



Supplementary Figure 14. Probable QTL genotype at the signal peak from all progenies for flower dry weight and fresh weight QTLs of diploid rose populations *qDWT.TX2WOB-LG1*(A) and *qDWT.TX2WOB-LG3* (B) for TX2WOB and *qFWT.TX2WSE-LG3* (C) for TX2WSE.

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