Corrigendum

Corrigendum to “Some aspects of the venom proteome of the Colubridae snake Philodryas olfersii revealed from a Duvernoy’s (venom) gland transcriptome”


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An unfortunate error occurred during the preparation of Fig. 4B. In the second block of the alignment, the amino acid sequence segment “EALAGGTTALRDVEEQQQLVAEESG” should not be part of the AAC60341 protein, but part of the protein AAY47072 in the line below in the alignment. Note that the phylogenetic tree on Fig. 4C was constructed from the correct sequences.

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Fig. 4. Primary structure of the *P. olfersii* CNP precursor. (A) Schematic view of NP precursors in the different snake families and other vertebrates. (B) *P. olfersii* precursor (bold) is aligned to some representatives of such precursors, showing high conservation (shaded residues). (C) Bayesian inference of the phylogeny based upon the posterior probability distribution of the trees. MrBayes software version 3.1.1-p1 [27], with $10^6$ number of cycles for the Markov chain Monte Carlo algorithm, was used to the analysis of the proteins indicated by their accession numbers on the tree.