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## Correction to: epidemiological and genomic characteristics of *A. baumannii* from different infection sites using comparative genomics



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Following publication of the original article [1], it was reported that there was an error in Fig. 4. The image used was a duplicate of that for Fig. 5. The correct Fig. 4 is provided in this Correction article and the original article has been updated.

The original article can be found online at https://doi.org/10.1186/s12864-021-07842-5

Full list of author information is available at the end of the article



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**Fig. 4** Sequence types of 64 *A. baumannii* isolates from three infection sites (**A**). eBURST analysis of 64 *A. baumannii* isolates (**B**). Each solid circle represents one sequence type and its size represents the quantity of isolates of this type. Each line between solid circles indicates one allele variation

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