

Table S6 Nucleotide diversity (per site) across linkage groups of loblolly pine (*Pinus taeda* L.). Values are weighted averages (see Materials and Methods).

LG	<i>l</i>	<i>l</i>	<i>l</i>	<i>n</i>	<i>n</i>	<i>S</i>	<i>S</i>	<i>S</i>	<i>S</i>	<i>S</i>	θ_{π}	θ_{π}	θ_{π}	θ_{π}	θ_w	θ_w	θ_w	θ_w
	All	<i>n</i> >1	Ann	<i>n</i> >1	Ann	All	Ann	NS	SY	NC	All	NS	SY	NC	All	NS	SY	NC
1	72	72	36	13.0	13.6	3.6	3.5	1.4	1.1	1.4	0.00323	0.00167	0.00550	0.00194	0.00320	0.00169	0.00478	0.00192
2	115	111	55	12.5	13.2	5.6	4.4	1.4	1.6	1.8	0.00467	0.00167	0.00792	0.00269	0.00486	0.00192	0.00708	0.00282
3	116	111	55	12.2	12.0	4.4	4.0	0.9	1.7	1.7	0.00383	0.00116	0.00892	0.00271	0.00381	0.00137	0.00855	0.00303
4	94	94	58	12.8	12.7	5.1	4.8	1.1	1.7	2.2	0.00421	0.00180	0.00839	0.00357	0.00418	0.00164	0.00784	0.00358
5	117	113	64	12.7	12.8	5.4	5.2	1.4	1.8	2.4	0.00390	0.00150	0.00843	0.00427	0.00430	0.00171	0.00895	0.00447
6	118	114	59	13.2	13.5	4.8	4.6	1.5	1.4	2.5	0.00394	0.00165	0.00710	0.00246	0.00431	0.00201	0.00771	0.00299
7	113	108	57	12.5	12.9	4.7	4.5	1.3	1.6	2.2	0.00369	0.00127	0.00793	0.00312	0.00390	0.00161	0.00749	0.00345
8	136	131	70	12.7	13.4	4.5	4.5	1.1	1.8	2.4	0.00389	0.00154	0.00865	0.00270	0.00394	0.00148	0.00929	0.00272
9	99	97	54	12.7	12.6	5.3	4.9	1.0	2.0	2.4	0.00399	0.00126	0.00802	0.00259	0.00422	0.00129	0.01039	0.00281
10	114	112	59	13.0	12.6	5.1	5.1	1.5	1.8	2.5	0.00458	0.00205	0.00859	0.00411	0.00473	0.00215	0.00790	0.00400
11	101	99	49	12.6	13.4	5.3	6.1	2.2	2.3	2.9	0.00416	0.00241	0.00910	0.00356	0.00434	0.00259	0.00998	0.00365
12	111	106	57	12.6	13.5	4.7	4.3	1.3	1.8	1.9	0.00414	0.00143	0.00838	0.00283	0.00422	0.00166	0.00847	0.00259

Abbreviations: All, all aligned sites; Ann, annotated; *l*, number of loci; LG, linkage group; *n*, sample size; NC, noncoding; NS, nonsynonymous; *S*, segregating sites; SY, synonymous; θ_{π} , nucleotide diversity based on the average number of pairwise differences; θ_w , nucleotide diversity based on the number of segregating sites following Watterson (1975).