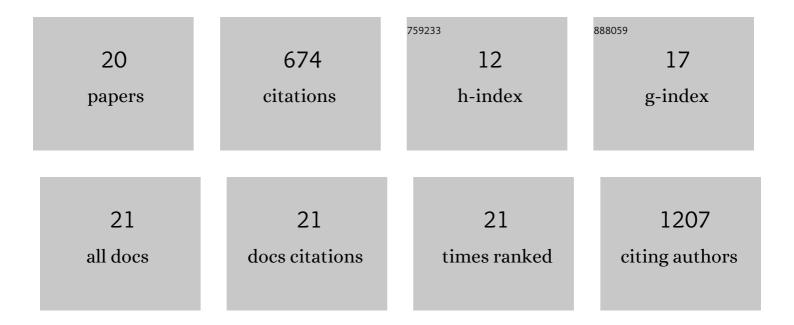
Marianna Aprile

List of Publications by Year in descending order

Source: https://exaly.com/author-pdf/983659/publications.pdf Version: 2024-02-01



#	Article	IF	CITATIONS
1	TNFα Mediates Inflammation-Induced Effects on PPARG Splicing in Adipose Tissue and Mesenchymal Precursor Cells. Cells, 2022, 11, 42.	4.1	6
2	Hepatic Insulin Resistance in Hyperthyroid Rat Liver: Vitamin E Supplementation Highlights a Possible Role of ROS. Antioxidants, 2022, 11, 1295.	5.1	13
3	Subcellular Localization of uc.8+ as a Prognostic Biomarker in Bladder Cancer Tissue. Cancers, 2021, 13, 681.	3.7	12
4	PPARÎ ³ and Diabetes: Beyond the Genome and Towards Personalized Medicine. Current Diabetes Reports, 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. International Journal of Obesity, 2021, 45, 1811-1820.	3.4	9
6	LncRNAs in Cancer: From garbage to Junk. Cancers, 2020, 12, 3220.	3.7	41
7	In Vitro-Generated Hypertrophic-Like Adipocytes Displaying PPARG Isoforms Unbalance Recapitulate Adipocyte Dysfunctions In Vivo. Cells, 2020, 9, 1284.	4.1	14
8	Tumorigenesis-Related Long Noncoding RNAs and Their Targeting as Therapeutic Approach in Cancer. RNA Technologies, 2020, , 277-303.	0.3	0
9	PPARÎ ³ Δ5, a Naturally Occurring Dominant-Negative Splice Isoform, Impairs PPARÎ ³ Function and Adipocyte Differentiation. Cell Reports, 2018, 25, 1577-1592.e6.	6.4	58
10	Unique true predicted neoantigens (TPNAs) correlates with anti-tumor immune control in HCC patients. Journal of Translational Medicine, 2018, 16, 286.	4.4	24
11	Heart failure: Pilot transcriptomic analysis of cardiac tissue by RNA-sequencing. Cardiology Journal, 2017, 24, 539-553.	1.2	54
12	Identification and Validation of HCC-specific Gene Transcriptional Signature for Tumor Antigen Discovery. Scientific Reports, 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. Journal of Immunology Research, 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. European Journal of Human Genetics, 2013, 21, 134-142.	2.8	216
17	Non-coding RNA and pseudogenes in neurodegenerative diseases: "The (un)Usual Suspects― Frontiers in Genetics, 2012, 3, 231.	2.3	40
18	Massive-scale RNA-Seq experiments in human genetic diseases. EMBnet Journal, 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