

CORRECTION

# Correction: A rapid multiplex PCR assay for presumptive species identification of rhinoceros horns and its implementation in Vietnam

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There are errors in [Table 1](#). The amplicon lengths for primers RID\_FWD/RID\_REV and Mac\_FWD/Mac\_REV do not include primer regions. Please see the corrected [Table 1](#) here.



## OPEN ACCESS

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**Table 1. Cytochrome b markers and their corresponding primers.**

Genetic marker for:	Primer name:	Primer sequence (5'-3'):	Annealing temperature (°C):	Amplicon length* (bp):	Reference:
<i>Diceros bicornis</i> (black rhino)	Rh_BR_FWD (forward)	AATCTGCCTAATCCTACAAATC	60	222	This study
	Rh_BR_REV (reverse)	GGTTTCTAGGAAGGTGTAGG			
<i>Ceratotherium simum</i> (white rhino)	Rh_WR_FWD (forward)	CCACTCATTTCATCGATCTGC	60	266	This study
	Rh_WR_REV (reverse)	TAATAGATACCGCGTCCTAC			
<i>Rhinoceros unicornis</i> (Indian rhino)	Rh_IR_FWD (forward)	TCTCACCCACTAGTTAAAATCA	60	310	This study
	Rh_IR_REV (reverse)	AGGAAGGTGTAAGATCCATAG			
All rhino species	RID_FWD (forward)	AACATCCGTAATCYCACCCA	55	274	[6]
	RID_REV (reverse)	GGCAGATRAARAATATGGATGCT			
All rhino species	Mac_FWD (forward)	CAYTATACACCAGACACAACAAC	55	226	This study
	Mac_REV (reverse)	TGAAYGCDGTGGCTATTAGRG			

\* Amplicon lengths are inclusive of primer regions.

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## Reference

1. Ewart KM, Frankham GJ, McEwing R, The DT, Hogg CJ, Wade C, et al. (2018) A rapid multiplex PCR assay for presumptive species identification of rhinoceros horns and its implementation in Vietnam. PLoS ONE 13(6): e0198565. <https://doi.org/10.1371/journal.pone.0198565> PMID: 29902212