

Figure S3 Tissue specific expression of causative gene candidates. The gene expression was retrieved from the genome-wide tissue transcription by Sekhon et al. (Sekhon et al. 2011). Cob, immature cob at V18; Pericarp, pericarp at 18DAP; Root_VE, primary root at VE; Silks, silks at R1; Leaf_base, leaf at base of stage-2; Leaf_V9, eleventh leaf at V9.

Sekhon, R.S., H. Lin, K.L. Childs, C.N. Hansey, C.R. Buell et al., 2011 Genome-wide atlas of transcription during maize development. Plant J. 66: 553-563.