

**Supplementary Figure 11. Phenotypic variation explained by genic and non-genic TASs.** Results are based on the maize Working Gene Set. (A) genic TASs versus all other TASs; (B) genic region plus promoter 1kb versus all other TASs, and (C) genic region plus promoter 5kb versus all other TASs. Phenotypic variation explained by all QTLs is shown for comparison.

