



Supplemental Figure SF5. Conserved motif sequences of the Cdc28 cyclins from *C. neoformans* and *S. cerevisiae*. Identification of the 10 protein sequence motifs was performed using MEME Suite 4.10.1 using the following parameters: zero or one occurrence per sequence, 10 motifs, motif width set as between 6 and 50, site number of each motif set as between 2 and 600.