

Marianna Aprile

List of Publications by Year in descending order

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20
papers

674
citations

759233

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888059

17
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times ranked

1207
citing authors

#	ARTICLE	IF	CITATIONS
1	TNF α Mediates Inflammation-Induced Effects on PPAR γ Splicing in Adipose Tissue and Mesenchymal Precursor Cells. <i>Cells</i> , 2022, 11, 42.	4.1	6
2	Hepatic Insulin Resistance in Hyperthyroid Rat Liver: Vitamin E Supplementation Highlights a Possible Role of ROS. <i>Antioxidants</i> , 2022, 11, 1295.	5.1	13
3	Subcellular Localization of uc.8+ as a Prognostic Biomarker in Bladder Cancer Tissue. <i>Cancers</i> , 2021, 13, 681.	3.7	12
4	PPAR δ and Diabetes: Beyond the Genome and Towards Personalized Medicine. <i>Current Diabetes Reports</i> , 2021, 21, 18.	4.2	23
5	In severe obesity, subcutaneous adipose tissue cell-derived cytokines are early markers of impaired glucose tolerance and are modulated by quercetin. <i>International Journal of Obesity</i> , 2021, 45, 1811-1820.	3.4	9
6	LncRNAs in Cancer: From garbage to Junk. <i>Cancers</i> , 2020, 12, 3220.	3.7	41
7	In Vitro-Generated Hypertrophic-Like Adipocytes Displaying PPAR γ Isoforms Unbalance Recapitulate Adipocyte Dysfunctions In Vivo. <i>Cells</i> , 2020, 9, 1284.	4.1	14
8	Tumorigenesis-Related Long Noncoding RNAs and Their Targeting as Therapeutic Approach in Cancer. <i>RNA Technologies</i> , 2020, , 277-303.	0.3	0
9	PPAR $\delta^{\Delta 5}$, a Naturally Occurring Dominant-Negative Splice Isoform, Impairs PPAR δ Function and Adipocyte Differentiation. <i>Cell Reports</i> , 2018, 25, 1577-1592.e6.	6.4	58
10	Unique true predicted neoantigens (TPNAs) correlates with anti-tumor immune control in HCC patients. <i>Journal of Translational Medicine</i> , 2018, 16, 286.	4.4	24
11	Heart failure: Pilot transcriptomic analysis of cardiac tissue by RNA-sequencing. <i>Cardiology Journal</i> , 2017, 24, 539-553.	1.2	54
12	Identification and Validation of HCC-specific Gene Transcriptional Signature for Tumor Antigen Discovery. <i>Scientific Reports</i> , 2016, 6, 29258.	3.3	22
13	Abstract A046: Identification and validation of HCC-specific gene transcriptional signature for tumor antigen discovery. , 2016, , .		0
14	Stimulation of Innate and Adaptive Immunity by Using Filamentous Bacteriophage fd Targeted to DEC-205. <i>Journal of Immunology Research</i> , 2015, 2015, 1-11.	2.2	15
15	AnaLysis of Expression on human chromosome 21, ALE-HSA21: a pilot integrated web resource. Database: the Journal of Biological Databases and Curation, 2014, 2014, bau009.	3.0	12
16	RNA-Seq and human complex diseases: recent accomplishments and future perspectives. <i>European Journal of Human Genetics</i> , 2013, 21, 134-142.	2.8	216
17	Non-coding RNA and pseudogenes in neurodegenerative diseases: "The (un)Usual Suspects". <i>Frontiers in Genetics</i> , 2012, 3, 231.	2.3	40
18	Massive-scale RNA-Seq experiments in human genetic diseases. <i>EMBnet Journal</i> , 2012, 17, 39.	0.6	0

#	ARTICLE	IF	CITATIONS
19	Massive-Scale RNA-Seq Analysis of Non Ribosomal Transcriptome in Human Trisomy 21. PLoS ONE, 2011, 6, e18493.	2.5	62
20	PPARG: Gene Expression Regulation and Next-Generation Sequencing for Unsolved Issues. PPAR Research, 2010, 2010, 1-17.	2.4	52