



Supplementary Figure S1: Hierarchical clustering heatmap depicting overall results of the FPKM clustering of differentially expressing genes (DEGs) of WBPH (using $\log_2(\text{FPKM}+1)$ values) while feeding on different RILs (TN1, TR145 and TR152). The scaffold IDs are provided on the right. The color scale represents an increase in gene expression from blue (-1) to red (1).