Supplementary Figure S1. Alignment of amino acid sequences of *A. fumigatus* zinc importers, ZrfA, ZrfB, ZrfC and ZrfD.
Figure S2. Alignment of *C. albicans* Pra1 and *A. fumigatus* AspF2.

For both figures, sequences were aligned using ClustalW (www.ebi.ac.uk/Tools/msa/clustalw2/) and rendered using ESPript [1].