



S1 Figure. Conservation of cryptococcal nucleotide sugar transporters.

(A) Protein sequence alignment of Uxt1 and Uxt2 (CNAG_02036 and CNAG_03695) with conserved residues highlighted (black, identical residues; grey, conserved substitutions). (B) Phylogenetic relationships of *C. neoformans* (Cn) NSTs (including Uxt1 and Uxt2, in bold), and UDP-Xyl, UDP-Galf, and UDP-Arap transporters from other organisms (*Hs*, *Homo sapiens*; *Af*, *Aspergillus fumigatus*; *At*, *Arabidopsis thaliana*) using MUSCLE, PhyML, and TreeDyn software (see Materials and Methods). Branch lengths are drawn to scale.