Claudia Coronnello

List of Publications by Year in descending order

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623734 526287 33 787 14 27 citations g-index h-index papers 34 34 34 1243 docs citations times ranked citing authors all docs

#	Article	IF	CITATIONS
1	Moving Towards Induced Pluripotent Stem Cell-based Therapies with Artificial Intelligence and Machine Learning. Stem Cell Reviews and Reports, 2022, 18, 559-569.	3.8	22
2	A multivariate statistical test for differential expression analysis. Scientific Reports, 2022, 12, 8265.	3.3	3
3	Repetitive sequences in aging. Aging, 2021, 13, 10816-10817.	3.1	1
4	Support Vector Machine as a Supervised Learning for the Prioritization of Novel Potential SARS-CoV-2 Main Protease Inhibitors. International Journal of Molecular Sciences, 2021, 22, 7714.	4.1	11
5	Transcriptomic Changes Following Partial Depletion of CENP-E in Normal Human Fibroblasts. Genes, 2021, 12, 1322.	2.4	1
6	miRNA expression analysis in the human heart: Undifferentiated progenitors vs. bioptic tissuesâ€"Implications for proliferation and ageing. Journal of Cellular and Molecular Medicine, 2021, 25, 8687-8700.	3.6	5
7	A Radioactive-Free Method for the Thorough Analysis of the Kinetics of Cell Cytotoxicity. Journal of Imaging, 2021, 7, 222.	3.0	2
8	Usefulness of regional right ventricular and right atrial strain for prediction of early and late right ventricular failure following a left ventricular assist device implant: A machine learning approach. International Journal of Artificial Organs, 2020, 43, 297-314.	1.4	16
9	Cross-talk between Colon Cells and Macrophages Increases ST6GALNAC1 and MUC1-sTn Expression in Ulcerative Colitis and Colitis-Associated Colon Cancer. Cancer Immunology Research, 2020, 8, 167-178.	3.4	61
10	Lung Segmentation on High-Resolution Computerized Tomography Images Using Deep Learning: A Preliminary Step for Radiomics Studies. Journal of Imaging, 2020, 6, 125.	3.0	31
11	miR-1207-5p Can Contribute to Dysregulation of Inflammatory Response in COVID-19 via Targeting SARS-CoV-2 RNA. Frontiers in Cellular and Infection Microbiology, 2020, 10, 586592.	3.9	32
12	Residue analysis of a synthetic glucocorticoid in liver samples by a 1HMR spectroscopy approach: An exploratory study on animal model. Food Additives and Contaminants - Part A Chemistry, Analysis, Control, Exposure and Risk Assessment, 2020, 37, 1640-1650.	2.3	1
13	An improvement of ComiR algorithm for microRNA target prediction by exploiting coding region sequences of mRNAs. BMC Bioinformatics, 2020, 21, 201.	2.6	12
14	Aneuploid IMR90 cells induced by depletion of pRB, DNMT1 and MAD2 show a common gene expression signature. Genomics, 2020, 112, 2541-2549.	2.9	4
15	Tissue Classification to Support Local Active Delineation of Brain Tumors. Communications in Computer and Information Science, 2020, , 3-14.	0.5	23
16	Radiomics: A New Biomedical Workflow to Create a Predictive Model. Communications in Computer and Information Science, 2020, , 280-293.	0.5	16
17	NK cell anti-tumor and anti-viral function can be enhanced in vitro and exploited for successful cell mediated therapy. Cytotherapy, 2019, 21, S29.	0.7	O
18	RIP-Chip analysis supports different roles for AGO2 and GW182 proteins in recruiting and processing microRNA targets. BMC Bioinformatics, 2019, 20, 120.	2.6	17

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19	MTGO-SC, A Tool to Explore Gene Modules in Single-Cell RNA Sequencing Data. Frontiers in Genetics, 2019, 10, 953.	2.3	3
20	Predicting Acute and Chronic Right Ventricular Failure in Patients Undergoing Left Ventricular Assist Device Implant: The Importance of Right Atrial Strain and Regional Deformation of the Right Ventricular Free Wall. Journal of Heart and Lung Transplantation, 2018, 37, S381-S382.	0.6	0
21	On the prospect of serum exosomal miRNA profiling and protein biomarkers for the diagnosis of ascending aortic dilatation in patients with bicuspid and tricuspid aortic valve. International Journal of Cardiology, 2018, 273, 230-236.	1.7	36
22	Gene-based and semantic structure of the Gene Ontology as a complex network. Physica A: Statistical Mechanics and Its Applications, 2016, 458, 313-328.	2.6	2
23	NK Cells Functional Skew: Implications in Innate Immunity to HCV Infection. Journal of Hepatology, 2016, 64, S511-S512.	3.7	0
24	The Challenge of Using CB-HSCs As Source for Gene Therapy: Lentiviral Vector Transduction, Phenotypic Characterization and Global Gene Expression Profile of Ex-Vivo Expanded CB CD34+ Cells. Blood, 2015, 126, 5548-5548.	1.4	0
25	Aldosterone Regulates MicroRNAs in the Cortical Collecting Duct to Alter Sodium Transport. Journal of the American Society of Nephrology: JASN, 2014, 25, 2445-2457.	6.1	42
26	Overexpression of microRNA-1 promotes cardiomyocyte commitment from human cardiovascular progenitors via suppressing WNT and FGF signaling pathways. Journal of Molecular and Cellular Cardiology, 2013, 63, 146-154.	1.9	62
27	Estrogen represses gene expression through reconfiguring chromatin structures. Nucleic Acids Research, 2013, 41, 8061-8071.	14.5	17
28	ComiR: combinatorial microRNA target prediction tool. Nucleic Acids Research, 2013, 41, W159-W164.	14.5	174
29	Expression of Regulatory Platelet MicroRNAs in Patients with Sickle Cell Disease. PLoS ONE, 2013, 8, e60932.	2.5	21
30	Novel Modeling of Combinatorial miRNA Targeting Identifies SNP with Potential Role in Bone Density. PLoS Computational Biology, 2012, 8, e1002830.	3.2	38
31	SPANNING TREES AND BOOTSTRAP RELIABILITY ESTIMATION IN CORRELATION-BASED NETWORKS. International Journal of Bifurcation and Chaos in Applied Sciences and Engineering, 2007, 17, 2319-2329.	1.7	124
32	Economic sector identification in a set of stocks traded at the New York Stock Exchange: a comparative analysis., 2007,,.		8
33	Time evolution of the microwave second-order response of YBaCuO powder. Physica C: Superconductivity and Its Applications, 2004, 402, 309-316.	1.2	2