

Supplemental Figure 3. Histogram of RNAseq data describing mRNA accumulation for *Lgn* (GRMZM2G134382) in multiple tissue types for both B73 and Mo17. RNAseq data was generated using the qTeller tool (<http://qteller.com/>). Tissue types are as follows: root, shoot, tassel and shoot apex. The HTML link for regenerating this analysis at the qTeller website is:

http://qteller.com/NAM/bar_chart.php?name=GRMZM2G134382&info=B73_root%7CMo17_root%7CMo17_sam_apex%7CB73_sam_apex%7CMo17_tassel%7CB73_tassel%7CMo17_shoot%7CB73_shoot

