## **CORRIGENDUM**

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## Corrigendum: Genomic evolution and local epidemiology of Klebsiella pneumoniae from a major hospital in Beijing, China, over a 15 year period: dissemination of known and novel highrisk clones

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In the published version of this article, the corresponding author was listed incorrectly. The corresponding author statement should have read as follows:

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Additionally, in the first paragraph of the discussion section some text and accompanying references were mistakenly omitted. The omitted references and corrected text with corresponding citations are detailed below:

'This study aimed to understand the population structure of *K. pneumoniae* within the People's Liberation Army General Hospital (H301) in Beijing over a 15 year period. While several studies have investigated the genetic epidemiology of CR-Kp in China in general [19, 25, 26] and in Hospital 301 in particular [1–6], this study represents the first longitudinal investigation focusing on the broad *K. pneumoniae* population from China.'

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- 3. Luo Y, Wang Y, Ye L, Yang J. Molecular epidemiology and virulence factors of pyogenic liver abscess causing *Klebsiella pneumoniae* in China. *Clin Microbiol Infect*. 2014;20(11):O818–24
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- 6. Lai K, Ma Y, Guo L, An J, Ye L, *et al.* Molecular characterization of clinical IMP-producing *Klebsiella pneumoniae* isolates from a Chinese Tertiary hospital. *Ann Clin Microbiol Antimicrob*. 2017;16(1):42

The authors apologise for any inconvenience caused.

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