinal Goblet Cell-Like LS174T via PLC/PKCα/ERK1-2 Pathway. Frontiers in Physiology, 2018, 9, 357.	1.3	46
6	Differential p63 and p53 Expression in Human Keloid Fibroblasts and Hypertrophic Scar Fibroblasts. DNA and Cell Biology, 2007, 26, 541-547.	0.9	40
7	Telomerase expression in amyotrophic lateral sclerosis (ALS) patients. Journal of Human Genetics, 2014, 59, 555-561.	1.1	36
8	Small non-coding RNA signature in multiple sclerosis patients after treatment with interferon- \hat{l}^2 . BMC Medical Genomics, 2014, 7, 26.	0.7	36
9	Wide-Ranging Analysis of MicroRNA Profiles in Sporadic Amyotrophic Lateral Sclerosis Using Next-Generation Sequencing. Frontiers in Genetics, 2018, 9, 310.	1.1	34
10	T cell activation induces CuZn superoxide dismutase (SOD)-1 intracellular re-localization, production and secretion. Biochimica Et Biophysica Acta - Molecular Cell Research, 2014, 1843, 265-274.	1.9	33
11	Human miR-26a-5p regulates the glutamate transporter SLC1A1 (EAAT3) expression. Relevance in multiple sclerosis. Biochimica Et Biophysica Acta - Molecular Basis of Disease, 2018, 1864, 317-323.	1.8	33
12	Telomere shortening in women resident close to waste landfill sites. Gene, 2012, 500, 101-106.	1.0	29
13	Telomere shortening may be associated with human keloids. BMC Medical Genetics, 2009, 10, 110.	2.1	28
14	A transcriptionally active copia-like retroelement in Citrus limon. Cellular and Molecular Biology Letters, 2009, 14, 289-304.	2.7	25
15	Gene expression profiling in zebrafish embryos exposed to diclofenac, an environmental toxicant. Molecular Biology Reports, 2012, 39, 2119-2128.	1.0	24
16	Association between exposure to dioxin-like polychlorinated biphenyls and miR-191 expression in human peripheral blood mononuclear cells. Mutation Research - Genetic Toxicology and Environmental Mutagenesis, 2013, 753, 36-41.	0.9	23
17	MicroRNA Expression Signature in Mild Cognitive Impairment Due to Alzheimer's Disease. Molecular Neurobiology, 2020, 57, 4408-4416.	1.9	23
18	Cigarette Smoke Condensate Causes a Decrease of the Gene Expression of Cu–Zn Superoxide Dismutase, Mn Superoxide Dismutase, Glutathione Peroxidase, Catalase, and Free Radical-Induced Cell Injury in SH-SY5Y Human Neuroblastoma Cells. Neurotoxicity Research, 2011, 19, 49-54.	1.3	20

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19	Genetic fingerprint of microorganisms associated with the deterioration of an historical tuff monument in Italy. Journal of Genetics, 2010, 89, 253-257.	0.4	17
20	The IFN- \hat{I}^2 1b effect on Cu Zn superoxide dismutase (SOD1) in peripheral mononuclear blood cells of relapsing-remitting multiple sclerosis patients and in neuroblastoma SK-N-BE cells. Brain Research Bulletin, 2015, 118, 1-6.	1.4	13
21	miR-34 modulates apoptotic gene expression in Ingenol mebutate treated keloid fibroblasts. Molecular Medicine Reports, 2018, 17, 7081-7088.	1.1	12
22	Modulation of 3-Hydroxy-3-methylglutaryl-CoA Reductase Gene Expression by CuZn Superoxide Dismutase in Human Fibroblasts and HepG2 Cells. Gene Expression, 2004, 12, 29-38.	0.5	10
23	Novel methylation at GpC dinucleotide in the fish Sparus aurata genome. Molecular Biology Reports, 2000, 27, 225-230.	1.0	9
24	Molecular Characterization of Microbial Population Dynamics during Sildenafil Citrate Degradation. Molecular Biotechnology, 2009, 41, 123-132.	1.3	9
25	A novel repeated sequence DNA originated from a Tc1-like transposon in water green frog Rana esculenta. Gene, 2000, 261, 205-210.	1.0	8
26	Effect of selenocystine on gene expression profiles in human keloid fibroblasts. Genomics, 2011, 97, 265-276.	1.3	8
27	Comparative transcriptional analysis reveals differential gene expression between Sand Daffodil tissues. Genetica, 2013, 141, 443-452.	0.5	8
28	Assessment of DNA Damage by RAPD inParacentrotus lividusEmbryos Exposed to Amniotic Fluid from Residents Living Close to Waste Landfill Sites. Journal of Biomedicine and Biotechnology, 2010, 2010, 1-7.	3.0	7
29	Ingenol mebutate treatment in keloids. BMC Research Notes, 2015, 8, 466.	0.6	7
30	Variability of DNA methylation pattern in somatic and germ cells in male newt (Amphibia,) Tj ETQq0 0 0 rgBT /Ov	verlogk 10	Tf 50 302 Td
31	Molecular characterization of a novel pathogen-responsive receptor kinase-like in Citrus limon. Tree Genetics and Genomes, 2010, 6, 47-56.	0.6	5
32	Microbial diversity of landslide soils assessed by RFLP and SSCP fingerprints. Journal of Applied Genetics, 2014, 55, 403-415.	1.0	5
33	Characterization of DIP1, a novel nuclear protein in Drosophila melanogaster. Biochemical and Biophysical Research Communications, 2003, 307, 224-228.	1.0	4
34	Characterization of a novel satellite DNA sequence from Flying Dragon (Poncirus trifoliata). Genetica, 2006, 127, 45-53.	0.5	4
35	Molecular analysis of a novel tandemly organized repetitive DNA sequence inCitrus limon (L.) Burm. Journal of Applied Genetics, 2007, 48, 233-239.	1.0	2
36	Assessment of genetic diversity between wild and cultivated artichokes using SSR markers. Genetic Resources and Crop Evolution, 2016, 63, 1363-1369.	0.8	2

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	37	Genetic structure of a novel biofuel-producing microorganism community. Journal of Genetics, 2012, 91, 183-191.	0.4	1
	38	Identification of a cDNA clone encoding DIP1-binding protein in Drosophila melanogaster. Molecular Biology Reports, 2004, 31, 165-169.	1.0	0