

# Anastasia Conti

## List of Publications by Year in descending order

Source: <https://exaly.com/author-pdf/6544497/publications.pdf>

Version: 2024-02-01

13  
papers

598  
citations

1040056

9  
h-index

1125743

13  
g-index

13  
all docs

13  
docs citations

13  
times ranked

1138  
citing authors

#	ARTICLE	IF	CITATIONS
1	Follicular helper T cell signature of replicative exhaustion, apoptosis, and senescence in common variable immunodeficiency. <i>European Journal of Immunology</i> , 2022, 52, 1171-1189.	2.9	9
2	Premature Senescence and Increased Oxidative Stress in the Thymus of Down Syndrome Patients. <i>Frontiers in Immunology</i> , 2021, 12, 669893.	4.8	15
3	Oncogene-induced senescence in hematopoietic progenitors features myeloid restricted hematopoiesis, chronic inflammation and histiocytosis. <i>Nature Communications</i> , 2021, 12, 4559.	12.8	17
4	Alu RNA Modulates the Expression of Cell Cycle Genes in Human Fibroblasts. <i>International Journal of Molecular Sciences</i> , 2019, 20, 3315.	4.1	10
5	Precise Gene Editing Preserves Hematopoietic Stem Cell Function following Transient p53-Mediated DNA Damage Response. <i>Cell Stem Cell</i> , 2019, 24, 551-565.e8.	11.1	237
6	An early senescence state in aged mesenchymal stromal cells contributes to hematopoietic stem and progenitor cell clonogenic impairment through the activation of a proinflammatory program. <i>Aging Cell</i> , 2019, 18, e12933.	6.7	114
7	Epigenetic and Transcriptional Modifications in Repetitive Elements in Petrol Station Workers Exposed to Benzene and MTBE. <i>International Journal of Environmental Research and Public Health</i> , 2018, 15, 735.	2.6	22
8	p53 activation: a checkpoint for precision genome editing?. <i>Genome Medicine</i> , 2018, 10, 66.	8.2	11
9	Whole-genome expression analysis of mammalian-wide interspersed repeat elements in human cell lines. <i>DNA Research</i> , 2016, 24, dsw048.	3.4	16
10	Hydroquinone induces DNA hypomethylation-independent overexpression of retroelements in human leukemia and hematopoietic stem cells. <i>Biochemical and Biophysical Research Communications</i> , 2016, 474, 691-695.	2.1	15
11	Identification of RNA polymerase III-transcribed Alu loci by computational screening of RNA-Seq data. <i>Nucleic Acids Research</i> , 2015, 43, 817-835.	14.5	55
12	Identification of RNA polymerase III-transcribed genes in eukaryotic genomes. <i>Biochimica Et Biophysica Acta - Gene Regulatory Mechanisms</i> , 2013, 1829, 296-305.	1.9	75
13	The Murine PSE/TATA-Dependent Transcriptome: Evidence of Functional Homologies with Its Human Counterpart. <i>International Journal of Molecular Sciences</i> , 2012, 13, 14813-14827.	4.1	2