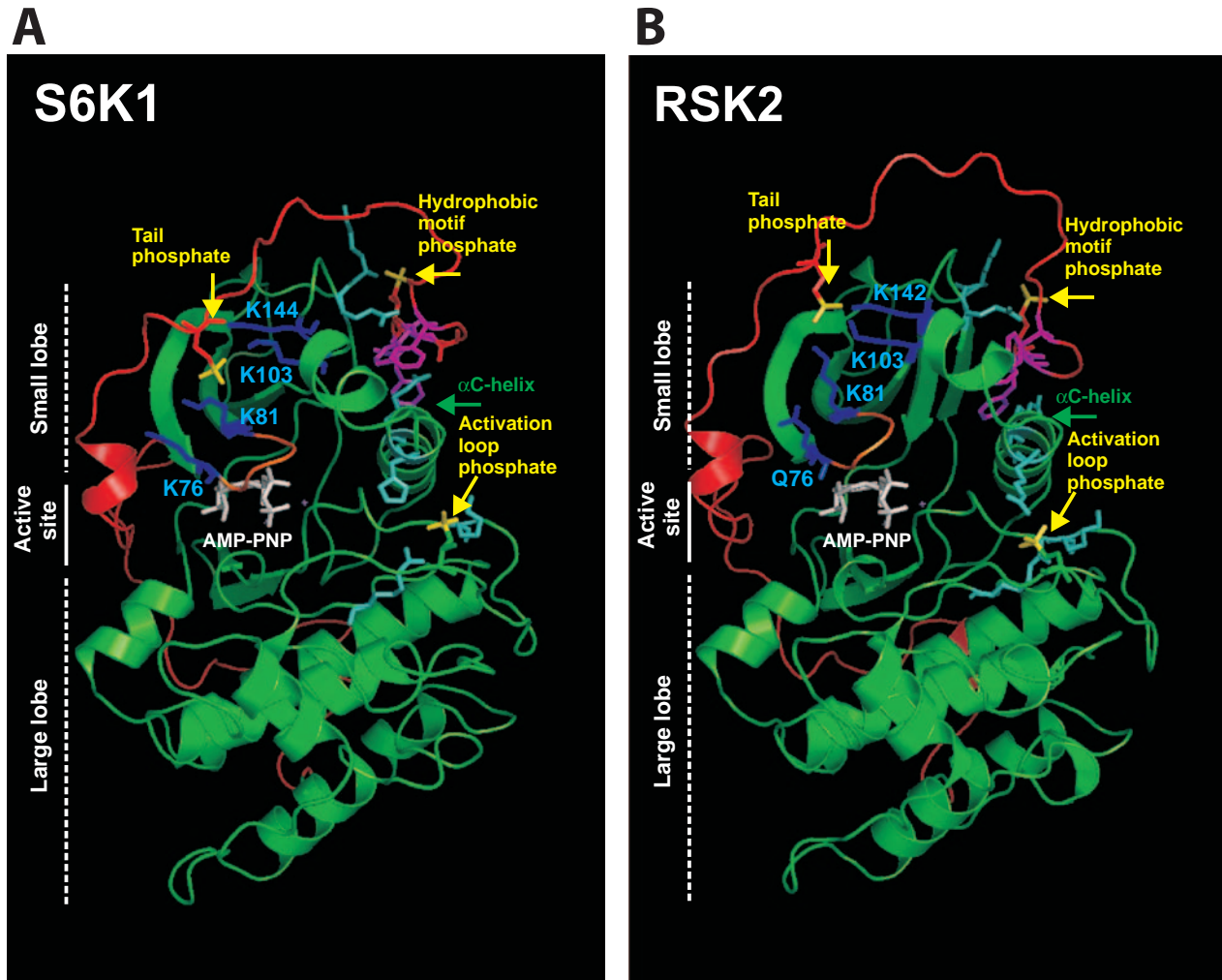


Supplementary Fig. 3



Supplementary Fig. 3. Molecular modelling of S6K1 and RSK2 suggests a homologous binding site for the tail phosphoSer/Thr within the small lobe of the kinase domain. The kinase domain and linker of S6K1 (A) and RSK2 (B) were modelled as described in Materials and Methods. Colour codes are as described in the legend to Fig. 1A.