

Simone Puccio

List of Publications by Year in descending order

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Version: 2024-02-01

19
papers

722
citations

686830

13
h-index

794141

19
g-index

20
all docs

20
docs citations

20
times ranked

1476
citing authors

#	ARTICLE	IF	CITATIONS
1	Two subsets of stem-like CD8+ memory T cell progenitors with distinct fate commitments in humans. <i>Nature Immunology</i> , 2020, 21, 1552-1562.	7.0	167
2	IRF4 instructs effector Treg differentiation and immune suppression in human cancer. <i>Journal of Clinical Investigation</i> , 2020, 130, 3137-3150.	3.9	103
3	Antioxidant metabolism regulates CD8+ T memory stem cell formation and antitumor immunity. <i>JCI Insight</i> , 2018, 3, .	2.3	84
4	Circulating mucosal-associated invariant T cells identify patients responding to anti-PD-1 therapy. <i>Nature Communications</i> , 2021, 12, 1669.	5.8	48
5	Genomic and transcriptomic comparison between <i>Staphylococcus aureus</i> strains associated with high and low within herd prevalence of intra-mammary infection. <i>BMC Microbiology</i> , 2017, 17, 21.	1.3	45
6	Mutation and Suppressor Analysis of the Essential Lipopolysaccharide Transport Protein LptA Reveals Strategies To Overcome Severe Outer Membrane Permeability Defects in <i>Escherichia coli</i> . <i>Journal of Bacteriology</i> , 2018, 200, .	1.0	36
7	Comprehensive mapping of the <i>Helicobacter pylori</i> NikR regulon provides new insights in bacterial nickel responses. <i>Scientific Reports</i> , 2017, 7, 45458.	1.6	35
8	Insight into the essential role of the <i>Helicobacter pylori</i> HP1043 orphan response regulator: genome-wide identification and characterization of the DNA-binding sites. <i>Scientific Reports</i> , 2017, 7, 41063.	1.6	34
9	RhoTermPredict: an algorithm for predicting Rho-dependent transcription terminators based on <i>Escherichia coli</i> , <i>Bacillus subtilis</i> and <i>Salmonella enterica</i> databases. <i>BMC Bioinformatics</i> , 2019, 20, 117.	1.2	32
10	The Lack of the Essential LptC Protein in the Trans-Envelope Lipopolysaccharide Transport Machine Is Circumvented by Suppressor Mutations in LptF, an Inner Membrane Component of the <i>Escherichia coli</i> Transporter. <i>PLoS ONE</i> , 2016, 11, e0161354.	1.1	26
11	Single-cell profiling identifies impaired adaptive NK cells expanded after HCMV reactivation in haploidentical HSCT. <i>JCI Insight</i> , 2021, 6, .	2.3	19
12	Involvement of 5-Lipoxygenase in the Corticosteroid-Dependent Amyloid Beta Formation: In Vitro and In Vivo Evidence. <i>PLoS ONE</i> , 2011, 6, e15163.	1.1	19
13	WoPPER: Web server for Position Related data analysis of gene Expression in Prokaryotes. <i>Nucleic Acids Research</i> , 2017, 45, W109-W115.	6.5	16
14	Selecting soluble/foldable protein domains through single-gene or genomic ORF filtering: structure of the head domain of <i>Burkholderia pseudomallei</i> antigen BPSL2063. <i>Acta Crystallographica Section D: Biological Crystallography</i> , 2015, 71, 2227-2235.	2.5	15
15	The <i>Helicobacter pylori</i> CagY Protein Drives Gastric Th1 and Th17 Inflammation and B Cell Proliferation in Gastric MALT Lymphoma. <i>International Journal of Molecular Sciences</i> , 2021, 22, 9459.	1.8	14
16	The <i>Helicobacter pylori</i> Heat-Shock Repressor HspR: Definition of Its Direct Regulon and Characterization of the Cooperative DNA-Binding Mechanism on Its Own Promoter. <i>Frontiers in Microbiology</i> , 2018, 9, 1887.	1.5	9
17	InteractomeSeq: a web server for the identification and profiling of domains and epitopes from phage display and next generation sequencing data. <i>Nucleic Acids Research</i> , 2020, 48, W200-W207.	6.5	7
18	Defining the <i>Helicobacter pylori</i> Disease-Specific Antigenic Repertoire. <i>Frontiers in Microbiology</i> , 2020, 11, 1551.	1.5	6

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19	Interactome-Seq: A Protocol for Domainome Library Construction, Validation and Selection by Phage Display and Next Generation Sequencing. Journal of Visualized Experiments, 2018, , .	0.2	5