

Addendum

Whole-genome sequencing for neonatal intensive care unit outbreak investigations: Insights and lessons learned – ADDENDUM

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In the above article¹, WGS Lesson #3 was originally omitted; it reads:

WGS Lesson #3: WGS provides high resolution for strain typing in outbreak investigations and can provide support for transmission clusters that would have otherwise been miscategorized

Reference

Sansom, S., Logan, L., Green, S., Moore, N., & Hayden, M. (2021). Whole-genome sequencing for neonatal intensive care unit outbreak investigations: Insights and lessons learned. *Antimicrobial Stewardship & Healthcare Epidemiology*, 1(1), E2. doi: [10.1017/ash.2021.161](https://doi.org/10.1017/ash.2021.161)

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