

ERRATUM

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Erratum: The evolution and population structure of *Lactobacillus fermentum* from different naturally fermented products as determined by multilocus sequence typing (MLST)

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Erratum

The original version of this article [1] unfortunately contained a mistake in the section entitled “Nucleotide sequence accession numbers”. The accession numbers of two housekeeping genes (*clpX* and *pyrG*) used in this study were wrongly provided. KR078446-KR078647 should read KU512980 - KU513181 and KR079456 - KR079657 should read KU513182 - KU513383.

The correct text is provided below:

Nucleotide sequence accession numbers

The partial sequences of the 11 MLST loci used in this study have been deposited in the GenBank/EMBL databases under accession numbers KR078648-KR079455, KR079658-KR080061, KP224504-KP225109, KU512980 - KU513181 and KU513182 - KU513383.

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Reference

1. Dan T, Liu W, Song Y, Xu H, Menghe B, Zhang H and Zhihong Sun. *BMC Microbiology*. 2015;15:107. doi:10.1186/s12866-015-0447-z.

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