

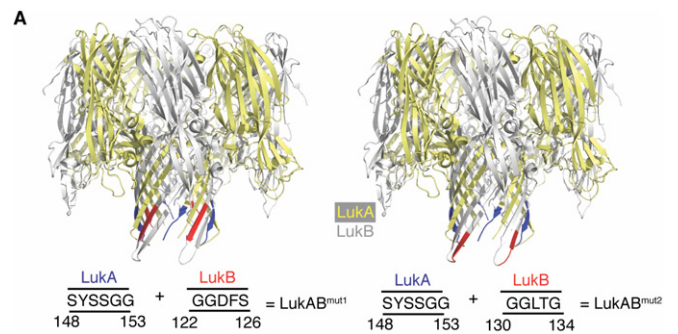
# Exploiting dominant-negative toxins to combat *Staphylococcus aureus* pathogenesis

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The authors of the above article regret a mistake in the labeling of the LukAB<sup>mut2</sup> mutant in Figure 4A and Table EV1. The correct mutated residues are 130–134 instead of residues 122–126 indicated in the current version of the article.

The correct Figure 4A is presented here, and the correct Table EV1 is now available online. The results and conclusions of the article remain unchanged.



**Figure 4.**