Management and investigation of a Serratia marcescens outbreak in a neonatal unit in Switzerland - the role of hand hygiene and whole genome sequencing - R1, ARIC-D-17-00143

ZINGG, Walter, et al.

Abstract
Many outbreaks due to Serratia marcescens among neonates have been described in the literature but little is known about the role of whole genome sequencing in outbreak analysis and management.

Reference

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Correction to: Management and investigation of a *Serratia marcescens* outbreak in a neonatal unit in Switzerland – the role of hand hygiene and whole genome sequencing

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**Correction**

The original article [1] contained an error whereby an erroneous article ID was displayed at the end of the title; the original article has now been corrected to remove this ID.

In addition, this error was mistakenly carried forward by the production team that handled this article, and thus, was not the fault of the authors.

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