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Accessibility
High-Quality Draft Genome Sequence of Vagococcus lutrae Strain LBD1, Isolated from the Largemouth Bass Micropterus salmoides

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Vagococci are usually isolated from marine hosts and occasionally from endodontic infections. Using 16S rRNA gene comparison, the closest relatives are members of the genera Enterococcus and Carnobacterium. A draft sequence of Vagococcus lutrae was generated to clarify the relationship of Vagococcus to these and other related low-G+C Gram-positive bacteria.

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