

David Kornspan

List of Publications by Year in descending order

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Version: 2024-02-01

10
papers

52
citations

2148532

4
h-index

1905433

7
g-index

10
all docs

10
docs citations

10
times ranked

80
citing authors

#	ARTICLE	IF	CITATIONS
1	Transcriptomic Analysis of the <i>Brucella melitensis</i> Rev.1 Vaccine Strain in an Acidic Environment: Insights Into Virulence Attenuation. <i>Frontiers in Microbiology</i> , 2019, 10, 250.	1.5	14
2	Genomic analysis of the original Elberg <i>Brucella melitensis</i> Rev.1 vaccine strain reveals insights into virulence attenuation. <i>Virulence</i> , 2018, 9, 1436-1448.	1.8	10
3	Complete Genome Sequence of the Live Attenuated Vaccine Strain <i>Brucella melitensis</i> Rev.1. <i>Genome Announcements</i> , 2018, 6, .	0.8	9
4	Genomic Analysis of Natural Rough <i>Brucella melitensis</i> Rev.1 Vaccine Strains: Identification and Characterization of Mutations in Key Genes Associated with Bacterial LPS Biosynthesis and Virulence. <i>International Journal of Molecular Sciences</i> , 2020, 21, 9341.	1.8	5
5	Transcriptomic analysis of smooth versus rough <i>Brucella melitensis</i> Rev.1 vaccine strains reveals insights into virulence attenuation. <i>International Journal of Medical Microbiology</i> , 2020, 310, 151363.	1.5	4
6	Protein Biomarker Identification for the Discrimination of <i>Brucella melitensis</i> Field Isolates From the <i>Brucella melitensis</i> Rev.1 Vaccine Strain by MALDI-TOF MS. <i>Frontiers in Microbiology</i> , 2021, 12, 712601.	1.5	4
7	Whole Genome Sequence Analysis of <i>Brucella melitensis</i> Phylogeny and Virulence Factors. <i>Microbiology Research</i> , 2021, 12, 698-710.	0.8	3
8	Genomic Epidemiology of Clinical <i>Brucella melitensis</i> Isolates from Southern Israel. <i>Microorganisms</i> , 2022, 10, 238.	1.6	2
9	Differential functions of TLE1 and TLE3 depending on a specific phosphorylation site. <i>Biochemical and Biophysical Research Communications</i> , 2021, 545, 164-170.	1.0	1
10	The Acidic Stress Response of the Intracellular Pathogen <i>Brucella melitensis</i> : New Insights from a Comparative, Genome-Wide Transcriptome Analysis. <i>Genes</i> , 2020, 11, 1016.	1.0	0