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Due to an arithmetic error, an active site Serine residue at position 272 in the deduced amino acid sequence of the A subunit of the toxin described in this paper was incorrectly stated as being at position 271. This has resulted in incorrect terminology for the critical residue and for mutations introduced at this position, as listed below.

codon 271	*should be* codon 272	*on page 37*
S_{271}	*should be* S_{272}	*on pages 37 and 40*
Ser_{271}-Ala	*should be* Ser_{272}-Ala	*on page 45*
pK184*subA*$_{A271}$B	*should be* pK184*subA*$_{A272}$B	*throughout*
SubA$_{A271}$B	*should be* SubA$_{A272}$B	*in Figure 8 B*

The residue numbers of the terminal amino acids of the Ser catalytic domain (269 and 279) *should be* 270 and 280 *in Figure 2*.