

Juan Marzosa Fandiño

List of Publications by Year in descending order

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Version: 2024-02-01

16
papers

532
citations

759233

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940533

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16
times ranked

802
citing authors

| # | ARTICLE | IF | CITATIONS |
|----|---|-----|-----------|
| 1 | Poultry as reservoir for extraintestinal pathogenic <i>Escherichia coli</i> O45:K1:H7-B2-ST95 in humans. <i>Veterinary Microbiology</i> , 2013, 167, 506-512. | 1.9 | 87 |
| 2 | Four Main Virotypes among Extended-Spectrum-β-Lactamase-Producing Isolates of <i>Escherichia coli</i> O25b:H4-B2-ST131: Bacterial, Epidemiological, and Clinical Characteristics. <i>Journal of Clinical Microbiology</i> , 2013, 51, 3358-3367. | 3.9 | 76 |
| 3 | Molecular epidemiology and virulence of <i>Escherichia coli</i> O16:H5-ST131: Comparison with H30 and H30-Rx subclones of O25b:H4-ST131. <i>International Journal of Medical Microbiology</i> , 2014, 304, 1247-1257. | 3.6 | 64 |
| 4 | Emergence of new variants of ST131 clonal group among extraintestinal pathogenic <i>Escherichia coli</i> producing extended-spectrum β-lactamases. <i>International Journal of Antimicrobial Agents</i> , 2013, 42, 347-351. | 2.5 | 55 |
| 5 | Detection of quinolone-resistant <i>Escherichia coli</i> isolates belonging to clonal groups O25b:H4-B2-ST131 and O25b:H4-D-ST69 in raw sewage and river water in Barcelona, Spain. <i>Journal of Antimicrobial Chemotherapy</i> , 2013, 68, 758-765. | 3.0 | 44 |
| 6 | Virulence Patterns in a Murine Sepsis Model of ST131 <i>Escherichia coli</i> Clinical Isolates Belonging to Serotypes O25b:H4 and O16:H5 Are Associated to Specific Virotypes. <i>PLoS ONE</i> , 2014, 9, e87025. | 2.5 | 44 |
| 7 | Impact of human-associated <i>Escherichia coli</i> clonal groups in Antarctic pinnipeds: presence of ST73, ST95, ST141 and ST131. <i>Scientific Reports</i> , 2018, 8, 4678. | 3.3 | 37 |
| 8 | Diversity of Shiga toxin-producing <i>Escherichia coli</i> in sheep flocks of Paraná State, southern Brazil. <i>Veterinary Microbiology</i> , 2015, 175, 150-156. | 1.9 | 34 |
| 9 | Analysis of outer membrane porin complexes of <i>Neisseria meningitidis</i> in wild-type and specific knock-out mutant strains. <i>Proteomics</i> , 2009, 9, 648-656. | 2.2 | 17 |
| 10 | Identification of <i>Neisseria meningitidis</i> Outer Membrane Vesicle Complexes Using 2-D High Resolution Clear Native/SDS-PAGE. <i>Journal of Proteome Research</i> , 2010, 9, 611-619. | 3.7 | 17 |
| 11 | Lambs are an important source of atypical enteropathogenic <i>Escherichia coli</i> in southern Brazil. <i>Veterinary Microbiology</i> , 2016, 196, 72-77. | 1.9 | 15 |
| 12 | Induction of immune responses by purified outer membrane protein complexes from <i>Neisseria meningitidis</i> . <i>Vaccine</i> , 2012, 30, 2387-2395. | 3.8 | 13 |
| 13 | A cross-reactive neisserial antigen encoded by the NMB0035 locus shows high sequence conservation but variable surface accessibility. <i>Journal of Medical Microbiology</i> , 2008, 57, 80-87. | 1.8 | 11 |
| 14 | Characterisation and immune responses to meningococcal recombinant porin complexes incorporated into liposomes. <i>Vaccine</i> , 2009, 27, 5338-5343. | 3.8 | 7 |
| 15 | High Resolution Clear Native Electrophoresis (hrCNE) Allows a Detailed Analysis of the Heterotrimeric Structure of Recombinant <i>Neisseria meningitidis</i> Porins Inserted into Liposomes. <i>Journal of Proteome Research</i> , 2013, 12, 777-784. | 3.7 | 6 |
| 16 | Liposomes or traditional adjuvants: induction of bactericidal activity by the macrophage infectivity potentiator protein (Mip) of <i>Neisseria meningitidis</i> . <i>Apmis</i> , 2017, 125, 725-731. | 2.0 | 5 |