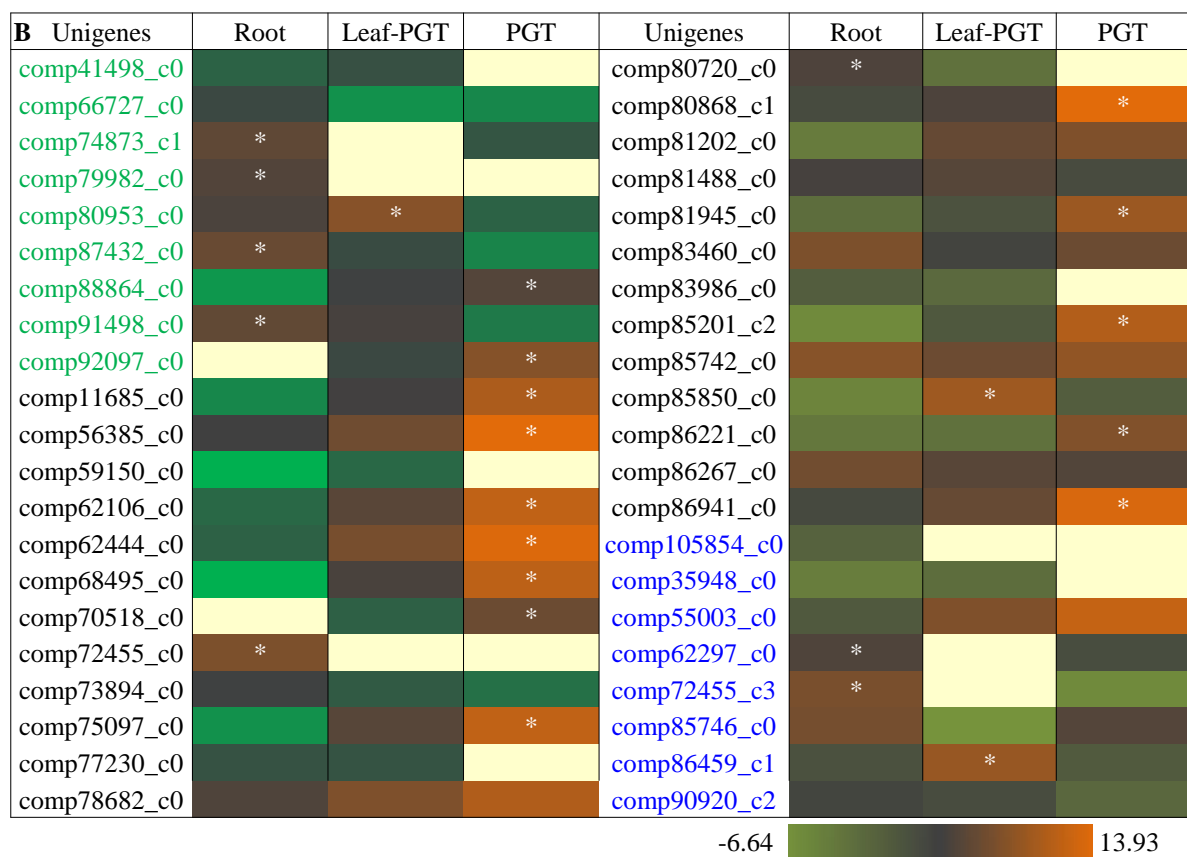
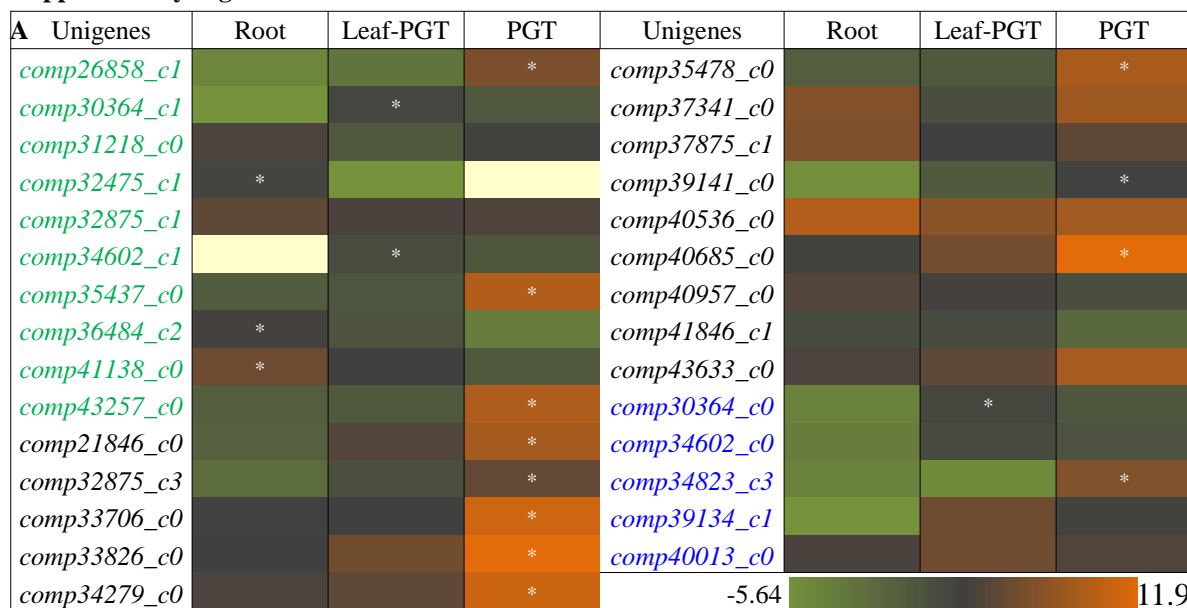


Supplementary Figure S4



Supplementary Figure S4. Expression profiling of TPSs in mint and basil based on RNA-Seq analysis. (A) Heat map showing the expression patterns of mint TPSs among three different tissues including root, leaf-PGT and PGT. (B) Expression patterns of basil TPSs in three different tissues including root, leaf-PGT and PGT. In (A) and (B), the stars “*” indicated the tissue-preferred genes. The yellow cells indicate no expression of this gene in the tissue (FPKM=0). The green and blue locus names indicated TPSs encoding either PF01397 or PF03936 domain only, respectively. The black locus names indicated that the TPSs encode both domains.