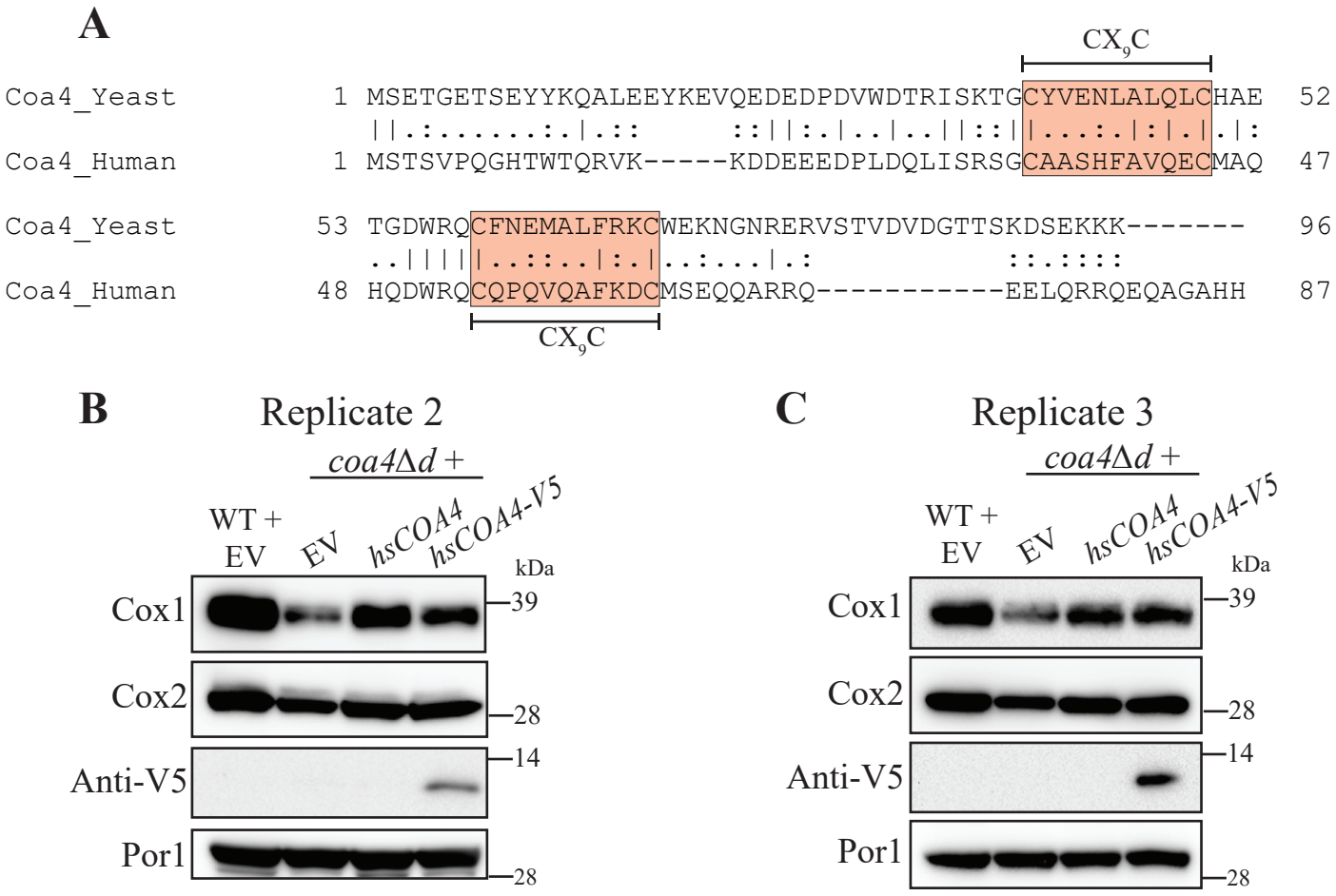

Supplementary Figure S7. (A) Pairwise sequence alignment of yeast and human Coa4 sequences. The yeast and human Coa4 protein sequences were obtained from UniProt and pairwise global sequence alignment was performed using EMBOSS needle program. The conserved twin CX_9C motif is highlighted in orange box. (B) Replicate 2 and (C) replicate 3 of figure 6D.