



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.