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High Prevalence of Klebsiella pneumoniae in European Food Products: a Multicentric Study Comparing Culture and Molecular Detection Methods.

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| 11 | Plasmid overlap and evolution between Enterobacterales isolates from bloodstream infections and non-human compartments.   |     |           |
| 10 | Phenotypic and Genetic Characterization of Klebsiella pneumoniae Isolates from Wild Animals in Central Italy. <i>Animals</i> , <b>2022</b> , 12, 1347   | 3.1 | 2         |
| 9  | Detection of Klebsiella pneumoniae human gut carriage: a comparison of culture, qPCR, and whole metagenomic sequencing methods. <b>2022</b> , 14,   |     | O         |
| 8  | Limited Transmission of Klebsiella pneumoniae among Humans, Animals, and the Environment in a Caribbean Island, Guadeloupe (French West Indies).  |     | 0         |
| 7  | A Meta-Analysis to Estimate Prevalence of Resistance to Tetracyclines and Third Generation Cephalosporins in Enterobacteriaceae Isolated from Food Crops. <b>2022</b> , 11, 1424                |     | O         |
| 6  | Geno- and Phenotypic Characteristics of a Klebsiella pneumoniae ST20 Isolate with Unusual Colony Morphology. <b>2022</b> , 10, 2063   |     | 0         |
| 5  | Repeated cross-sectional study identifies differing risk factors associated with microbial contamination in common food products in the United Kingdom. <b>2023</b> , 111, 104196               |     | O         |
| 4  | KbvR mutant of Klebsiella pneumoniae affects the synthesis of type 1 fimbriae and provides protection to mice as a live attenuated vaccine. <b>2022</b> , 53,                                   |     | О         |
| 3  | Draft Genomic Sequence for Klebsiella pneumoniae 060517CS3-g, a Bacterium Harboring Multidrug<br>Resistance Genes, Isolated from Retail Ground Chicken Meat.                                    |     | O         |
| 2  | Multidrug resistant Klebsiella Pneumoniae reservoir and their capsular resistance genes in cow farms of district Peshawar, Pakistan. <b>2023</b> , 18, e0282245                                 |     | O         |
| 1  | Profiling bacterial communities and foodborne pathogens on food-associated surface following contact with raw beef, chicken and pork using 16S amplicon metagenomics. <b>2023</b> , 149, 109698 |     | О         |