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Author Correction: Whole rumen metagenome sequencing allows classifying and predicting feed efficiency and intake levels in cattle

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Correction to: *Scientific Reports* <https://doi.org/10.1038/s41598-018-36673-w>, published online 09 January 2019

This Article contains a typographical error in the Methods section under subheading ‘Metagenome ensemble’ where,

“The annotation was performed with options –compliant –centre UoN –norrna –notrna –metagenome, and a bioproject was submitted to NCBI database with number PRJNA423102.”

should read:

“The annotation was performed with options –compliant –centre UoN –norrna –notrna –metagenome, and a bioproject was submitted to NCBI database with number PRJNA423103.”



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Published online: 13 February 2020