

Fabiana Bigi

List of Publications by Year in descending order

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Version: 2024-02-01

52
papers

1,783
citations

331670

21
h-index

276875

41
g-index

52
all docs

52
docs citations

52
times ranked

2245
citing authors

#	ARTICLE	IF	CITATIONS
1	Development and evaluation of a low cost IgG ELISA test based in RBD protein for COVID-19. <i>Journal of Immunological Methods</i> , 2022, 500, 113182.	1.4	8
2	Replication and transmission features of two experimental vaccine candidates against bovine tuberculosis subcutaneously administrated in a murine model. <i>Tuberculosis</i> , 2022, 134, 102203.	1.9	0
3	Role of PhoPR in the response to stress of <i>Mycobacterium bovis</i> . <i>Comparative Immunology, Microbiology and Infectious Diseases</i> , 2021, 74, 101593.	1.6	6
4	Recent advances in non-specific immune memory against bovine tuberculosis. <i>Comparative Immunology, Microbiology and Infectious Diseases</i> , 2021, 75, 101615.	1.6	5
5	The subunit vaccine H65+CAF01 increased the BCG- protection against <i>Mycobacterium bovis</i> infection in a mouse model of bovine tuberculosis. <i>Research in Veterinary Science</i> , 2021, 136, 595-597.	1.9	7
6	Identifying Bacterial and Host Factors Involved in the Interaction of <i>Mycobacterium bovis</i> with the Bovine Innate Immune Cells. <i>Frontiers in Immunology</i> , 2021, 12, 674643.	4.8	3
7	Semi-stable production of bovine IL-4 and GM-CSF in the mammalian episomal expression system. <i>Journal of Veterinary Research (Poland)</i> , 2021, 65, 315-321.	1.0	1
8	H65 fusion protein fails to improve the protection of a rationally attenuated live vaccine candidate against bovine tuberculosis in a mouse model of tuberculosis. <i>International Journal of Mycobacteriology</i> , 2021, 10, 411.	0.6	1
9	Does <i>Mycobacterium bovis</i> persist in cattle in a non-replicative latent state as <i>Mycobacterium tuberculosis</i> in human beings?. <i>Veterinary Microbiology</i> , 2020, 247, 108758.	1.9	16
10	FasR Regulates Fatty Acid Biosynthesis and Is Essential for Virulence of <i>Mycobacterium tuberculosis</i> . <i>Frontiers in Microbiology</i> , 2020, 11, 586285.	3.5	1
11	A Phenotypic Characterization of Two Isolates of a Multidrug-Resistant Outbreak Strain of <i>Mycobacterium tuberculosis</i> with Opposite Epidemiological Fitness. <i>BioMed Research International</i> , 2020, 2020, 1-9.	1.9	2
12	Elimination of ESAT-6 and CFP-10 from a candidate vaccine against bovine tuberculosis impaired its protection efficacy in the BALBc mouse model. <i>International Journal of Mycobacteriology</i> , 2020, 9, 417.	0.6	5
13	Editorial: Cellular and Molecular Mechanisms of <i>Mycobacterium tuberculosis</i> Virulence. <i>Frontiers in Cellular and Infection Microbiology</i> , 2019, 9, 331.	3.9	11
14	Production of <i>Mycobacterium bovis</i> Antigens Included in Recombinant Occlusion Bodies of Baculovirus. <i>Journal of Molecular Microbiology and Biotechnology</i> , 2019, 29, 83-90.	1.0	1
15	Analysing nonsynonymous mutations between two <i>Mycobacterium bovis</i> strains with contrasting pathogenic profiles. <i>Veterinary Microbiology</i> , 2019, 239, 108482.	1.9	5
16	Rv2617c and P36 are virulence factors of pathogenic mycobacteria involved in resistance to oxidative stress. <i>Virulence</i> , 2019, 10, 1026-1033.	4.4	11
17	Identification of bovine tuberculosis biomarkers to detect tuberculin skin test and IFN- γ release assay false negative cattle. <i>Research in Veterinary Science</i> , 2019, 122, 7-14.	1.9	26
18	<i>Mycobacterium bovis</i> ESAT-6, CFP-10 and EspC antigens show high conservation among field isolates. <i>Tuberculosis</i> , 2018, 111, 143-146.	1.9	9

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19	Characterization of the two component regulatory system PhoPR in <i>Mycobacterium bovis</i> . <i>Veterinary Microbiology</i> , 2018, 222, 30-38.	1.9	17
20	Single nucleotide polymorphisms may explain the contrasting phenotypes of two variants of a multidrug-resistant <i>Mycobacterium tuberculosis</i> strain. <i>Tuberculosis</i> , 2017, 103, 28-36.	1.9	10
21	<i>Mycobacterium bovis</i> Requires P27 (LprG) To Arrest Phagosome Maturation and Replicate within Bovine Macrophages. <i>Infection and Immunity</i> , 2017, 85, .	2.2	25
22	<i>ERAP1</i> and <i>PDE8A</i> Are Downregulated in Cattle Protected against Bovine Tuberculosis. <i>Journal of Molecular Microbiology and Biotechnology</i> , 2017, 27, 237-245.	1.0	10
23	Metabolic profile of <i>Mycobacterium smegmatis</i> reveals Mce4 proteins are relevant for cell wall lipid homeostasis. <i>Metabolomics</i> , 2016, 12, 1.	3.0	8
24	Polymorphisms of 20 regulatory proteins between <i>Mycobacterium tuberculosis</i> and <i>Mycobacterium bovis</i>. <i>Microbiology and Immunology</i> , 2016, 60, 552-560.	1.4	14
25	Evaluation of <i>Mycobacterium bovis</i> double knockout mce2-phoP as candidate vaccine against bovine tuberculosis. <i>Tuberculosis</i> , 2015, 95, 186-189.	1.9	9
26	Identification and evaluation of new <i>Mycobacterium bovis</i> antigens in the in vitro interferon gamma release assay for bovine tuberculosis diagnosis. <i>Tuberculosis</i> , 2015, 95, 795-801.	1.9	12
27	Assessment of Deleted in p27-p55 Virulence Operon as Candidate Vaccine against Tuberculosis in Animal Models. <i>BioMed Research International</i> , 2014, 2014, 1-6.	1.9	3
28	IFNG-mediated immune responses enhance autophagy against <i>Mycobacterium tuberculosis</i> antigens in patients with active tuberculosis. <i>Autophagy</i> , 2014, 10, 2109-2121.	9.1	63
29	Role of the Mce1 transporter in the lipid homeostasis of <i>Mycobacterium tuberculosis</i> . <i>Tuberculosis</i> , 2014, 94, 170-177.	1.9	104
30	<i>Mycobacterium bovis</i> mce2 double deletion mutant protects cattle against challenge with virulent M. bovis. <i>Tuberculosis</i> , 2013, 93, 363-372.	1.9	31
31	Draft Genome Sequence of <i>Mycobacterium bovis</i> 04-303, a Highly Virulent Strain from Argentina. <i>Genome Announcements</i> , 2013, 1, .	0.8	9
32	Virulence factors of the <i>Mycobacterium tuberculosis</i> complex. <i>Virulence</i> , 2013, 4, 3-66.	4.4	543
33	Assessment of the Immune Responses Induced in Cattle after Inoculation of a <i>Mycobacterium bovis</i> Strain Deleted in Two <i>mce2</i> Genes. <i>Journal of Biomedicine and Biotechnology</i> , 2012, 2012, 1-8.	3.0	14
34	Impact of the deletion of the six mce operons in <i>Mycobacterium smegmatis</i> . <i>Microbes and Infection</i> , 2012, 14, 590-599.	1.9	52
35	Transcriptional Response of Peripheral Blood Mononuclear Cells from Cattle Infected with <i>Mycobacterium bovis</i> . <i>PLoS ONE</i> , 2012, 7, e41066.	2.5	34
36	Knockout mutation of <i>p27-p55</i> operon severely reduces replication of <i>Mycobacterium bovis</i> in a macrophagic cell line and survival in a mouse model of infection. <i>Virulence</i> , 2011, 2, 233-237.	4.4	24

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37	Increased IL-17 expression is associated with pathology in a bovine model of tuberculosis. <i>Tuberculosis</i> , 2011, 91, 57-63.	1.9	46
38	Role of P27-P55 operon from <i>Mycobacterium tuberculosis</i> in the resistance to toxic compounds. <i>BMC Infectious Diseases</i> , 2011, 11, 195.	2.9	38
39	Evaluation of pathogenesis caused in cattle and guinea pig by a <i>Mycobacterium bovis</i> strain isolated from wild boar. <i>BMC Veterinary Research</i> , 2011, 7, 37.	1.9	23
40	Identification of four novel DC-SIGN ligands on <i>Mycobacterium bovis</i> BCG. <i>Protein and Cell</i> , 2010, 1, 859-870.	11.0	48
41	Mce3R, a TetR-type transcriptional repressor, controls the expression of a regulon involved in lipid metabolism in <i>Mycobacterium tuberculosis</i> . <i>Microbiology (United Kingdom)</i> , 2009, 155, 2245-2255.	1.8	62
42	Identification of two proteins that interact with the Erp virulence factor from <i>Mycobacterium tuberculosis</i> by using the bacterial two-hybrid system. <i>BMC Molecular Biology</i> , 2009, 10, 3.	3.0	20
43	Study of the immunological profile towards <i>Mycobacterium bovis</i> antigens in naturally infected cattle. <i>Microbiology and Immunology</i> , 2009, 53, 460-467.	1.4	21
44	Differential transcriptome profiles of attenuated and hypervirulent strains of <i>Mycobacterium bovis</i> . <i>Microbes and Infection</i> , 2009, 11, 956-963.	1.9	31
45	Study of the role of Mce3R on the transcription of mce genes of <i>Mycobacterium tuberculosis</i> . <i>BMC Microbiology</i> , 2008, 8, 38.	3.3	40
46	Immunogenicity and protection induced by <i>Mycobacterium tuberculosis</i> mce-2 and mce-3 mutants in a Balb/c mouse model of progressive pulmonary tuberculosis. <i>Vaccine</i> , 2006, 24, 2333-2342.	3.8	27
47	Mutation in mce operons attenuates <i>Mycobacterium tuberculosis</i> virulence. <i>Microbes and Infection</i> , 2005, 7, 325-334.	1.9	148
48	Identification of genetic markers for <i>Mycobacterium pinnipedi</i> through genome analysis. <i>FEMS Microbiology Letters</i> , 2005, 248, 147-152.	1.8	9
49	Mutation in the P36 gene of <i>Mycobacterium bovis</i> provokes attenuation of the bacillus in a mouse model. <i>Tuberculosis</i> , 2005, 85, 221-226.	1.9	14
50	The knockout of the lprG-Rv1410 operon produces strong attenuation of <i>Mycobacterium tuberculosis</i> . <i>Microbes and Infection</i> , 2004, 6, 182-187.	1.9	61
51	Negative transcriptional regulation of the mce3 operon in <i>Mycobacterium tuberculosis</i> . <i>Microbiology (United Kingdom)</i> , 2002, 148, 2997-3006.	1.8	52
52	A novel 27 kDa lipoprotein antigen from <i>Mycobacterium bovis</i> . <i>Microbiology (United Kingdom)</i> , 1997, 143, 3599-3605.	1.8	43