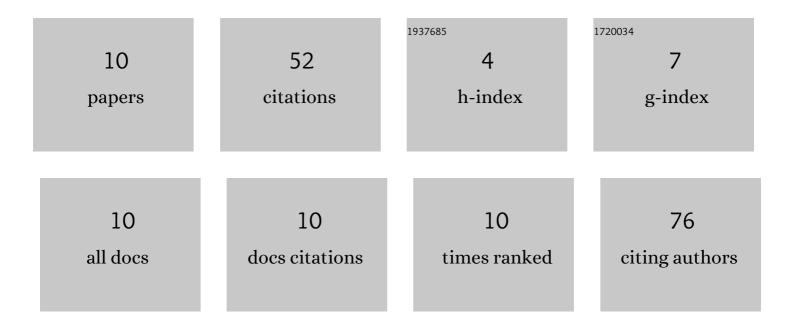
## David Kornspan

List of Publications by Year in descending order

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DAVID KODNEDAN

| #  | Article   | IF  | CITATIONS |
|----|---|-----|-----------|
| 1  | Transcriptomic Analysis of the Brucella melitensis Rev.1 Vaccine Strain in an Acidic Environment:<br>Insights Into Virulence Attenuation. Frontiers in Microbiology, 2019, 10, 250.   | 3.5 | 14        |
| 2  | Genomic analysis of the original Elberg <i>Brucella melitensis</i> Rev.1 vaccine strain reveals insights into virulence attenuation. Virulence, 2018, 9, 1436-1448.   | 4.4 | 10        |
| 3  | Complete Genome Sequence of the Live Attenuated Vaccine Strain Brucella melitensis Rev.1. Genome<br>Announcements, 2018, 6, .   | 0.8 | 9         |
| 4  | Genomic Analysis of Natural Rough Brucella melitensis Rev.1 Vaccine Strains: Identification and<br>Characterization of Mutations in Key Genes Associated with Bacterial LPS Biosynthesis and Virulence.<br>International Journal of Molecular Sciences, 2020, 21, 9341. | 4.1 | 5         |
| 5  | Transcriptomic analysis of smooth versus rough Brucella melitensis Rev.1 vaccine strains reveals<br>insights into virulence attenuation. International Journal of Medical Microbiology, 2020, 310, 151363.  | 3.6 | 4         |
| 6  | Protein Biomarker Identification for the Discrimination of Brucella melitensis Field Isolates From the<br>Brucella melitensis Rev.1 Vaccine Strain by MALDI-TOF MS. Frontiers in Microbiology, 2021, 12, 712601.  | 3.5 | 4         |
| 7  | Whole Genome Sequence Analysis of Brucella melitensis Phylogeny and Virulence Factors.<br>Microbiology Research, 2021, 12, 698-710.   | 1.9 | 3         |
| 8  | Genomic Epidemiology of Clinical Brucella melitensis Isolates from Southern Israel. Microorganisms, 2022, 10, 238.  | 3.6 | 2         |
| 9  | Differential functions of TLE1 and TLE3 depending on a specific phosphorylation site. Biochemical and Biophysical Research Communications, 2021, 545, 164-170.  | 2.1 | 1         |
| 10 | The Acidic Stress Response of the Intracellular Pathogen Brucella melitensis: New Insights from a<br>Comparative, Genome-Wide Transcriptome Analysis. Genes, 2020, 11, 1016.  | 2.4 | 0         |