

Table S4 Comparisons of the MergeMap and LPmerge *P. taeda* consensus genetic maps.

	MergeMap	LPmerge
Number of markers mapped	3856	3856
Number of unique marker positions	3810	2959
Average marker spacing (cM)	0.599	0.469
Total map length (cM)	2305.42	1802.59
RMSE with input maps averaged across LGs	0.97	1.34