

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

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all docs

21
docs citations

21
times ranked

3794
citing authors

#	ARTICLE	IF	CITATIONS
1	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
2	A Community Study of SARS-CoV-2 Detection by RT-PCR in Saliva: A Reliable and Effective Method. Viruses, 2022, 14, 313.	3.3	10
3	The altered transcriptome of pediatric myelodysplastic syndrome revealed by RNA sequencing. Journal of Hematology and Oncology, 2020, 13, 135.	17.0	4
4	ddSeeker: a tool for processing Bio-Rad ddSEQ single cell RNA-seq data. BMC Genomics, 2018, 19, 960.	2.8	22
5	Mother-to-Infant Microbial Transmission from Different Body Sites Shapes the Developing Infant Gut Microbiome. Cell Host and Microbe, 2018, 24, 133-145.e5.	11.0	822
6	The short-term impact of probiotic consumption on the oral cavity microbiome. Scientific Reports, 2018, 8, 10476.	3.3	36
7	Genomic characterization of Nontuberculous Mycobacteria. Scientific Reports, 2017, 7, 45258.	3.3	176
8	The new phylogeny of the genus Mycobacterium : The old and the news. Infection, Genetics and Evolution, 2017, 56, 19-25.	2.3	128
9	Unexplored diversity and strain-level structure of the skin microbiome associated with psoriasis. Npj Biofilms and Microbiomes, 2017, 3, 14.	6.4	159
10	The <i>CDKN2A/p16^{INK4a}</i> 5'UTR sequence and translational regulation: impact of novel variants predisposing to melanoma. Pigment Cell and Melanoma Research, 2016, 29, 210-221.	3.3	9
11	HIV-1 Nef promotes infection by excluding SERINC5 from virion incorporation. Nature, 2015, 526, 212-217.	27.8	376
12	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
13	Characterization of 17 strains belonging to the Mycobacterium simiae complex and description of Mycobacterium paraense sp. nov.. International Journal of Systematic and Evolutionary Microbiology, 2015, 65, 656-662.	1.7	31
14	Enhanced microbial diversity in the saliva microbiome induced by short-term probiotic intake revealed by 16S rRNA sequencing on the IonTorrent PGM platform. Journal of Biotechnology, 2014, 190, 30-39.	3.8	34
15	OncomiR detection in circulating body fluids: a PDMS microdevice perspective. Lab on A Chip, 2014, 14, 4067-4075.	6.0	24
16	Identification of new p53 target microRNAs by bioinformatics and functional analysis. BMC Cancer, 2013, 13, 552.	2.6	51
17	Interaction between p53 and estradiol pathways in transcriptional responses to chemotherapeutics. Cell Cycle, 2013, 12, 1211-1224.	2.6	32
18	Gene expression pathways induced by axotomy and decentralization of rat superior cervical ganglion neurons. European Journal of Neuroscience, 2006, 23, 65-74.	2.6	22

#	ARTICLE	IF	CITATIONS
19	Distinct DNA Elements Contribute to Rap1p Affinity for its Binding Sites. <i>Journal of Molecular Biology</i> , 2004, 338, 877-893.	4.2	12
20	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
21	Manganese water-soluble porphyrin senses DNA conformation. <i>Journal of Molecular Biology</i> , 1998, 282, 43-57.	4.2	22