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Effect of seedling stock on the early stand development and physiology of improved loblolly pine (*Pinus taeda* L.) seedlings

Shakuntala Sharma ⁽¹⁾, Joshua P Adams ⁽²⁾, Jamie L Schuler ⁽³⁾, Robert L Ficklin ⁽⁴⁾, Don C Bragg ⁽⁵⁾ This study assessed the effects of spacing and genotype on the growth and physiology of improved loblolly pine (*Pinus taeda* L.) seedlings from three distinct genotypes planted in Drew County, Arkansas (USA). Genotype had a significant effect on survival and height. Clone CF Var 1 showed greater height and survival compared to other seedlings. Genotype had significant effects on uniformity in height both years and ground line diameter (GLD) first year. However, genotype had no significant effects on leaf water potential and coefficient variation of leaf water potential. These growth and physiology should be further studied to assess potential genetic differences among seedlings and to determine if they can be identified early for improved growth at later ages.

Keywords: Loblolly Pine, Genotype, Leaf Water Potential, Coefficient of Variation

Introduction

Loblolly pine (Pinus taeda L.) is the native and most commercially important pine species in the southern United States and has been intensively and extensively managed for pulp and timber throughout the southeastern US (Fox et al. 2007). As consumer demands for wood timber products increase, greater production will be required to ensure supply meets demand (Prestemon & Abt 2002). Given that this increased production must come from a gradually declining timber base, intensive management that combines the best genetically improved planting stock and best silvicultural practices likely represent the most effective option to meet these future demands (Fox et al. 2007, Zhao et al. 2011, Aspinwall et al. 2012, Subedi et al. 2012).

Deployment of improved varieties could result in greater stand-level uniformity and enhanced productivity potentially aiding in the sustainability of southern forests to meet market demands (Jansson & Li 2004). An increase of 15 to 20% in volume and improvements to stem quality and disease

resistance have been reported in first generation loblolly and slash (Pinus elliottii Mill.) pine (Hodge et al. 1989, Talbert et al. 1985). This increase is even more pronounced when comparing the mean annual increment (MAI) of naturally regenerated stands (2 to 3 m³ ha¹ yr¹) to high intensity plantations with MAI of 21 to 28 m³ ha⁻¹ yr1 (Fox et al. 2007). Large scale planting of genetically improved stock loblolly pine will inevitably have a significant impact on the future wood supply and forest management practices across the southern US (Li et al. 1999b). Across the southeastern US improved genetics and intensive culture such as N and P fertilization and completion control have significantly increased the productivity of loblolly pine (Albaugh et al. 1998, McKeand et al. 2006, Fox et al. 2007, Subedi et al. 2012).

Genetically improved loblolly pine plantations, with their greater physiological uniformity, could result in more resource capture and more stand level productivity during the early stages of stand development (Bettinger et al. 2009). Few studies have compared the uniformity in growth and physiology of improved loblolly pine (Aspinwall et al. 2011a, 2011b). Therefore, the goal of this study was to compare the growth, uniformity in growth and leaf water potential among three loblolly pine genotype. Our hypothesis was that less genetically diverse genotypes (i.e., clones) would have more uniform growth and physiological traits than a more diverse genotype.

Material and methods

Site and loblolly pine genotype descriptions

The study area is located in Drew County, Arkansas (USA), on the University of Arkansas-Monticello (UAM) Teaching and Research School Forest (Latitude: 33° 37′ 1″ N, Longitude: 91° 43′ 9″ W). Based on NOAA weather station data, mean annual precipitation is 135.9 cm, with an average January temperature of 6.3 °C, and an average July temperature of 27.7 °C (Larance et al. 1976, NOAA 2013). The study area is characterized by predominant Calloway silt loams (fine-silty, mixed, active, thermic aquic Fraglossudalfs) with strongly acidic soil with level to gentle slope with an estimated 50-year loblolly pine site index 25 m (Larance et al. 1976).

The study site was a mature (approximately 55-year-old) pine-hardwood stand prior to destruction *via* tornado and subsequent clearing in 2010. Three families of loblolly pine seedlings were hand-planted in January 2012: an Arkansas-origin improved, open-pollinated family and two Atlantic Coastal Plain-origin full-sibling cloned families. Banded herbicide was applied post planting in late winter 2012 and contained 2% solution of glyphosate and Oust® XP 0.14 kg ha¹. An additional applica-

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tion was made during late winter of 2013. No fertilization treatment was done.

The three types of seedlings included a half-sibling bare-root seedlings (hereafter, "half-sib") were Arkansas Forestry Commission 3-star loblolly pine produced from seed sources of Arkansas, and selected by the Western Gulf Tree Improvement Cooperative. The half-sib seedlings are reported to have a 41-51% genetic gain over woodsrun stock (AFC 2014). Also included were, two full-sib clones planted as containerized seedlings and developed by CellFor (currently owned by ArborGen). The first clone, CellFor Variety 1 (CF Var 1) was products of Carolina parents (father tree South Carolina and mother tree North Carolina). These clones are advertised as having fast growth rate, high resistant to fusiform rust and pitch canker, stem straightness, and medium to wide crown (CellFor 20). CellFor Variety 2 (CF Var 2) is the progeny of two South Carolina parents and has exceptional tree form with small branches, narrow crown, stem straightness, fast growth rate, and high resistance to fusiform rust.

Experimental design

The experimental design was a split plot design. Wide (3.05 m \times 4.27 m = 13 m²) and narrow (3.05 m \times 3.05 m = 9.3 m²) spacing treatments were randomly assigned to the whole plot units. Genotypes (half-sib, CF Var 1, and CF Var 2) were randomly assigned to the subplots resulting into six treatment combinations. Each combination was replicated 3 times, for a total of 18 genotype by spacing with 140 trees per plot. Within one of the subplots in each of the spacing strips, the subplot was further split into a mixture planting that included a clone (35 tree) and half-sib planting (35 tree) mixture in alternate row. Mixed plots were also segregated by stock and considered subplots in this study, given no treeto-tree competition was occurring at this early stage. A total 3360 trees comprised the study which covered 0.3 ha in area.

Data collection

Ground line diameter (GLD) to the nearest mm and height (HT) to the nearest half cm were measured for all seedlings at the end of first and second growing season. Survival percentage at the end of each growing season also was determined dur-

ing these sampling periods with a census of each subplot. Trees were counted as alive if green needles were present (n=3360).

Protocols were developed based on available measuring devices, labor and limited time frame (an hour) for leaf water potential (MPa) data collection to ensure maximal environmental homogeneity. Needles were collected from six randomly selected trees per plot (n=180). Needles along the stem between first flush and first branch were excised, placed into plastic bags, and stored in a cooler until they were analyzed in the lab. Leaf water potential was measured in one randomly selected needle per tree with a pressure bomb (PMS Instrument Company, Albany, OR, USA).

Leaf water potential were measured every month, at mid-month, throughout the growing season (i.e., March to September 2013) in similar weather conditions, i.e., full sun. To compare different genotypes for leaf water potential, measurements were done within an hour and under consistent weather in terms of wind speed, temperature, solar radiation etc. when the plants were at a relatively stable hydraulic state; at dawn plants were under minimal water deficit while at midday they were at peak stress (Blum 2011). PMS Instrument Company has given ranges of values to determine the stress level of loblolly pine. Predawn/midday plant moisture stress (PMS) value ranges from 0.5 to 0.8 MPa reflect no limitation for growth, and under this condition plants have adequate water supply to maintain the maximum shoot growth; predawn to mid-day PMS value of 1.0 to 1.2 MPa reflects the slight to moderate shoot growth reductions; such stress limits phloem transport, leaf expansion, and diameter growth. Whereas PMS value of 1.2 to 1.4 MPa during sunrise to midday can cause stomatal closure (PMS 2014). These positive pressures reflect the reciprocal tension (negative energy/pressure) at which water is held in the leaf.

Data analysis and model development

Effects of spacing, genotype, and their interaction on mean height, GLD, survival, height growth, GLD growth, and leaf water potential were analyzed using a generalized linear model approach (PROC GLM in SAS® version 9.2) with the block as a ran-

Tab. 1 - ANOVA table of arcsine transformed loblolly pine second year survival and change in mortality from first-year to second-year per spacing-by-genotype sub-plots. (*): Denotes significance at α = 0.05.

Effects	df	Survival (age 2)	Mortality (age 1 to age 2)	
		P > F	P > F	
Spacing	1	0.23	0.84	
Block	1	0.19	0.03*	
Block × Spacing	1	0.63	0.37	
Genotype	2	<0.01*	0.28	
Block × Genotype	2	0.39	0.96	
Genotype × Spacing	2	0.81	0.95	

dom effect and both genotype and spacing as fixed effects. The study design was a blocked split-plot design, in which the main effect (spacing) was split into the genotypes (subplots). The statistical model used was (eqn. 1):

$$Y_{ijk} = \mu + \alpha_i + \beta_j + \gamma_k + (\alpha \beta)_{ij} + (\alpha \gamma)_{jk} + (\beta \gamma)_{ik} + e_{ijk}$$

where Y_{ijk} represents response variable at i^{th} block, j^{th} spacing, and k^{th} genotype, μ represents the overall mean effect, α_i was the effect of the i^{th} block, β_i was the fixed effect of the j^{th} main plot spacing, γ_k was the fixed effect of the k^{th} subplot genotype, $\alpha\beta$, $\alpha\gamma$, and $\beta\gamma$ were the interaction terms, and e_{iik} was the error term of this model. Survival was expressed as percent survival per sub-plot and transformed using the arcsine function prior to performing ANOVA. Changes in height growth and GLD growth were calculated by subtracting the first year from the second year measurements. Individual trees within a plot were averaged to obtain a plot mean and all the analysis was performed at the sub-plot level. In order to determine the uniformity, the coefficient of variation (CV) for GLD, height, leaf water potential was estimated for each genotype growing within each spacing for each growing season. ANOVA was used to test for differences in the CV among genotype and spacing. When the ANOVA identified significant genotype or interaction treatment effects at $\alpha = 0.05$, a least squares means approach was used to compare the genotypes for significant differences. Assumptions of all analyses were checked before and after analysis.

Results

Effect of genetics, spacing, and their interaction on survival, growth and uniformity

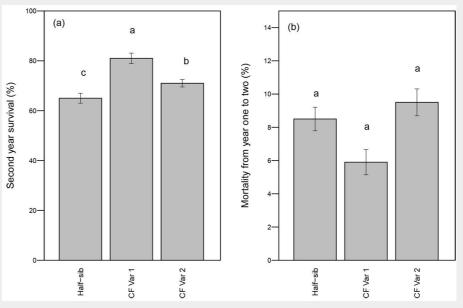
Survival after two growing seasons was significantly affected by genotype (p <0.01 - Tab. 1). CF Var 1 had significantly greater survival rate (81%) than either CF Var 2 (71%) or half-sib seedlings (65% - Fig. 1a). While CF Var 2 had experienced greater mortality from year one to year two relative to the half-sib and CF Var 1 seedlings (Fig. 1b), genotype did not significantly affect mortality (p-value = 0.28 - Tab. 1).

First year total height was a highly significant covariate for total height after two growing seasons (p<0.01 - Tab. 2). Total height was not affected by spacing or spacing-by-genotype interactions, but varied by genotypes (p<0.01 - Tab. 2). Both varietals were taller than the half-sib seedlings, and CF Var 1 was approximately 10% taller than CF Var 2 (Fig. 2a). Height growth for the second growing season did not differ significantly among genotype and between spacings (Tab. 2), and averaged around 50 cm (Fig. 2b).

First year GLD was not a significant covariate (p = 0.08) in the ANOVA for total

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Fig. 1 - Mean percent (a) survival after the second growing season and (b) mortality (year 2 - year 1, \pm standard error) by genotype. Different letters indicate significant differences in average survival percentages and mortality among genotypes by growing season using least squares means test (α = 0.05).



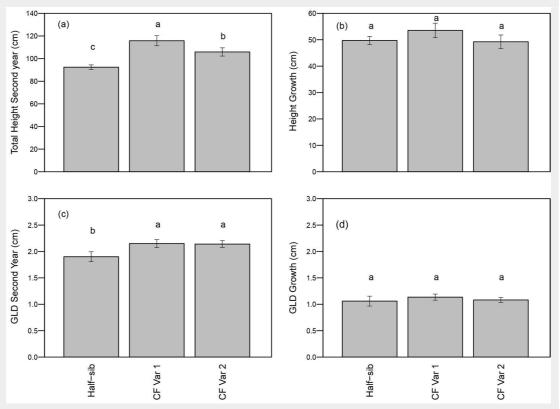
GLD at second year (Tab. 2). GLD of half-sib seedlings, CF Var 1, and CF Var 2 were 1.9 cm, 2.15 cm, and 2.14 cm, respectively (Fig. 2c). GLD and GLD growth during the second growing season were not affected by spacing or genotype (Tab. 2 and Fig. 2d).

In both years, the CV for height differed significantly among genotypes (p <0.01 - Tab. 3). As expected, clonal genotypes had greater uniformity (i.e., lower CVs) in height both years compared to the half-sib seedlings. CV for total height of half-sib seedlings was significantly higher (28% in the first year and 35% in the second year) than CF Var 1 (25% first year and 28% second year) or CF Var 2 (22% first year and 31% sec-

Tab. 2 - ANOVA table of loblolly pine second year GLD, height, GLD growth, height growth for spacing-by- genotype sub-plots. First year seedling GLD and height were used as covariates in the second year GLD and height respectively. (*): Denotes significance at α = 0.05; (1): first year GLD was the covariate for second year GLD; (2): first year height was the covariate for the second year total height.

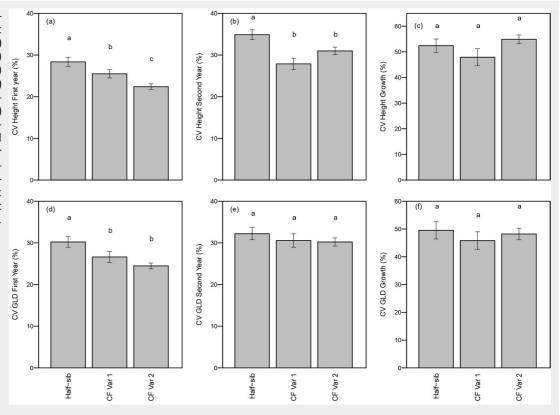
Effects	df	GLD (age 2) ¹	Height (age 2) ²	GLD Growth	Height Growth	
		P > F	P > F	P > F	P > F	
First Year (GLD/Height)	1	0.08 ¹	<0.01*2	=	-	
Spacing	1	0.15	0.58	0.14	0.53	
Block	1	0.86	0.67	0.90	0.07	
Block × Spacing	1	0.89	0.01*	0.86	0.01*	
Genotype	2	0.86	0.01*	0.8	0.31	
Block × Genotype	2	0.61	0.16	0.5	0.97	
Genotype × Spacing	2	0.47	0.37	0.35	0.30	

Fig. 2 - Mean (± standard error) (a) height after second growing season, (b) height growth, (c) GLD after second growing season, and (d) GLD growth by genotype. Means followed by a different letter indicate significant difference within growing season using least squares means test (α = 0.05).



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Fig. 3 - Mean (± standard error) coefficient of variation (CV) (a) first year height, (b) second year height, (c) height growth, (d) first year GLD, (e) second year GLD, and (f) GLD growth by genotype. Means followed by a different letter indicate significant differences within growing season using least squares means test $(\alpha = 0.05).$



Tab. 3 - ANOVA table of loblolly pine CV of GLD and height at first, second year, growth, and leaf water potential for spacing-by-genotype sub-plots. (*): Denotes significance at $\alpha = 0.05$.

Effects	df	GLD			Height			Leaf water
		age 1	age 2	Growth	age 1 P > F	age 2 P > F	Growth P > F	potential P > F
		P > F	P > F	P > F				
Spacing	1	0.36	0.4	0.57	0.70	0.97	0.99	0.81
Block	1	0.75	0.38	0.33	0.41	0.93	0.97	0.48
Block × Spacing	1	0.56	0.58	0.07	0.21	0.67	0.45	0.16
Genotype	2	0.01*	0.54	0.65	<0.01*	<0.01*	0.23	0.36
Block × Genotype	2	0.63	0.62	0.93	0.34	0.13	0.71	0.15
Genotype × Spacing	2	0.22	0.68	0.61	0.22	0.98	0.81	0.81

ond year - Fig. 3a and Fig. 3b). Furthermore, CV for height growth was not significantly affected by genotype (Tab. 3, Fig. 3c).

Tab. 4 - ANOVA table of effects on leaf water potential of measured loblolly pine seedlings composed of three genotypes and planted at two spacing across the second growing season (March to September 2013). (*): Denotes significance at α = 0.05.

Effects	DF	Leaf water potential		
		P > F		
Spacing	1	0.72		
Block	1	0.81		
Block × Spacing	1	0.30		
Genotype	2	0.97		
Block × Genotype	2	0.45		
Genotype × Spacing	2	0.33		
Month	6	<0.01*		
Genotype × Month	12	<0.01*		

Within a clone there is no tree-to-tree genetic variation so clonal seedlings are expected to be more uniform with a lower coefficient of variation (CV) than half-sib seedlings. First year CV for GLD differed significantly among genotypes (p = 0.01) (Tab. 3). CV for GLD of half-sib seedlings (30%) was significantly greater than CF Var 1 (26%) and CF Var 2 (24% - Fig. 3d). CVs for GLD and GLD growth were not significantly different among genotypes (Tab. 3, Fig. 3d, Fig. 3e, Fig. 3f).

Effects of genotype on leaf water potential and uniformity

Leaf water potential was not significantly affected by genotype and spacing (Tab. 4). Mean leaf water potential was -9.0 MPa across all genotypes (Fig. S1 in Supplementary material). Although leaf water potential was not significantly affected by genotype, when month was added to the model the month and month × genotype interaction effects significantly affected leaf water potential (Tab. 4). From March

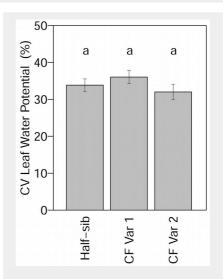


Fig. 4 - Mean (\pm standard error) CV of leaf water potential by genotype in 2013. Means followed by a different letter indicate significant differences in each month using least squares means test (α = 0.05).

through September, leaf water potential in June and leaf water potential of half-sib in August were significantly lower than other months. Leaf water potential was significantly higher for CF Var 2 in August (Fig. S1 in Supplementary material). The highest leaf water potential values were recorded in September, which corresponds to precipitation in September that was less than the precipitation average for the month in the past ten years. Coefficient of variation for leaf water potential did not differ significantly among genotypes (Tab. 3). The CV for leaf water potential of half-sib, CF Var 1, and CF Var 2 were 34%, 36%, and 32% (Fig. 4), respectively.

Discussion

Supporting our starting hypothesis, a greater growth and higher uniformity were observed in the clones. Although stands in this study are in their early stages of stand development, clones exhibited greater growth and uniformity compared to half-sib seedlings. However, significant difference among leaf water potential was not observed in this study.

During the initial stage of stand development, genotype had significant influences on survival and height (Tab. 1 and Tab. 2); however, GLD did not differ significantly among genotypes (Tab. 1). Significant genetic effects on GLD among half-sib, fullsib, and clonal seedling stocks has been reported (Aspinwall et al. 2011b). Although the percentage of survival and the height of clonal seedlings were significantly greater than half-sib seedlings (Fig. 1 and Fig. 2), these may have been influenced by factors associated with their production. For instance, the initial seedling size was unknown in this study; therefore, first year height was used as a covariate in the year 2 analysis and was found to be significant. In addition, height growth and GLD growth from the first to the second year were not significantly different among families (Tab. 2). These traits may indicate that the size differential among genotype was still affected by the initial size at time of planting. Other studies have found that initial seedling size affects total tree height and GLD after six years (South & Mason 1993) and after two years (Aspinwall et al. 2011b).

It is also likely that seedling type (i.e., containerized vs. bare-root) may have affected the growth and survival rate. In the current study, clonal seedlings were containerized whereas the half-sib seedlings were bareroot, which may have confounded the observed differences in height, GLD, and survivorship. A number of studies of southern pines on marginal or adverse sites have found that containerized stock survives and grows better than bare-root stocks (Barnett & Brissette 1986, 2004, Boyer 1989, Cram et al. 1999, Gwaze et al. 2006). Moreover, the two clonal stocks were grown and processed similarly and yet had a 10% difference in survival and a 10 cm difference in total height at the end of year two, potentially indicating a pure genetic response (Fig. 2). Interestingly, differences among the same planting stock were not evident in either GLD at the end of second year or GLD growth.

For individual clones which have no genetic variation (McKeand et al. 2006) from one tree to the next, uniformity in GLD and height is expected to be high (Martin et al. 2001, Bettinger et al. 2009) compared to the uniformity within half-sib genotypes, which is what was observed for height but not for height growth and GLD growth (Fig. 3). In contrast to this, Aspinwall et al. (2011b) found more variability in height growth in clones compared to open pollinated and full-sib genotypes, and they reported no consistent difference in GLD uniformity over time. Furthermore, they attributed the observed variability to microsite conditions on their minimally prepared, highly heterogeneous site. During the early stages of loblolly pine development in operational plantation conditions, genotypes with little or no genetic diversity may show greater sensitivity to changes in site environmental conditions (Aspinwall et al. 2011b). Under more intensive management conditions, clones may show more uniform growth and development (Martin et al. 2001, Bettinger et al. 2009).

Although leaf water potential among genotype was not significant, more uniformity in the leaf water potential was seen in the clones than in half-sib seedling. These findings may be due to variation in light condition, competing vegetation, nutrient availability, and water availability in this study site. Given the range of stress levels provided by the PMS instrument company for loblolly pine (PMS 2014), leaf water potential in this study site had a minimal impact in June (-o.6 MPa) and August (-o.7 MPa). In March, April, May, July, and September a slight to moderate impact of leaf water potential may have occurred, though no extreme values (greater than -1.2 MPa) were recorded.

One clone did outperform the half-sib in survival and height after two years, but whether this is due to genetic differences or seedlings type is unclear. If this performance is due to genetics, many studies have suggested that sustained productivity from genetic improvement can be sustained with the help of site-specific management (Fox 2000, Li et al. 1999, Nambiar 1996).

Conclusions

In this study, differences among the three planting stocks appeared very quickly, including between the two containerized clones, on the test site in southeastern Arkansas and provided further (albeit limited) support for the use of genetically improved planting stock. There also were seedling stock differences (i.e., bare-root vs. containerized) which confounded genetic effects. Moreover, CF Var 1 was found to have consistently higher rates of survival

as well as greater height when compared to the other containerized clone seedlings and bare-root half-sib seedlings. In terms of leaf water potential, neither clonal variety significantly outperformed the other in all measures. Furthermore, evidence that levels of genetic diversity would result in corresponding stand level uniformity for measured traits was not detected consistently. This finding suggest that less genetic diversity can lead to no difference in uniformity on a heterogeneous site which potentially decreases expected gains from the superior stock.

More time is needed to determine if early trends hold especially after inter-tree competition begins. Although one clone did outperform the improved half-sib seedlings both in survival and growth, the lower cost of the half-sib stock make it a popular choice for landowners in southeastern Arkansas who want to plant improved loblolly pine. With additional data on stand productivity among clonal, full-sib, and half-sib loblolly pine seedlings and with improvements in production and distribution of clonal stock, a greater proportion of forest landowners may seek to invest in genetically improved loblolly pine.

Results from this study suggest it is important to understand the response of an individual clone in growth and physiology under operational conditions with site specific management for the sustained productivity of the site. Adequate testing of highly selected genotypes should be continued to minimize risks and maximize productivity and uniformity.

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Supplementary Material

Fig. S1 - Mean (\pm standard error) leaf water potential by genotype by month in 2013. Means followed by a different letter indicate significant differences in each month using least squares means test (α = 0.05).

Link: Sharma 1725@supploo1.pdf

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