



Supplementary Figure 16. Probable QTL genotype at the signal peak from all progenies for number of petaloid QTLs of diploid rose populations $qPD.TX2WOB-LG3$ (A) and $qPD.TX2WSE-LG3$ (B) for TX2WOB and TX2WSE, respectively.

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