

Sonja M K Schoenfelder

List of Publications by Year in descending order

Source: <https://exaly.com/author-pdf/9760906/publications.pdf>

Version: 2024-02-01

8

papers

310

citations

1307594

7

h-index

1588992

8

g-index

8

all docs

8

docs citations

8

times ranked

627

citing authors

| # | ARTICLE | IF | CITATIONS |
|---|--|------|-----------|
| 1 | Another layer of complexity in <i>S. aureus</i> methionine biosynthesis control: unusual RNase III-driven T-box riboswitch cleavage determines <i>met</i> operon mRNA stability and decay. <i>Nucleic Acids Research</i> , 2021, 49, 2192-2212. | 14.5 | 7 |
| 2 | A non-coding RNA from the intercellular adhesion (<i>ica</i>) locus of <i>S. epidermidis</i> controls polysaccharide intercellular adhesion (PIA)-mediated biofilm formation. <i>Molecular Microbiology</i> , 2019, 111, 1571-1591. | 2.5 | 25 |
| 3 | The small non-coding RNA RsaE influences extracellular matrix composition in <i>S. epidermidis</i> biofilm communities. <i>PLoS Pathogens</i> , 2019, 15, e1007618. | 4.7 | 33 |
| 4 | Antibiotic resistance profiles of coagulase-negative staphylococci in livestock environments. <i>Veterinary Microbiology</i> , 2017, 200, 79-87. | 1.9 | 55 |
| 5 | Genotyping of community-associated methicillin resistant <i>S. aureus</i> (CA-MRSA) in a tertiary care centre in Mysore, South India: ST2371-SCCmec IV emerges as the major clone. <i>Infection, Genetics and Evolution</i> , 2015, 34, 230-235. | 2.3 | 32 |
| 6 | Methionine Biosynthesis in <i>S. aureus</i> Is Tightly Controlled by a Hierarchical Network Involving an Initiator tRNA-Specific T-box Riboswitch. <i>PLoS Pathogens</i> , 2013, 9, e1003606. | 4.7 | 23 |
| 7 | Hypervariability of Biofilm Formation and Oxacillin Resistance in a <i>S. epidermidis</i> Strain Causing Persistent Severe Infection in an Immunocompromised Patient. <i>Journal of Clinical Microbiology</i> , 2010, 48, 2407-2412. | 3.9 | 15 |
| 8 | Success through diversity – How <i>S. epidermidis</i> establishes as a nosocomial pathogen. <i>International Journal of Medical Microbiology</i> , 2010, 300, 380-386. | 3.6 | 120 |