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Sandra Reuter, M. Estée Török, Matthew T.G. Holden, et al.

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Corrigendum: Building a genomic framework for prospective MRSA surveillance in the United Kingdom and the Republic of Ireland

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The authors would like to correct the omission of a reference in the first introductory paragraph of the initial publication of this article. The reference has been included in the following sentence:

However, for both lineages, the acquisition of the *SCC_{mec}* element conferring methicillin resistance and the presence of mutations in *gyrA* and *grrA* conferring fluoroquinolone resistance are considered to be major contributors to the success of these epidemic lineages (Horváth et al. 2012; Knight et al. 2012; McAdam et al. 2012; Holden et al. 2013).

The previously omitted reference is:

Horváth A, Dobay O, Kardos S, Ghidán Á, Tóth Á, Pászti J, Ungvári E, Horváth P, Nagy K, Zissman S, et al. 2012. Varying fitness cost associated with resistance to fluoroquinolones governs clonal dynamic of methicillin-resistant *Staphylococcus aureus*. *Eur J Clin Microbiol Infect Dis* **31**: 2029–2036.

The authors apologize for this omission, which has been corrected online.

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