



Figure S9 Distributions of gene categories, linkage disequilibrium, nucleotide diversity and nucleotide divergence across the consensus linkage map of loblolly pine based on sliding windows (5 cM size in steps of 2 cM). (A) Stacked bar plot of gene categories across the consensus linkage map. (B) Intragenic linkage disequilibrium, as assessed using Kelly's Z_{ns} statistic, across the consensus linkage map. (C) Nucleotide diversity (θ_{π} = red, θ_w = blue) and nucleotide divergence with respect to *P. radiata* (green) across the consensus linkage map. The number of amplicons where nucleotide divergence relative to *P. lambertiana* was defined was too small to plot across linkage groups.