



Appendix S12. Heat map of expression differences for the 10 genes that were identified as having a significant genotype by treatment interaction for all pairwise comparisons to SW. The value of each cell represents the log₂ fold-change in gene expression before and after cold acclimation (counts per million reads). Yellow-red color represents genes that are more highly expressed after cold acclimation, while black-purple color represents genes that are more highly expressed before cold acclimation. Plot was generated using the heatmap.2 function in the R package plots (Warnes et al., 2019).