Overestimated discriminatory power of MALDI-TOF mass spectrometry for typing of carbapenem-resistant \textit{Klebsiella pneumonia} clones – CORRIGENDUM

Fei Jiang, Ziyian Kong, Chen Cheng, Haiquan Kang, Bing Gu and Ping Ma

doi: https://doi.org/10.1017/S0950268819002097, Published online by Cambridge University Press, 17 December 2019

Original:

\begin{figure}
\centering
\includegraphics[width=\textwidth]{Fig_1.png}
\caption{Typing results of PFGE and MLST from 44 CRKP isolates. Two PFGE clones belong to ST337 and ST11, respectively.}
\end{figure}

© The Author(s), 2020. Published by Cambridge University Press. This is an Open Access article, distributed under the terms of the Creative Commons Attribution licence (http://creativecommons.org/licenses/by/4.0/), which permits unrestricted re-use, distribution, and reproduction in any medium, provided the original work is properly cited.
Correction:

Fig. 1. Typing results of PFGE and MLST from 44 CRKP isolates. Two PFGE clones belong to ST337 and ST11, respectively.
Fig. 2. Three-dimensional plot generated by MALDI-TOF MS. Four PCA clusters were identified. Strains from distinct PFGE clusters (PC) were grouped into one MALDI-TOF cluster, and strains belonging to the same PFGE cluster were assigned to more than one MALDI-TOF clusters.
Correction:

Reference