**Finished Genome Sequence of the Highly Multidrug-Resistant Human Urine Isolate *Citrobacter freundii* Strain SL151**

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*Citrobacter freundii* is a Gram-negative opportunistic pathogen that is increasingly being recognized as a causative agent of hospital-acquired urinary tract infections and an important reservoir of antimicrobial resistance determinants. In this report, we describe the finished genome sequence of *C. freundii* strain SL151, a highly multidrug-resistant human urine isolate.

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**Citation**


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