

Reviewer Report

Title: Draft genome of the gayal, Bos frontalis

Version: Original Submission **Date:** 6/16/2017

Reviewer name: Madhu Tantia

Reviewer Comments to Author:

Average exon in text is 3.27 where as in corresponding table it is 7.19 Reference for buffalo assembly is missing from references References need to be rechecked as per text There is a need to re-look into figure of 3.7gb as the genome size of Mithun

Level of Interest

Please indicate how interesting you found the manuscript: An article of importance in its field

Quality of Written English

Please indicate the quality of language in the manuscript: Needs some language corrections before being published

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