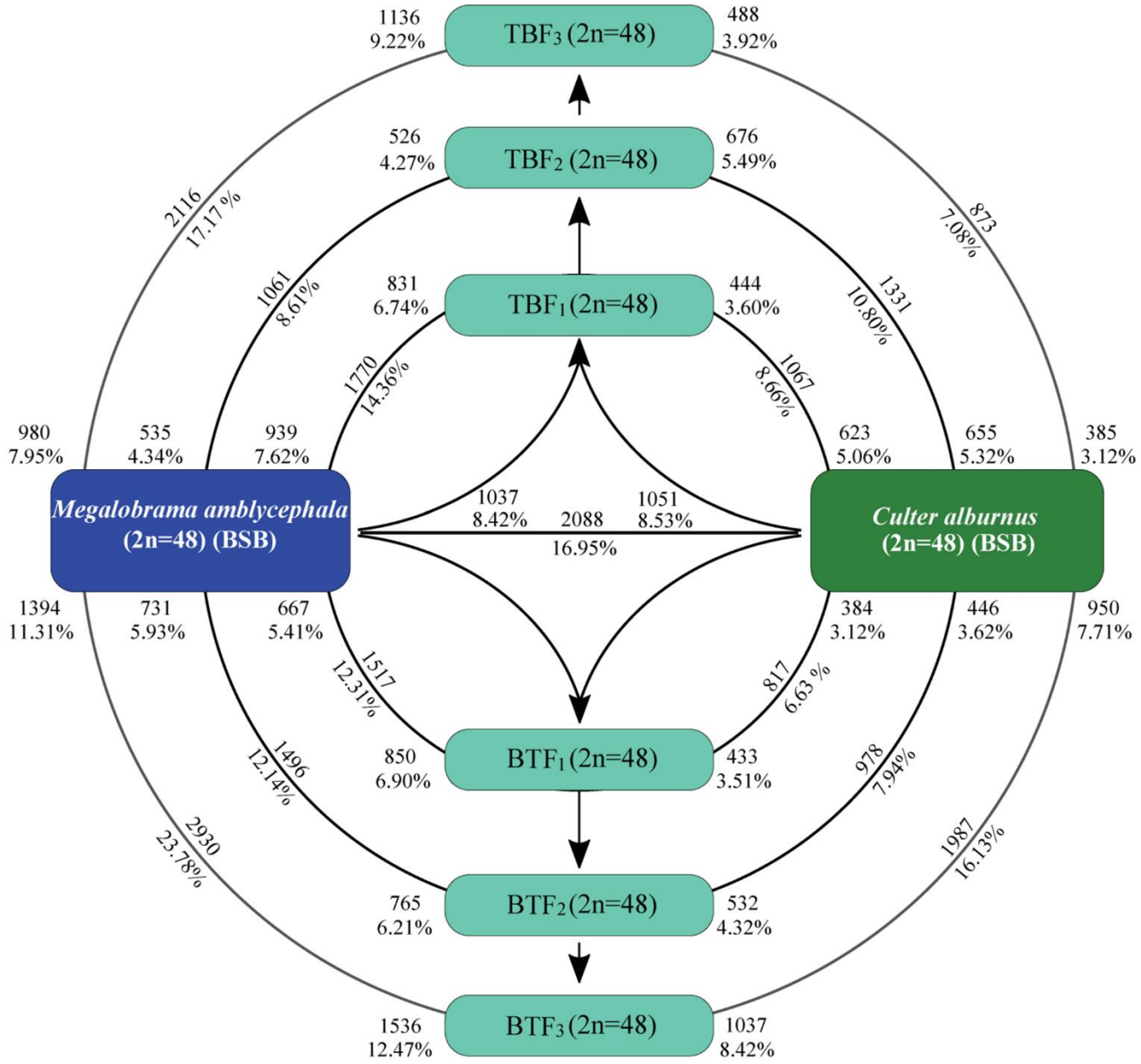

Supplemental Fig. S24. Differentially expressed genes between the two parents and their reciprocal cross hybrids. The number indicates the total number of genes identified as differentially expressed in liver tissue in each comparison based on RNA-seq. Also shown for each comparison is the proportion of the total number of differentially expressed genes with upregulation.