Erratum to:

Lalremsanga, H. T., A. K. Bal, G. Vogel & L. Biakzuala: Molecular phylogenetic analyses of lesser known colubrid snakes reveal a new species of *Herpetoreas* (Squamata: Colubridae: Natricinae), and new insights into the systematics of *Gongylosoma scriptum* and its allies from northeastern India. – Salamandra, **58**: 101–115.

In our study (Lalremsanga et al. 2022), we have a minor misspecification regarding the uncorrected p-distance presented in the text. The correct values for the genetic divergence were, however, presented in the p-distance matrix in Table 3 (the table is presented here for reference). The sentences bearing the erratic values were embedded within the comparison section of the newly described *Herpetoreas murlen* (see Page 107). Therefore, the correct values (p-distance) are justified herein, and the erratic values are indicated in parenthesis.

From *H. murlen*, the estimated genetic divergence with *H. burbrinki* is 8.7% (not 8.3%); 14.5% with *H. pealii* (not 13.7%); 13% with *H. xenura* (not 13.4%); 11.1–11.5% with *H. platyceps* (not 13.3%); and the genetic divergence with respect to *H. sieboldii* (12.7–13.3%) was also erroneously presented because we do not sampled any genetic data for this species.

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Table 3. Uncorrected p-distances for members of the putative *Herpetoreas* clades. GenBank accession numbers are provided after the species names. Table adopted from Lalremsanga et al. (2022).

Species	1	2	3	4	5	6	7	8	9	10
1. *Herpetoreas murlen* sp. n. ON204025										
2. *Herpetoreas burbrinki* GQ281781	0.087									
3. *Herpetoreas platyceps* MT571587	0.111	0.109								
4. *Herpetoreas platyceps* KJ685690	0.115	0.123	0.043							
5. *Herpetoreas platyceps* MW111464	0.115	0.125	0.043	0.003						
6. *Herpetoreas xenura* MN993850	0.130	0.125	0.126	0.144	0.140					
7. *Herpetoreas xenura* MN993851	0.130	0.125	0.126	0.144	0.140	0.000				
8. *Herpetoreas pealii* MT571586	0.145	0.120	0.133	0.144	0.144	0.113	0.113			
9. *Herpetoreas tsper* MW111476	0.099	0.089	0.096	0.108	0.104	0.135	0.135	0.130		
10. *Hebius parallelus* MK201567	0.099	0.092	0.099	0.111	0.108	0.135	0.135	0.133	0.003	

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