

Daria Bottai

List of Publications by Year in descending order

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Version: 2024-02-01

27
papers

1,648
citations

430874

18
h-index

552781

26
g-index

27
all docs

27
docs citations

27
times ranked

1972
citing authors

#	ARTICLE	IF	CITATIONS
1	Maternal anthropometric variables and clinical factors shape neonatal microbiome. <i>Scientific Reports</i> , 2022, 12, 2875.	3.3	2
2	Advances in biosensing: The CRISPR/Cas system as a new powerful tool for the detection of nucleic acids. <i>Journal of Pharmaceutical and Biomedical Analysis</i> , 2021, 192, 113645.	2.8	63
3	Genetic Manipulation as a Tool to Unravel <i>Candida parapsilosis</i> Species Complex Virulence and Drug Resistance: State of the Art. <i>Journal of Fungi (Basel, Switzerland)</i> , 2021, 7, 459.	3.5	6
4	A label-free impedance biosensing assay based on CRISPR/Cas12a collateral activity for bacterial DNA detection. <i>Journal of Pharmaceutical and Biomedical Analysis</i> , 2021, 204, 114268.	2.8	31
5	CpALS4770 and CpALS4780 contribution to the virulence of <i>Candida parapsilosis</i> . <i>Microbiological Research</i> , 2020, 231, 126351.	5.3	16
6	Role of CpALS4790 and CpALS0660 in <i>Candida parapsilosis</i> Virulence: Evidence from a Murine Model of Vaginal Candidiasis. <i>Journal of Fungi (Basel, Switzerland)</i> , 2020, 6, 86.	3.5	9
7	TbD1 deletion as a driver of the evolutionary success of modern epidemic <i>Mycobacterium tuberculosis</i> lineages. <i>Nature Communications</i> , 2020, 11, 684.	12.8	68
8	Characterization of the <i>Candida orthopsilosis</i> agglutinin-like sequence (ALS) genes. <i>PLoS ONE</i> , 2019, 14, e0215912.	2.5	16
9	A CRISPR/Cas9-based strategy to simultaneously inactivate the entire <i>ALS</i> gene family in <i>Candida orthopsilosis</i> . <i>Future Microbiology</i> , 2019, 14, 1383-1396.	2.0	12
10	Multiplexed Quantitation of Intraphagocyte <i>Mycobacterium tuberculosis</i> Secreted Protein Effectors. <i>Cell Reports</i> , 2018, 23, 1072-1084.	6.4	28
11	CoERG11 A395T mutation confers azole resistance in <i>Candida orthopsilosis</i> clinical isolates. <i>Journal of Antimicrobial Chemotherapy</i> , 2018, 73, 1815-1822.	3.0	19
12	CORTOC04210 is required for <i>Candida orthopsilosis</i> adhesion to human buccal cells. <i>Fungal Genetics and Biology</i> , 2018, 120, 19-29.	2.1	24
13	RD5-mediated lack of PE_PGRS and PPE-MPTR export in BCG vaccine strains results in strong reduction of antigenic repertoire but little impact on protection. <i>PLoS Pathogens</i> , 2018, 14, e1007139.	4.7	36
14	The N-Terminus of Human Lactoferrin Displays Anti-biofilm Activity on <i>Candida parapsilosis</i> in Lumen Catheters. <i>Frontiers in Microbiology</i> , 2017, 8, 2218.	3.5	18
15	CD4+ T Cells Recognizing PE/PPE Antigens Directly or via Cross Reactivity Are Protective against Pulmonary <i>Mycobacterium tuberculosis</i> Infection. <i>PLoS Pathogens</i> , 2016, 12, e1005770.	4.7	50
16	The BCG Strain Pool: Diversity Matters. <i>Molecular Therapy</i> , 2016, 24, 201-203.	8.2	14
17	Revisiting the role of phospholipases C in virulence and the lifecycle of <i>Mycobacterium tuberculosis</i> . <i>Scientific Reports</i> , 2015, 5, 16918.	3.3	39
18	Impact of <i>Mycobacterium tuberculosis</i> RD1-locus on human primary dendritic cell immune functions. <i>Scientific Reports</i> , 2015, 5, 17078.	3.3	18

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19	Type VII Secretion Systems in Gram-Positive Bacteria. <i>Current Topics in Microbiology and Immunology</i> , 2015, 404, 235-265.	1.1	33
20	Increased protective efficacy of recombinant BCG strains expressing virulence-neutral proteins of the ESX-1 secretion system. <i>Vaccine</i> , 2015, 33, 2710-2718.	3.8	51
21	Mycobacterial Pathogenomics and Evolution. <i>Microbiology Spectrum</i> , 2014, 2, MGM2-0025-2013.	3.0	36
22	Strong Immunogenicity and Cross-Reactivity of Mycobacterium tuberculosis ESX-5 Type VII Secretion-Encoded PE-PPE Proteins Predicts Vaccine Potential. <i>Cell Host and Microbe</i> , 2012, 11, 352-363.	11.0	102
23	The ESX-5 Associated eccB5-eccC5 Locus Is Essential for Mycobacterium tuberculosis Viability. <i>PLoS ONE</i> , 2012, 7, e52059.	2.5	49
24	ESX-1 α -mediated translocation to the cytosol controls virulence of mycobacteria. <i>Cellular Microbiology</i> , 2012, 14, 1287-1298.	2.1	375
25	ESAT-6 Secretion-Independent Impact of ESX-1 Genes espF and espG1 on Virulence of Mycobacterium tuberculosis. <i>Journal of Infectious Diseases</i> , 2011, 203, 1155-1164.	4.0	66
26	Systematic Genetic Nomenclature for Type VII Secretion Systems. <i>PLoS Pathogens</i> , 2009, 5, e1000507.	4.7	233
27	Control of M. tuberculosis ESAT-6 Secretion and Specific T Cell Recognition by PhoP. <i>PLoS Pathogens</i> , 2008, 4, e33.	4.7	234