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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
pK184subA _{A271} B	should be pK184subA _{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.