



Supplementary Figure 13. Probable QTL genotype at the signal peak from all progenies for diameter QTLs of diploid rose populations *qDiam.TX2WOB-LG1* (A) and *qDiam.TX2WOB-LG2.2* (B) for TX2WOB, and *qDiam.TX2WSE-LG2* (C) for TX2WSE.

Means not connected by the same letter are significantly different ($P < 0.05$) within each linkage group.