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Corrigendum

Corrigendum to “Growth responses of narrow or broad site adapted tree species to a range of resource availability treatments after a full harvest rotation” [For. Ecol. Manage. 362(2016)107–119]



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The authors regret that we neglected to include the change in fine root standing crop from one year to the next in our calculations of belowground net primary production (BNPP) and total net primary production (NPP). This affected Table 1 in the manuscript, and Supplemental Tables C and D. The authors would like to apologise for any inconvenience caused.

Appendix A. Supplementary material

Supplementary data associated with this article can be found, in the online version, at <http://dx.doi.org/10.1016/j.foreco.2016.03.015>.

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Table 1

Mean growth and productivity of five tree genotypes grown in South Carolina, USA after nine (2008) and 11 (2010) growing seasons. Within a genotype × year, treatment means followed by the same letter are not significantly different from other treatments within that genotype (Fisher's LSD, $\alpha = 0.05$). Among genotype means across treatments (shown below the 2008 and 2010 data) genotype means followed by the same letter are not significantly different from other genotypes (Fisher's LSD, $\alpha = 0.05$).

Year	Genotype	Treatment	Ht (m)	DBH (cm)	BA (m ² ha ⁻¹)	Vol. index (m ³ ha ⁻¹)	ANPP (Mg ha ⁻¹ yr ⁻¹)	BNPP (Mg ha ⁻¹ yr ⁻¹)	NPP (Mg ha ⁻¹ yr ⁻¹)	
2008	Cottonwood ST66	C	6.2 ^d	6.3 ^d	4.3 ^c	41.2 ^c	2.6 ^b	4.8 ^b	7.4 ^b	
		I	7.8 ^c	7.9 ^c	6.7 ^c	76.5 ^c	3.3 ^b	7.4 ^{ab}	10.7 ^b	
		F	11.0 ^b	10.2 ^b	11.3 ^b	176.8 ^b	4.8 ^{ab}	4.1 ^b	8.9 ^b	
		IF	13.6 ^a	12.4 ^a	17.4 ^a	343.8 ^a	7.2 ^a	10.1 ^a	17.3 ^a	
	Cottonwood S7C15	C	6.5 ^d	6.2 ^d	4.1 ^c	40.8 ^c	2.2 ^b	1.7	3.9 ^b	
		I	8.1 ^c	8.1 ^c	7.0 ^c	87.5 ^c	2.5 ^b	2.7	5.2 ^{ab}	
		F	11.2 ^b	10.6 ^b	11.8 ^b	194.3 ^b	4.7 ^{ab}	5.0	9.7 ^a	
		IF	13.5 ^a	12.2 ^a	17.2 ^a	343.7 ^a	6.4 ^a	3.3	9.7 ^a	
	Sycamore	C	7.8 ^d	6.9 ^c	5.2 ^c	52.7 ^d	2.7 ^c	6.0	8.7	
		I	11.2 ^c	10.0 ^b	10.4 ^b	156.3 ^c	5.5 ^{bc}	6.7	12.1	
		F	14.1 ^b	12.3 ^a	15.6 ^a	285.9 ^b	6.8 ^{ab}	5.1	12.0	
		IF	16.1 ^a	13.6 ^a	17.4 ^a	367.4 ^a	8.6 ^a	3.3	11.9	
	Sweetgum	C	8.8 ^b	9.1 ^b	8.8 ^d	99.7 ^c	4.0 ^d	11.8 ^{ab}	15.7 ^b	
		I	10.0 ^b	10.4 ^b	12.2 ^c	167.9 ^c	7.7 ^c	12.2 ^a	19.9 ^{ab}	
		F	12.9 ^a	13.4 ^a	19.4 ^b	319.7 ^b	11.8 ^b	8.6 ^b	20.4 ^{ab}	
		IF	14.1 ^a	14.6 ^a	23.1 ^a	420.3 ^a	16.5 ^a	7.5 ^c	24.0 ^a	
	Loblolly Pine	C	11.9 ^b	15.1 ^b	23.6 ^c	370.4 ^c	14.3 ^c	9.5	23.8 ^b	
		I	11.5 ^b	14.2 ^b	20.9 ^c	315.6 ^c	14.9 ^c	8.9	23.7 ^b	
		F	12.3 ^b	17.2 ^a	32.0 ^b	508.4 ^b	19.1 ^b	8.4	27.5 ^{ab}	
		IF	14.6 ^a	18.3 ^a	35.1 ^a	671.9 ^a	22.4 ^a	9.9	32.3 ^a	
Genotype mean across treatment	ST66	–	9.7 ^b	9.2 ^c	10.0 ^c	159.9 ^b	4.5 ^c	6.6 ^b	11.1 ^c	
	S7C15	–	9.8 ^b	9.3 ^c	10.0 ^c	166.6 ^b	4.0 ^c	3.2 ^c	7.1 ^d	
	Sycamore	–	12.3 ^a	10.7 ^b	12.2 ^b	215.6 ^b	5.9 ^c	5.3 ^b	11.2 ^c	
	Sweetgum	–	11.5 ^a	11.9 ^b	15.9 ^b	251.9 ^b	10.0 ^b	10.0 ^a	20.0 ^b	
	Loblolly Pine	–	12.6 ^a	16.2 ^a	27.9 ^a	466.6 ^a	17.7 ^a	9.2 ^a	26.8 ^a	
2010	Sweetgum	C	9.5 ^d	9.7 ^b	10.1 ^c	123.3 ^c	4.8 ^c	11.9 ^a	16.8 ^a	
		I	11.1 ^c	11.2 ^b	14.3 ^b	220.6 ^c	7.4 ^c	10.9 ^{ab}	18.3 ^a	
		F	14.4 ^b	14.9 ^a	24.1 ^a	445.1 ^b	16.4 ^b	9.8 ^{ab}	26.2 ^b	
		IF	15.8 ^a	16.1 ^a	28.0 ^a	574.5 ^a	21.3 ^a	7.2 ^b	28.5 ^b	
	Loblolly Pine	C	13.7 ^b	16.6 ^b	28.7 ^b	518.0 ^c	13.8 ^b	12.5	26.3 ^{bc}	
		I	13.2 ^b	15.8 ^b	25.7 ^b	443.1 ^c	15.3 ^b	10.2	25.5 ^c	
		F	15.4 ^a	19.4 ^a	39.2 ^a	772.2 ^b	21.2 ^a	10.6	31.8 ^{ab}	
		IF	16.7 ^a	20.7 ^a	41.8 ^a	895.6 ^a	21.4 ^a	10.6	32.0 ^a	
	Genotype mean across treatment	Sweetgum	–	12.7 ^b	13.0 ^b	19.1 ^b	340.9 ^b	12.5 ^b	10.0	22.4 ^b
		Loblolly Pine	–	14.8 ^a	18.2 ^a	33.9 ^a	657.2 ^a	17.9 ^a	11.0	28.9 ^a