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Draft Genome Sequence for a Urinary Isolate of Nosocomiicoccus ampullae

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Draft Genome Sequence for a Urinary Isolate of Nosocomiicoccus ampullae

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A draft genome sequence for a urinary isolate of Nosocomiicoccus ampullae (UMB0853) was investigated. The size of the genome was 1,578,043 bp, with an observed G+C content of 36.1%. Annotation revealed 10 rRNA sequences, 40 tRNA genes, and 1,532 protein-coding sequences. Genome coverage was 727× and consisted of 32 contigs, with an N50 of 109,831 bp.

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