



Supplementary Figure S4. GO classification of common DEGs in response to *B. cinerea* and other stress groups. GO of *BUGs* and *BDGs* also affected by biotic (A), abiotic (B) and hormonal (C) stresses based on biological process, molecular function and cellular components. Gene identifications of 1,554 *BUGs* and 1,206 *BDGs* entered for this analysis. Error bars are SD. GO categories significantly over- or under-represented at $P < 0.05$ are shown. Normalized frequency of genes to the number of genes on the microarray chip was determined as previously described³⁰.