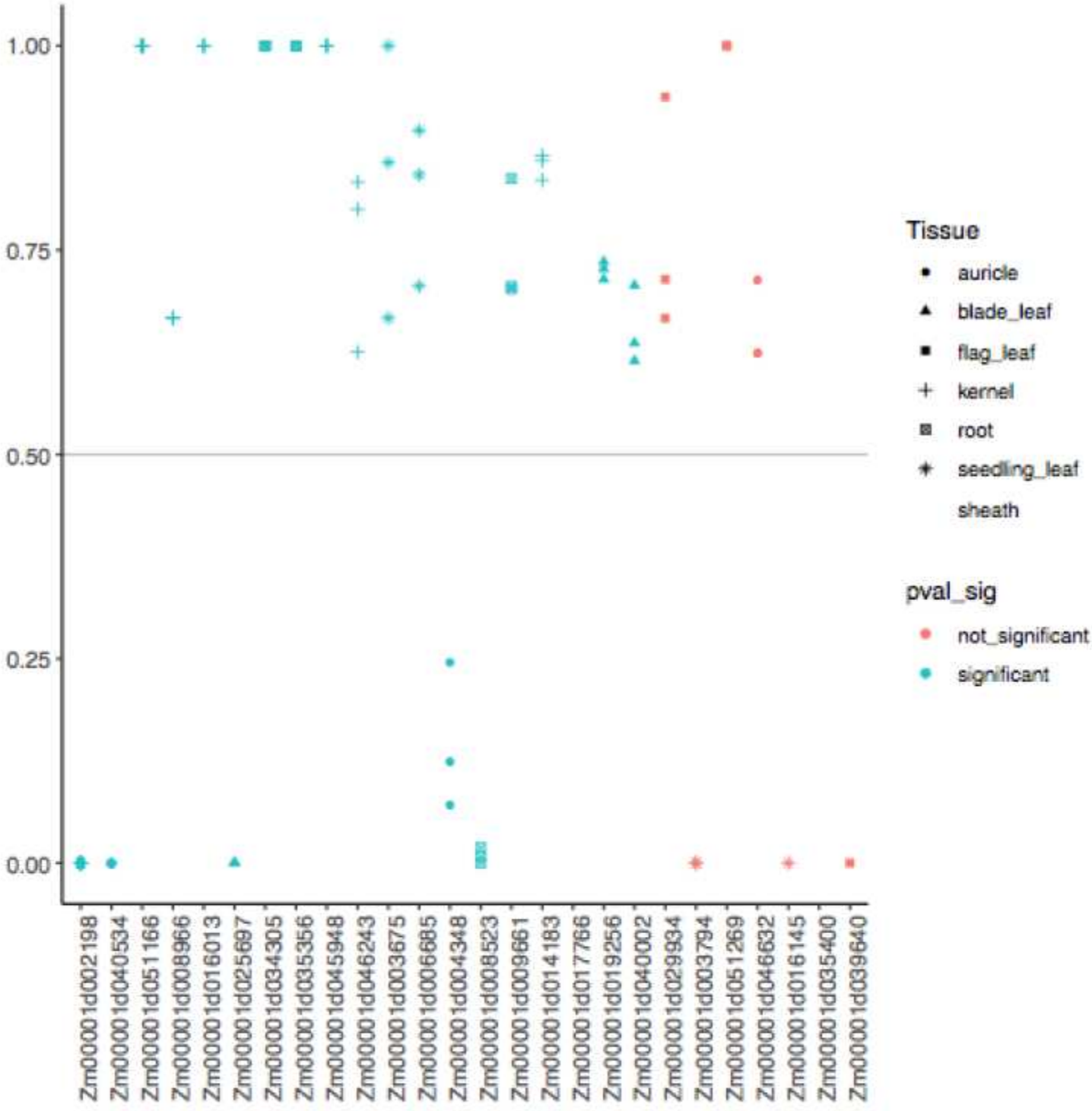

Figure S10: Allele specific expression of significant TE-Gene pairs. A subset of TE-ACRs with significant gene expression changes when present in B73 and absent in Mo17 and contain SNPs within the coding region (N=26) were assessed. Allele specific expression data from Zhou et al., 2019 was used to plot the ratio of the B73 to Mo17 alleles for the tissue determined as significant in the TE-ACR analysis. Tissue type is shown by shape and significant genes (based on a chi-square p-value < 0.1) is shown in blue.