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Comparative diversity analysis of ruminal methanogens in Murrah buffaloes (*Bubalus bubalis*) in four states of North India

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20 **Abstract**

- We compared the community structure of methanogens in Murrah breed of buffaloes of four 21
- states of north India using 16S rRNA gene clone library method. The results revealed the 22
- dominance of methanogens related to Methanobrevibacter in three states, while 23
- Methanomicrobium- related methanogens were abundant in one state. 24

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