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Stable antibiotic resistance and rapid human adaptation in livestock-associated MRSA

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12	Prophage-encoded immune evasion factors are critical for <i>Staphylococcus aureus</i> host infection, switching, and adaptation. 2022 , 100194		1
11	Absence of <i>Staphylococcus aureus</i> in wild populations of fish supports a spillover hypothesis.		0
10	Low-level colonization of methicillin-resistant <i>Staphylococcus aureus</i> in pigs is maintained by slowly evolving, closely related strains in Finnish pig farms. 2022 , 64,		0
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8	Clonal relatedness of coagulase-positive staphylococci among healthy dogs and dog-owners in Spain. Detection of multidrug-resistant-MSSA-CC398 and novel linezolid-resistant-MRSA-CC5. 14,		0
7	Genome sequencing and comparative genomic analysis of bovine mastitis-associated <i>Staphylococcus aureus</i> strains from India. 2023 , 24,		0
6	Comparative review of the nasal carriage and genetic characteristics of <i>Staphylococcus aureus</i> in healthy livestock: Insight into zoonotic and anthroponotic clones. 2023 , 109, 105408		1
5	High Prevalence of Livestock-Associated Methicillin-Resistant <i>Staphylococcus aureus</i> in Hungarian Pig Farms and Genomic Evidence for the Spillover of the Pathogen to Humans. 2023 , 2023, 1-13		0
4	High-Throughput Mutagenesis Reveals a Role for Antimicrobial Resistance- and Virulence-Associated Mobile Genetic Elements in <i>Staphylococcus aureus</i> Host Adaptation. 2023 , 11,		0
3	Immunogenicity and protective efficacy of a multi-epitope recombinant toxin antigen of <i>Pasteurella multocida</i> against virulent challenge in mice. 2023 , 41, 2387-2396		0
2	Monitoring Longitudinal Trends and Assessment of the Health Risk of <i>Shigella flexneri</i> Antimicrobial Resistance. 2023 , 57, 4971-4983		0
1	Molecular Epidemiology and Characterization of Multidrug-Resistant MRSA ST398 and ST239 in Himachal Pradesh, India. Volume 16, 2339-2348		0