Figure S2 RIC8 homologs in other fungi. ClustalW (http://www.ch.embnet.org/software/ClustalW.html) was used to align RIC8 sequences from *Neurospora crassa* (Nc; NCU02788.3), *Podospora anserina* (Pa; Pa_1_3900), *Magnaporthe grisea* (Mg; MGG_14008), *Fusarium graminearum* (Fg; FGSG_01511), *Sclerotinia sclerotiorum* (Ss; SS1G_02787.1), *Aspergillus nidulans* (An; AN1661.3), *Histoplasma capsulatum* (Hc; HCAG_03335.1), *Botrytis cinerea* (Bc; BC1G_02317.1), *Schizosaccharomyces japonicus* (Sj; SJAG_02027), *Yarrowia lipolytica* (Yl; ACCESSION# XP_505953). Shading indicates identical (black) and similar (grey) amino acid residues (http://www.ch.embnet.org/software/BOX_form.html).