



# Correction for Kennan et al., “Genomic Evidence for a Globally Distributed, Bimodal Population in the Ovine Footrot Pathogen *Dichelobacter nodosus*”

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It has been drawn to our attention that there are errors in the list of SRA accession numbers in Table S6 of this article, specifically in the last column of this table. Accordingly, we have provided a corrected copy of Table S6. The table has been replaced online. The authors apologize for any inconvenience that this error may have caused to other researchers.

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