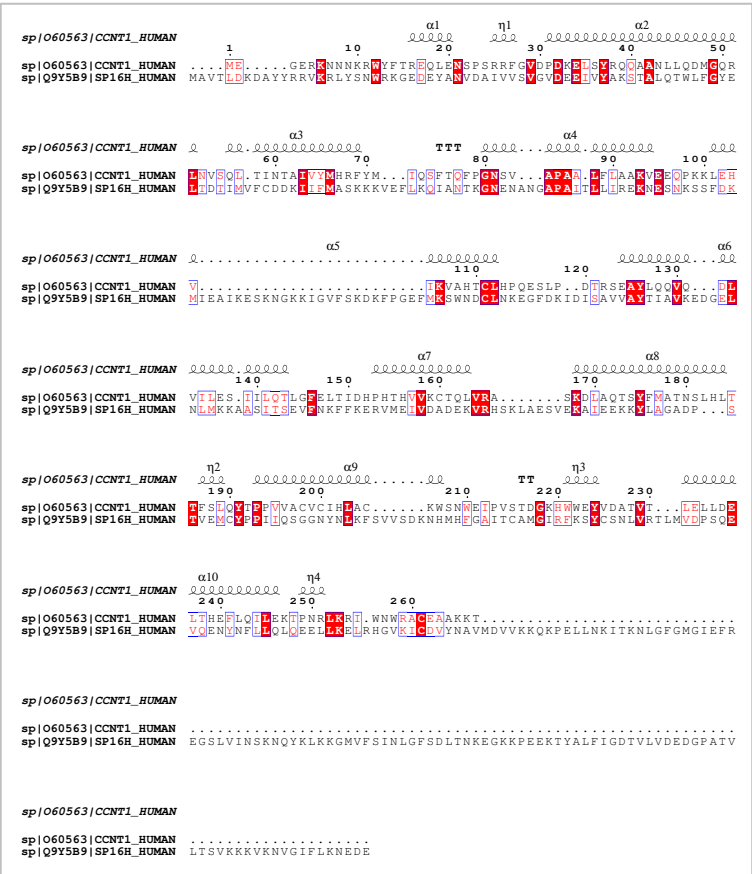
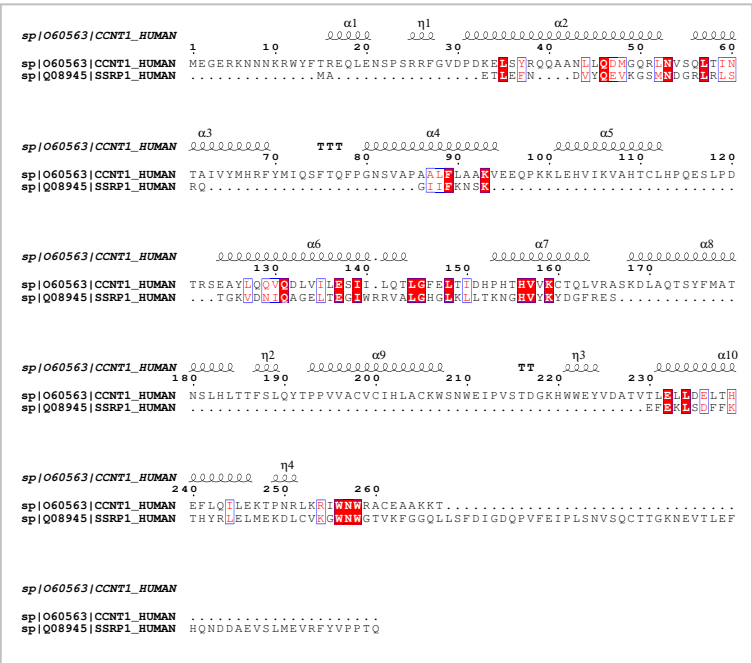


Supplementary Figure S4

A

**B**

Sequence alignment of Cyclin T1 residues 1-266 (Uniprot ID O60563) and N-terminal region of (A) SPT16 (Uniprot ID Q9Y5B9) or (B) SSRP1 (Uniprot ID Q08945) showing minimal sequence similarity between the two known ICP22-binding complexes.