Comparison of the Fecal Microbiota of Healthy Horses a Throughput Sequencing of the V3-V5 Region of the 16S

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Citation Report

#	Article	IF	Citations
1	Toward an Understanding of Changes in Diversity Associated with Fecal Microbiome Transplantation Based on 16S rRNA Gene Deep Sequencing. MBio, 2012, 3, .	1.8	151
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