



Correction to: Analysis of the complete genomic sequence of a novel virus, areca palm necrotic spindle-spot virus, reveals the existence of a new genus in the family *Potyviridae*

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putative recognized peptides by HC-Pro1 and HC-Pro2 were “LFYG/V” and “FRVG/G”, respectively.

The original version of this article unfortunately contained a mistake in the ‘panel c’ of Fig. 1 and its corresponding description in text. In the genomic 5’-terminus, ANSSV was predicted to contain two HC-Pro proteins (i.e., HC-Pro1 and HC-Pro2) in tandem, rather than one single large HC-Pro. HC-Pro1 and HC-Pro2 flank the regions nt. 206-979 and 980-1927, respectively. Consistent catalytic activity sites (Gly¹⁵⁹, Cys¹⁶¹ and His²³³ for HC-Pro1) and (Gly⁴⁶¹, Cys⁴⁶³ and His⁵³⁵ for HC-Pro2) were predicted. The

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