

CORRIGENDUM

High Prevalence of KPC-2-Producing Hypervirulent Klebsiella pneumoniae Causing Meningitis in Eastern China [Corrigendum]

Xu M, Fu Y, Fang Y, et al. Infect Drug Resist. 2019;12:641-653.

Page 647, Molecular characteristics of the *K. pneumoniae* isolates, first paragraph, right column, second sentence from the bottom, the text "The PFGE results showed that the homology of 44 isolates was diverse (Figure 3)" should read "The PFGE results showed that the homology of 44 isolates was diverse (Figure 4)".

Page 648, left column, last sentence, the text "In general, ST11 KPC-2-producing hvKP strains harbored more antibioticresistant genes and less virulence factors when compared with the non-ST11 KPC-2-producing hvKP (Figure 5)" should read "In general, ST11 KPC-2-producing hvKP strains harbored more antibioticresistant genes and less virulence factors when compared with the non-ST11 KPC-2-producing hvKP (Figure 6)".

Page 648, right column, first sentence, the text "Among the 15 KPC-2-producing hvKP strains, 4 strains (KP35, KP37, KP43, and KP44) shared >75% similarity and classified into one cluster (Figure 3), while the remaining 11 strains exhibited distinct genetic relationship" should read "Among the 15 KPC-2-producing hvKP strains, 4 strains (KP35, KP37, KP43, and KP44) shared >75% similarity and classified into one cluster (Figure 4), while the remaining 11 strains exhibited distinct genetic relationship".

Page 649, left column, second paragraph, first sentence, the text "Consistent with the epidemiology trend of *K. pneumoniae* meningitis in some Asian countries especially in Taiwan,^{3,4}" should read "Consistent with the epidemiology trend of *K. pneumoniae* meningitis in Taiwan,^{3,4}".

The authors apologize for these errors.

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