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Whole-Genome Sequence of a Stenotrophomonas maltophilia Isolate from Tap Water in an Intensive Care Unit
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Abstract
Here, we present a 4,508,936-bp complete genome sequence of Stenotrophomonas maltophilia strain HW002Y, which was isolated from the tap water in an intensive care unit at Sultan Ahmad Shah Medical Centre at the International Islamic University of Malaysia (Kuantan, Pahang, Malaysia). Sequencing was performed using a Nanopore Flongle flow cell. © 2023 American Society for Microbiology. All rights reserved.

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