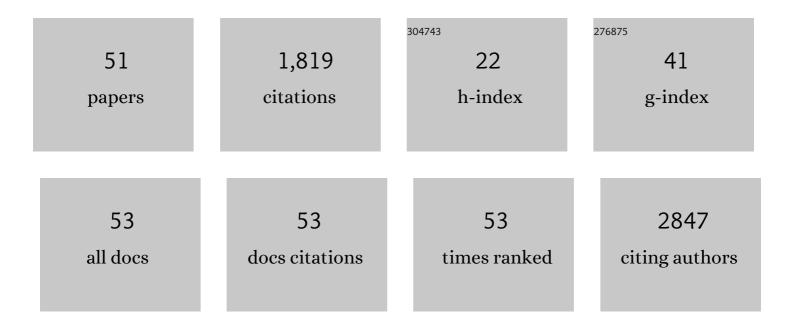
Apollonia Tullo

List of Publications by Year in descending order

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#	Article	IF	CITATIONS
1	Tissue-specific mtDNA abundance from exome data and its correlation with mitochondrial transcription, mass and respiratory activity. Mitochondrion, 2015, 20, 13-21.	3.4	146
2	Human Mitochondrial tRNA Processing. Journal of Biological Chemistry, 1995, 270, 12885-12891.	3.4	135
3	Identification and functional characterization of two new transcriptional variants of the human p63 gene. Nucleic Acids Research, 2009, 37, 6092-6104.	14.5	130
4	p53 mutation is a poor prognostic indicator for survival in patients with hepatocellular carcinoma undergoing surgical tumour ablation. British Journal of Cancer, 1998, 77, 776-782.	6.4	103
5	Whole transcriptome profiling of Late-Onset Alzheimer's Disease patients provides insights into the molecular changes involved in the disease. Scientific Reports, 2018, 8, 4282.	3.3	102
6	TRIM8 modulates p53 activity to dictate cell cycle arrest. Cell Cycle, 2012, 11, 511-523.	2.6	78
7	Impairment of F1F0-ATPase, adenine nucleotide translocator and adenylate kinase causes mitochondrial energy deficit in human skin fibroblasts with chromosome 21 trisomy. Biochemical Journal, 2010, 431, 299-310.	3.7	76
8	Respiratory complex I is essential to induce a Warburg profile in mitochondria-defective tumor cells. Cancer & Metabolism, 2013, 1, 11.	5.0	75
9	TRIM8 restores p53 tumour suppressor function by blunting N-MYC activity in chemo-resistant tumours. Molecular Cancer, 2017, 16, 67.	19.2	73
10	Complexity and Dynamics of the Winemaking Bacterial Communities in Berries, Musts, and Wines from Apulian Grape Cultivars through Time and Space. PLoS ONE, 2016, 11, e0157383.	2.5	60
11	p53FamTaG: a database resource of human p53, p63 and p73 direct target genes combining in silico prediction and microarray data. BMC Bioinformatics, 2007, 8, S20.	2.6	57
12	The complete and symmetric transcription of the main non coding region of rat mitochondrial genome: in vivo mapping of heavy and light transcripts. Current Genetics, 1990, 17, 247-253.	1.7	56
13	The Fatty Acid Synthase Gene is a Conserved p53 Family Target Gene from Worm to Human. Cell Cycle, 2006, 5, 750-758.	2.6	56
14	TRIM8 anti-proliferative action against chemo-resistant renal cell carcinoma. Oncotarget, 2014, 5, 7446-7457.	1.8	55
15	p73 and p63 Sustain Cellular Growth by Transcriptional Activation of Cell Cycle Progression Genes. Cancer Research, 2009, 69, 8563-8571.	0.9	51
16	Targeting Chemoresistant Tumors: Could TRIM Proteins-p53 Axis Be a Possible Answer?. International Journal of Molecular Sciences, 2019, 20, 1776.	4.1	49
17	Connecting p63 to Cellular Proliferation: The Example of the Adenosine Deaminase Target Gene. Cell Cycle, 2006, 5, 205-212.	2.6	46
18	Gamma rays induce a p53-independent mitochondrial biogenesis that is counter-regulated by HIF1α. Cell Death and Disease, 2013, 4, e663-e663.	6.3	31

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19	Transcription mapping of the Ori L region reveals novel precursors of mature RNA species and antisense RNAs in rat mitochondrial genome. FEBS Letters, 1992, 296, 311-316.	2.8	30
20	BMP-Mediated Functional Cooperation between Dlx5;Dlx6 and Msx1;Msx2 during Mammalian Limb Development. PLoS ONE, 2013, 8, e51700.	2.5	30
21	An update on the role of RANKL–RANK/osteoprotegerin and WNT-ß-catenin signaling pathways in pediatric diseases. World Journal of Pediatrics, 2019, 15, 4-11.	1.8	29
22	Pilot study on circulating miRNA signature in children with obesity born small for gestational age and appropriate for gestational age. Pediatric Obesity, 2018, 13, 803-811.	2.8	27
23	New p53 mutations in hilar cholangiocarcinoma. European Journal of Clinical Investigation, 2000, 30, 798-803.	3.4	25
24	TRIM8: Making the Right Decision between the Oncogene and Tumour Suppressor Role. Genes, 2017, 8, 354.	2.4	23
25	Transcription of rat mitochondrial NADH-dehydrogenase subunits presence of antisense and precursor RNA species. FEBS Letters, 1994, 354, 30-36.	2.8	22
26	The evolution of the RNase P- and RNase MRP-associated RNAs: Phylogenetic analysis and nucleotide substitution rate. Journal of Molecular Evolution, 1996, 43, 46-57.	1.8	20
27	Guinea Pig p53 mRNA: Identification of New Elements in Coding and Untranslated Regions and Their Functional and Evolutionary Implications. Genomics, 1999, 58, 50-64.	2.9	20
28	Characterization of p53 mutations in colorectal liver metastases and correlation with clinical parameters. Clinical Cancer Research, 1999, 5, 3523-8.	7.0	20
29	Genome-Wide Analysis of Differentially Expressed Genes and Splicing Isoforms in Clear Cell Renal Cell Carcinoma. PLoS ONE, 2013, 8, e78452.	2.5	19
30	Molecular strategies in Metazoan genomic evolution. Gene, 2002, 300, 195-201.	2.2	18
31	Adenosine deaminase, a key enzyme in DNA precursors control, is a new p73 target. Oncogene, 2003, 22, 8738-8748.	5.9	16
32	Emerging Roles of TRIM8 in Health and Disease. Cells, 2021, 10, 561.	4.1	16
33	Methods for screening tumors for p53 status and therapeutic exploitation. Expert Review of Molecular Diagnostics, 2003, 3, 289-301.	3.1	14
34	Detection of novel transcripts in the human mitochondrial DNA region coding for ATPase8-ATPase6 subunits. FEBS Letters, 1994, 344, 10-14.	2.8	13
35	TRIM Proteins in Colorectal Cancer: TRIM8 as a Promising Therapeutic Target in Chemo Resistance. Biomedicines, 2021, 9, 241.	3.2	12
36	RNase Mitochondrial RNA Processing Cleaves RNA from the Rat Mitochondrial Displacement Loop at the Origin of Heavy-Strand DNA Replication. FEBS Journal, 1995, 227, 657-662.	0.2	11

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37	TRIM8 Blunts the Pro-proliferative Action of ΔNp63α in a p53 Wild-Type Background. Frontiers in Oncology, 2019, 9, 1154.	2.8	8
38	RNase Mitochondrial RNA Processing Cleaves RNA from the Rat Mitochondrial Displacement Loop at the Origin of Heavy‧trand DNA Replication. FEBS Journal, 1995, 227, 657-662.	0.2	7
39	A platform independent RNA-Seq protocol for the detection of transcriptome complexity. BMC Genomics, 2013, 14, 855.	2.8	7
40	Leber's hereditary optic neuropathy, intellectual disability and epilepsy presenting with variable penetrance associated to the m.3460G >A mutation and a heteroplasmic expansion of the microsatellite in MTRNR1 gene – case report. BMC Medical Genetics, 2018, 19, 129.	2.1	7
41	Gene expression signature induced by grape intake in healthy subjects reveals wide-spread beneficial effects on peripheral blood mononuclear cells. Journal of Functional Foods, 2020, 64, 103705.	3.4	7
42	Regulation of IGFBP3 gene expression in short children born small for gestational age. Growth Hormone and IGF Research, 2011, 21, 349-355.	1.1	6
43	The p53 family member p73 modulates the proproliferative role of IGFBP3 in short children born small for gestational age. Molecular Biology of the Cell, 2015, 26, 2733-2741.	2.1	6
44	Tackling critical parameters in metazoan meta-barcoding experiments: a preliminary study based on <i>coxl</i> DNA barcode. PeerJ, 2018, 6, e4845.	2.0	6
45	Molecular Characterization of p53 Mutations in Primary and Secondary Liver Tumors. Molecular Biotechnology, 2002, 21, 265-278.	2.4	5
46	Identification of tumor-associated cassette exons in human cancer through EST-based computational prediction and experimental validation. Molecular Cancer, 2010, 9, 230.	19.2	5
47	Microarray data and pathway analyses of peripheral blood mononuclear cells from healthy subjects after a three weeks grape-rich diet. Data in Brief, 2020, 29, 105278.	1.0	5
48	EXPERIMENTAL COMPARISON OF MCF7 AND MCF10A RESPONSE TO LOW INTENSITY ULTRASOUND. Journal of Mechanics in Medicine and Biology, 2019, 19, 1950057.	0.7	2
49	Nuclear-mitochondrial coevolution of RNA processing enzymes and cellular function. Progress in Cell Research, 1995, , 143-147.	0.3	2
50	Mutation of p53 Tumor Suppressor Gene in Hepatocellular Carcinoma. , 2000, 45, 113-130.		1
51	Reorganization and merging of the EMBL and GenBank keyword indexes in a tree structure for more efficient retrieval of nucleic acid sequences. Protein Sequences & Data Analysis, 1990, 3, 327-34.	0.2	1