Figure S7

Gene count per OrthoMCL family for each species.

Genes were clustered into families using OrthoMCL for *S. sclerotiorum*, both strains of *B. cinerea*, and 7 other fungal genomes. The x-axis bins measure the number of genes per family for each species, where orphan genes not in families are in the 0 bin, genes in single copy in a family are in the 1 bin, and paralogous genes of 2 or more in a family are in the higher bins. The y-axis varies between the subplots, and counts the total number of genes in each bin per species.