

CITATION REPORT

List of articles citing

Whole-genome sequencing analysis of Shiga toxin-producing *Escherichia coli* O22:H8 isolated from cattle prediction pathogenesis and colonization factors and position in STEC universe phylogeny

DOI: 10.1007/s12275-022-1616-z
Journal of Microbiology, , , .

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#	Paper	IF	Citations
2	MEGARes and AMR++, v3.0: an updated comprehensive database of antimicrobial resistance determinants and an improved software pipeline for classification using high-throughput sequencing.		0
1	Genomes-based MLST, cgMLST, wgMLST and SNP analysis of Salmonella Typhimurium from animals and humans. 2023 , 96, 101973		0