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Candida_glabrata	.....
Rhizopus_oryzae	.....
Mucor_circinelloides	795 ALKDIKIKIYAPPERKYSTWIGGSILASLSTFKKMVSAEEYQEDPDIHKKTF
Phycomyces_blakesleanus	.....
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**Supplementary Figure 1. The C-terminal modulatory region of Mid1 is not conserved among fungi.** The multiple alignment of the primary structure of Mid1 from 28 species of fungi. The sequences were aligned with ClustalW multiple alignment program (22) and viewed with Jalview (43). Black filled triangles indicate 11 conserved cysteine residues among fungi and black filled ovals indicate two conserved CP (cysteine and proline) dipeptide motif. The 24 amino acid C-terminal modulatory domain is indicated by the grey box. Shaded regions correspond to amino acid similarity in Mid1 among fungi.