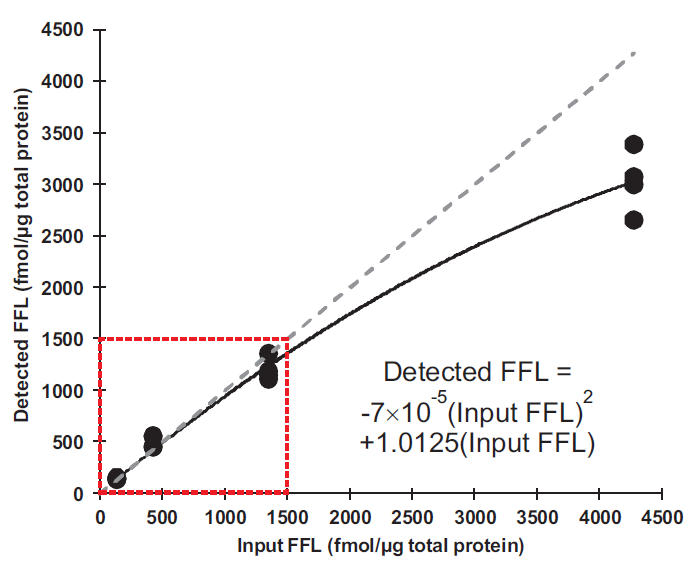
Supplementary Materials: Proteomic Analysis   
of Pathogenic Fungi Reveals Highly Expressed Conserved Cell Wall Proteins

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**Figure S1.** Dynamic range and linearity of MSE (Mass Spectrometry—Elevated Collision Energy) for label-free protein quantification in the presence of *Aspergillus fumigatus* cell extract. The dashed red box indicates the range of most quantified proteins, and the grey dashed line shows an ideal 1:1 ratio. Data from 16 independent LC-MSE analyses. FFL—recombinant Firefly Luciferase.