

Fig. S4 The distribution of differentially methylated genes in *Populus tomentosa* and *Populus simonii*. (a) Percentage distribution of differentially methylated genes in genomic features in *Populus tomentosa* and *Populus simonii*. (b) Overlaps between lncRNAs-target PCGs and differentially methylated genes in *P. tomentosa* and *P. simonii*. (c) Overlaps between differentially methylated lncRNAs-target PCGs and differentially expressed genes in *P. tomentosa* and *P. simonii*.

