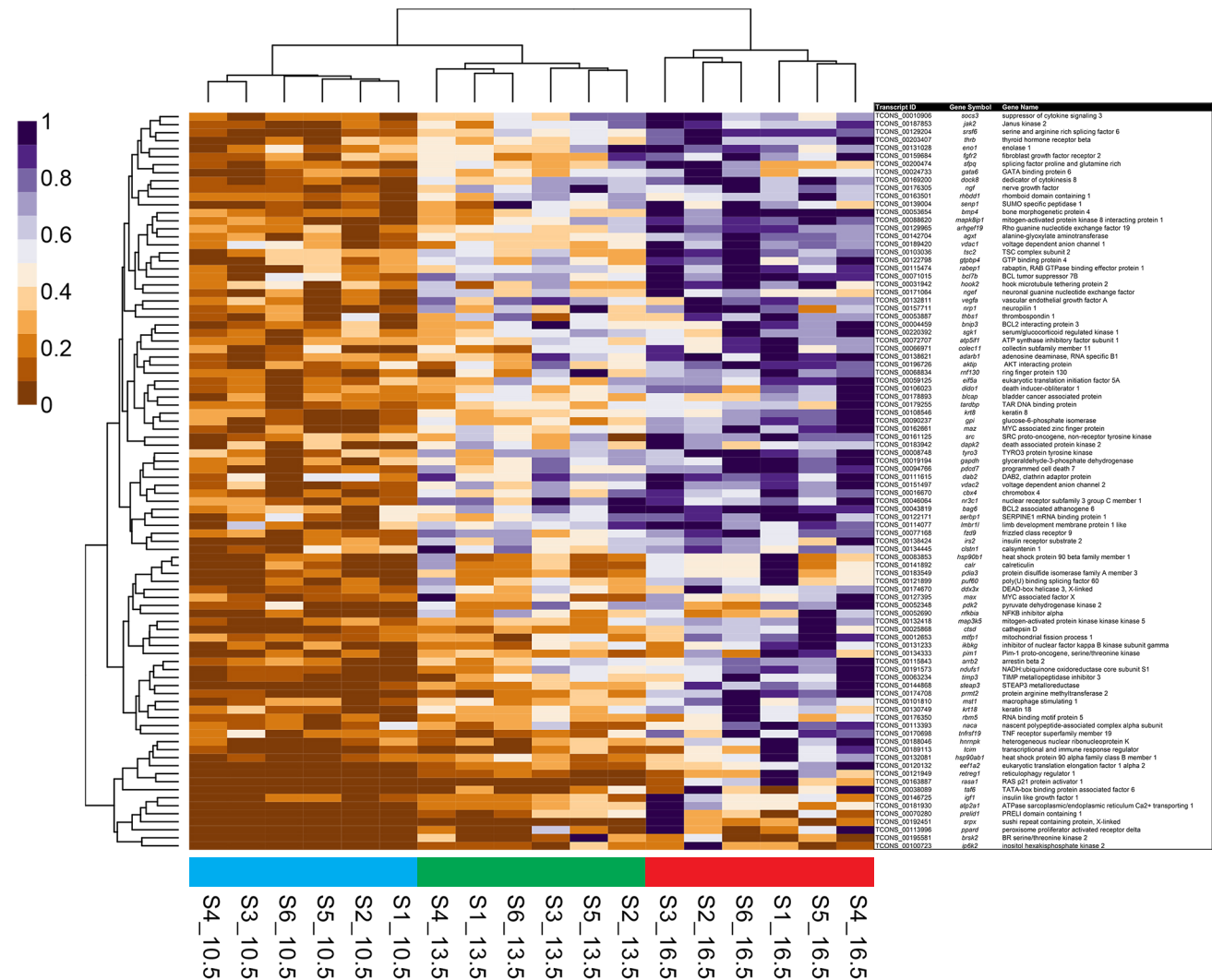




Supplementary Figure S1. Heatmap and hierarchal clustering of non-redundant differentially expressed transcripts (92 DETs) found in comparison of the 10.5 °C and 16.5 °C groups associated with GO:0006915 - Apoptotic Process. RNA samples used in sequencing are labeled on the base of the figure, with standardized fragments per kilobase of transcript per million mapped reads (FPKM) values for each DET clustered along the vertical axis.