

# Veronica De Sanctis

## List of Publications by Year in descending order

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Version: 2024-02-01

21  
papers

1,992  
citations

623734

14  
h-index

713466

21  
g-index

21  
all docs

21  
docs citations

21  
times ranked

3794  
citing authors

#	ARTICLE	IF	CITATIONS
1	Mother-to-Infant Microbial Transmission from Different Body Sites Shapes the Developing Infant Gut Microbiome. <i>Cell Host and Microbe</i> , 2018, 24, 133-145.e5.	11.0	822
2	HIV-1 Nef promotes infection by excluding SERINC5 from virion incorporation. <i>Nature</i> , 2015, 526, 212-217.	27.8	376
3	Genomic characterization of Nontuberculous Mycobacteria. <i>Scientific Reports</i> , 2017, 7, 45258.	3.3	176
4	Unexplored diversity and strain-level structure of the skin microbiome associated with psoriasis. <i>Npj Biofilms and Microbiomes</i> , 2017, 3, 14.	6.4	159
5	The new phylogeny of the genus <i>Mycobacterium</i> : The old and the news. <i>Infection, Genetics and Evolution</i> , 2017, 56, 19-25.	2.3	128
6	Identification of new p53 target microRNAs by bioinformatics and functional analysis. <i>BMC Cancer</i> , 2013, 13, 552.	2.6	51
7	The short-term impact of probiotic consumption on the oral cavity microbiome. <i>Scientific Reports</i> , 2018, 8, 10476.	3.3	36
8	Enhanced microbial diversity in the saliva microbiome induced by short-term probiotic intake revealed by 16S rRNA sequencing on the IonTorrent PGM platform. <i>Journal of Biotechnology</i> , 2014, 190, 30-39.	3.8	34
9	Interaction between p53 and estradiol pathways in transcriptional responses to chemotherapeutics. <i>Cell Cycle</i> , 2013, 12, 1211-1224.	2.6	32
10	Characterization of 17 strains belonging to the <i>Mycobacterium simiae</i> complex and description of <i>Mycobacterium paraense</i> sp. nov.. <i>International Journal of Systematic and Evolutionary Microbiology</i> , 2015, 65, 656-662.	1.7	31
11	OncomiR detection in circulating body fluids: a PDMS microdevice perspective. <i>Lab on A Chip</i> , 2014, 14, 4067-4075.	6.0	24
12	Manganese water-soluble porphyrin senses DNA conformation. <i>Journal of Molecular Biology</i> , 1998, 282, 43-57.	4.2	22
13	Gene expression pathways induced by axotomy and decentralization of rat superior cervical ganglion neurons. <i>European Journal of Neuroscience</i> , 2006, 23, 65-74.	2.6	22
14	ddSeeker: a tool for processing Bio-Rad ddSEQ single cell RNA-seq data. <i>BMC Genomics</i> , 2018, 19, 960.	2.8	22
15	In Vivo Topography of Rap1p-DNA Complex at <i>Saccharomyces cerevisiae</i> TEF2 UASRPG During Transcriptional Regulation. <i>Journal of Molecular Biology</i> , 2002, 318, 333-349.	4.2	16
16	Distinct DNA Elements Contribute to Rap1p Affinity for its Binding Sites. <i>Journal of Molecular Biology</i> , 2004, 338, 877-893.	4.2	12
17	A Community Study of SARS-CoV-2 Detection by RT-PCR in Saliva: A Reliable and Effective Method. <i>Viruses</i> , 2022, 14, 313.	3.3	10
18	The <i>CDKN2A/p16<sup>INK4a</sup></i> 5'UTR sequence and translational regulation: impact of novel variants predisposing to melanoma. <i>Pigment Cell and Melanoma Research</i> , 2016, 29, 210-221.	3.3	9

#	ARTICLE	IF	CITATIONS
19	The altered transcriptome of pediatric myelodysplastic syndrome revealed by RNA sequencing. Journal of Hematology and Oncology, 2020, 13, 135.	17.0	4
20	Exome sequencing in a patient with Catelâ€“Manzke-like syndrome excludes the involvement of the known genes and reveals a possible candidate. European Journal of Medical Genetics, 2015, 58, 597-602.	1.3	3
21	YES1 and MYC Amplifications as Synergistic Resistance Mechanisms to Different Generation ALK Tyrosine Kinase Inhibitors in Advanced NSCLC: Brief Report of Clinical and Preclinical Proofs. JTO Clinical and Research Reports, 2022, 3, 100278.	1.1	3