

Establishment of hybrid Poplar on a Reclaimed Mine site in West Virginia

A. Hass,¹ R.S. Zalesny Jr.¹, D. Patel², J. Vandevender³

¹ West Virginia State University; Agricultural & Environmental Research Station; Institute, WV

² U.S. Forest Service; Northern Research Station; Rhinelander, WI

³ U.S. USDA NRCS Plant Material Center, Alderson, WV



WEST VIRGINIA STATE
UNIVERSITY



Land Productivity



Picture by Dale K. Ritchey

Land Productivity



Land Productivity



Yield Gap (Actual [management])

NO FERT
NO LIME

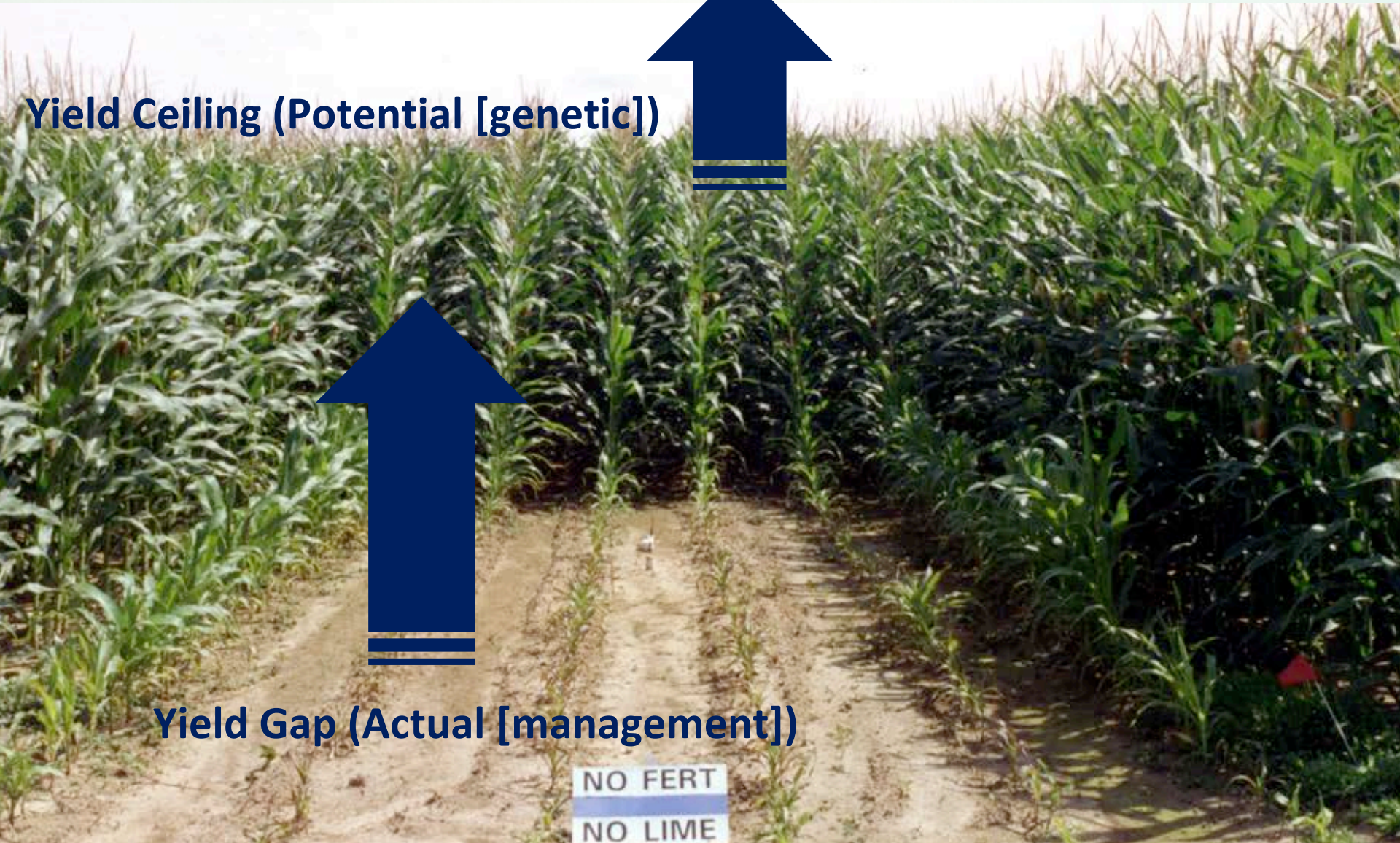
Land Productivity

Yield Ceiling (Potential [genetic])



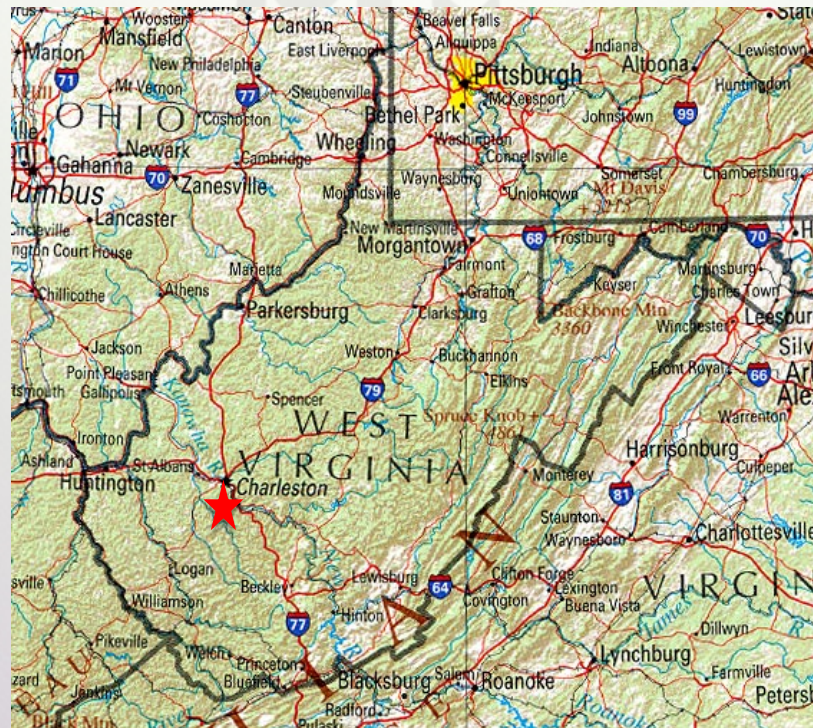
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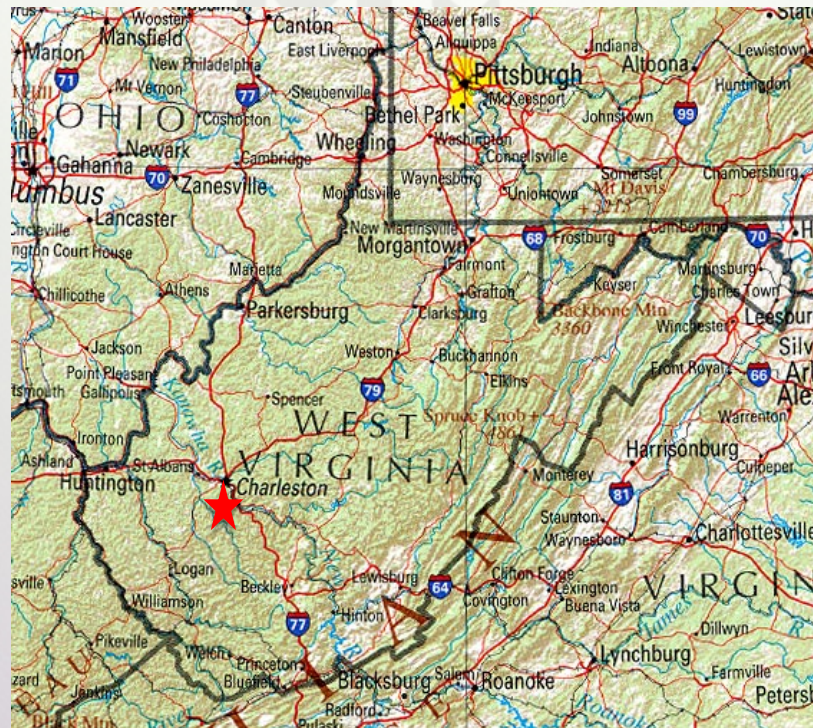
Objectives

- ❑ Use phyto-recurrent selection to identify *Populus* genotypes that grow better on reclaimed mine sites.



Objectives

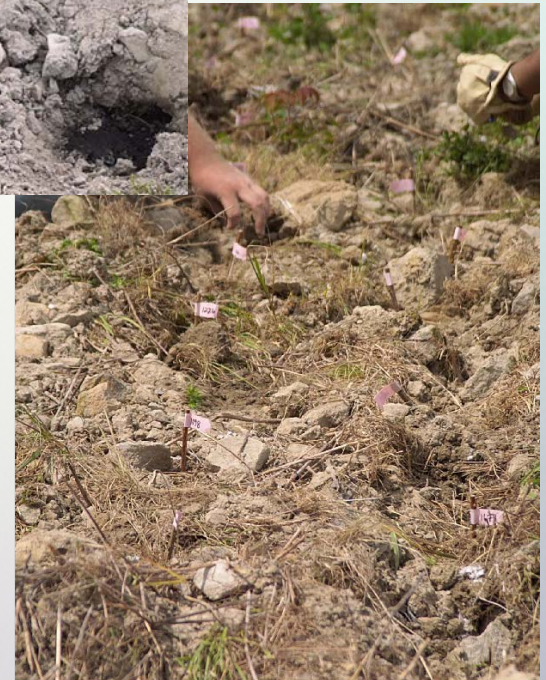
- ❑ Use phyto-recurrent selection to identify *Populus* genotypes that grow better on reclaimed mine sites.
- ❑ Test whether biochar can improve tree survival & growth on reclaim mine site.



Site Prep and Soil Treatment



Site Prep and Soil Treatment



Genomic Group	Clone
<i>P. deltoides</i> 'D'	8000105, C910809, D105, D109, D110, D112, D117, D118, D121, D125, D133, D134
<i>P. trichocarpa</i> 'T'	0.20.3, 0.4.3, 0.6.3, 0.7.5, 1.1.5, 1.2.5, 1.3.1, 2.1.2, 3.2.2, 4.1.3, 5.2.2
<i>P. deltoides</i> × <i>P. deltoides</i> 'DD'	119.16
<i>P. deltoides</i> × <i>P. nigra</i> 'DN'	DN154, DN164, DN17, DN170, DN177, DN182, DN34, DN5, DN70, NE264
<i>P. deltoides</i> × <i>P. suaveolens</i> subsp. <i>maximowiczii</i> 'DM'	113.64, 412.52, 202.37, 313.23, 313.55, DM105, DM108, DM111, DM112, DM113, DM114, DM115, DM117, NC14103, NC14104, NC14105, NC14106, NC14107
<i>P. nigra</i> × <i>P. suaveolens</i> subsp. <i>maximowiczii</i> 'NM'	NM2, NM6
<i>(P. trichocarpa</i> × <i>P. deltoides)</i> × <i>P. deltoides</i> 'TDD'	NC13536, NC13555, NC13624, NC13728, NC13740, NC13820



Algelros Duby – *P. deltoides* Bartr. ex Marsh, *P. nigra* L.

Tacamahaca Spach – *P. suaveolens* Fischer subsp. *maximowiczii* A. Henry, *P. trichocarpa* Torr. & Gray

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CYCLE 1

- Field (2014)
- Survival
- Ht, Diam
- Biomass
- Health

	Cycle 1	Cycle 2
Rep	8	8
Soil treatment	2	0
Clone	60	32

CYCLE 2

- Field (2015)
- Survival
- Ht, Diam
- Health

DEPLOYMENT



CYCLE 1

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- Survival
- Ht, Diam
- Biomass
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	Cycle 1	Cycle 2
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Soil treatment	2	0
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$$IV = 0.3 * MASS + 0.15 * HEIGHT + 0.15 * DIAMETER + 0.2 * SURVIVAL + 0.2 * HEALTH$$

CYCLE 2

- Field (2015)
- Survival
- Ht, Diam
- Health

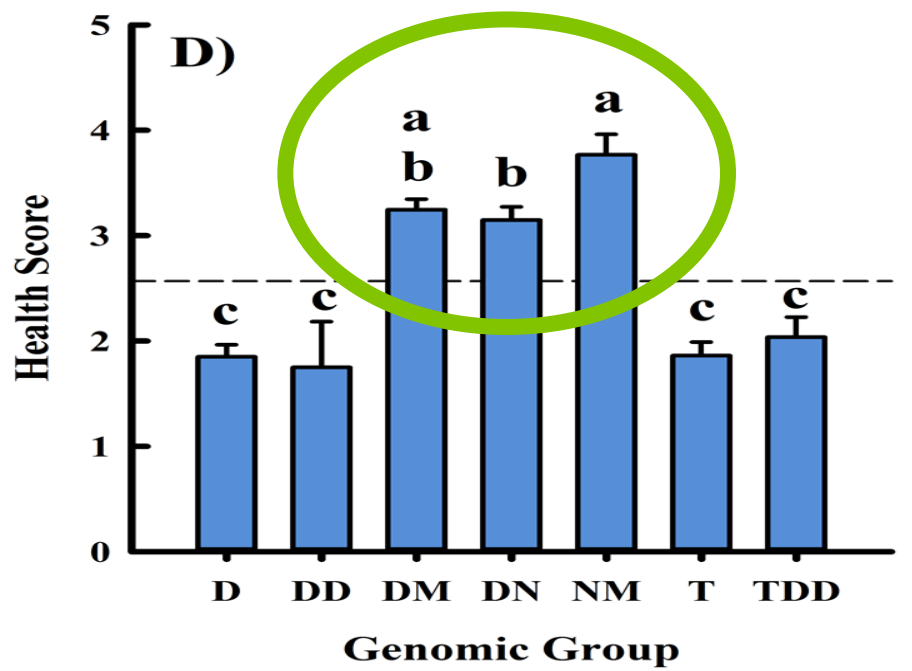
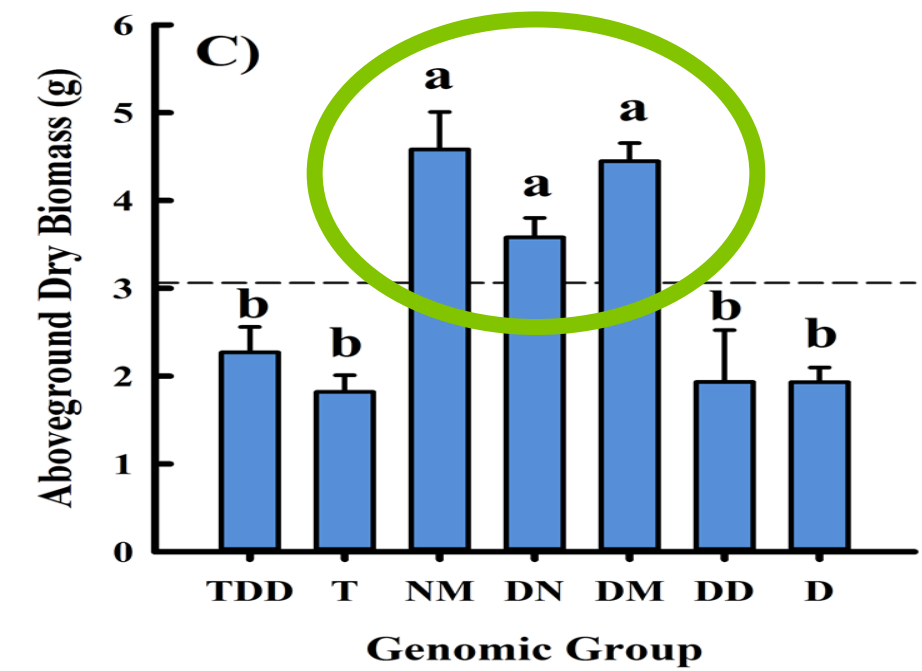
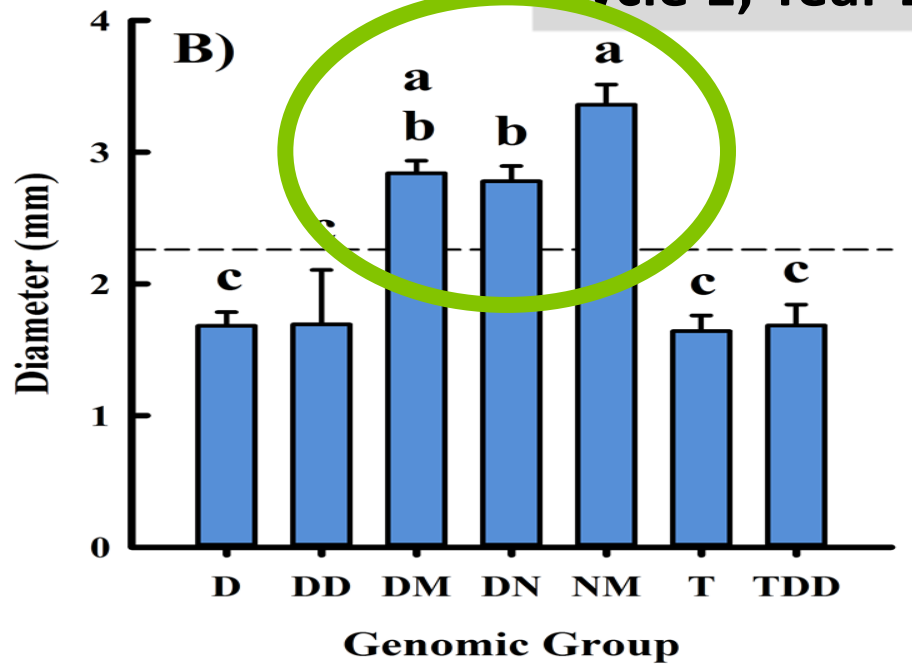
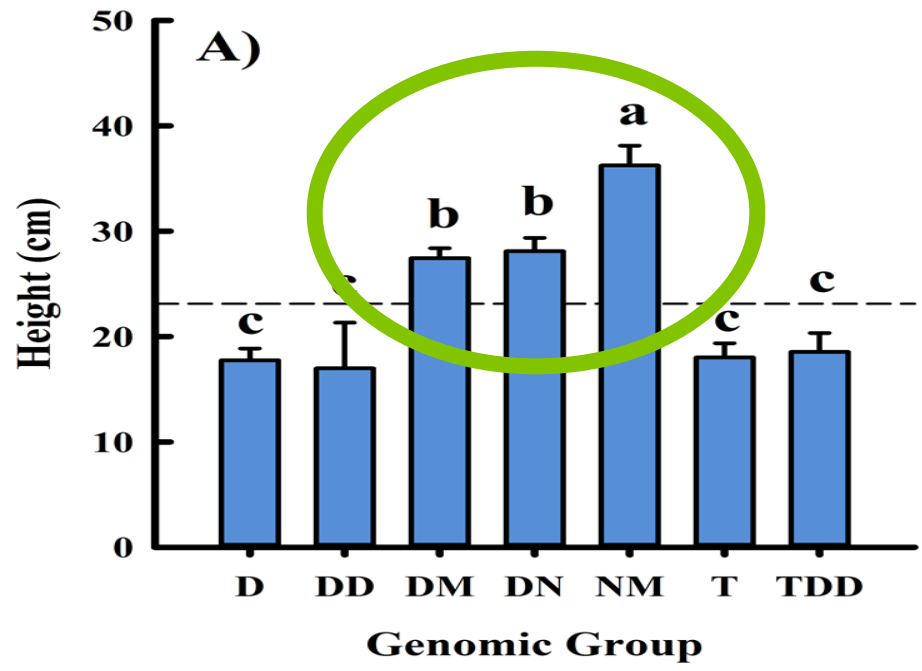
DEPLOYMENT

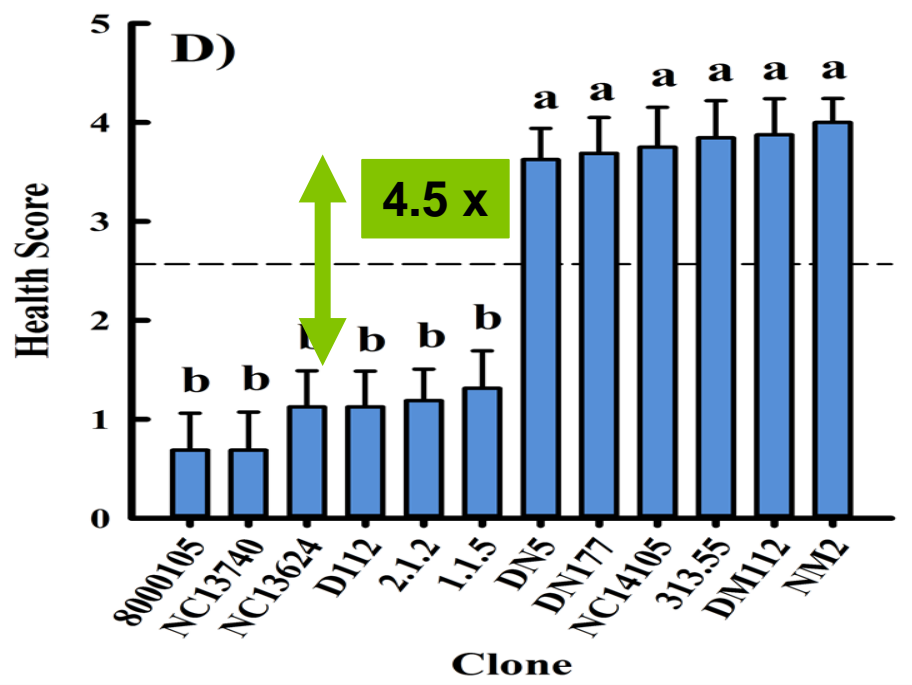
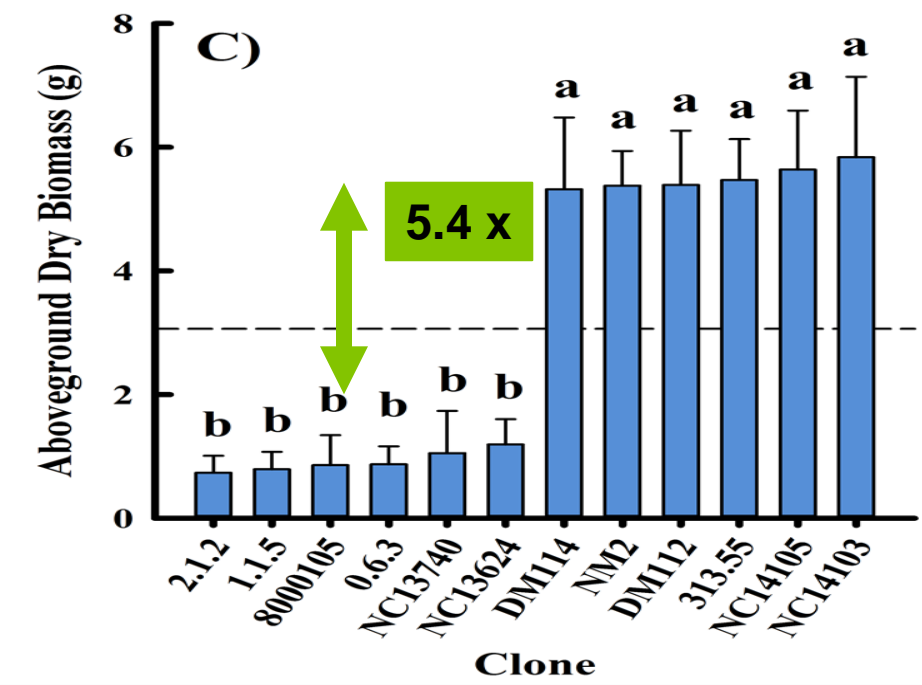
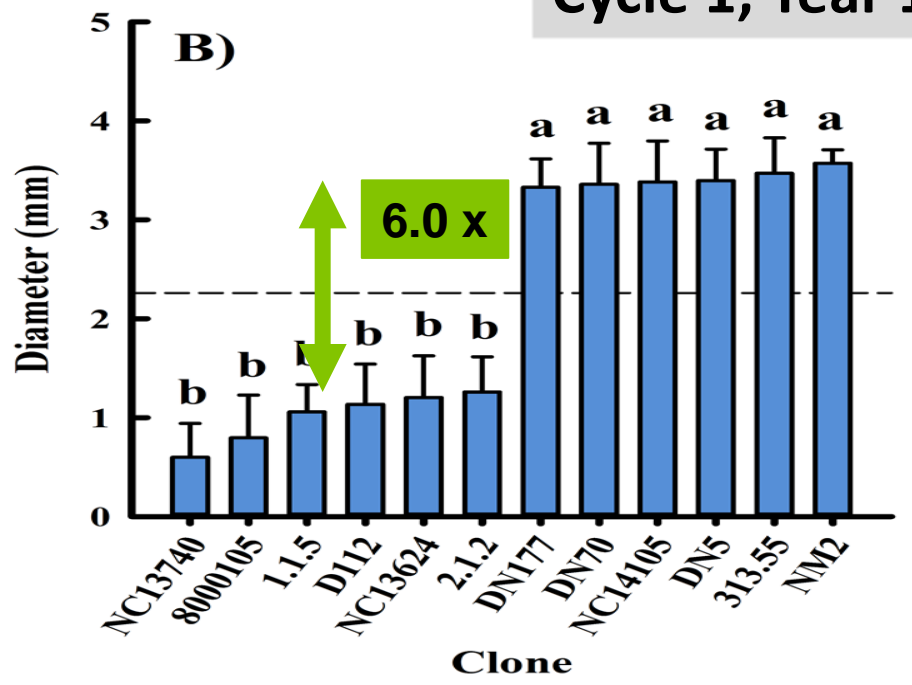
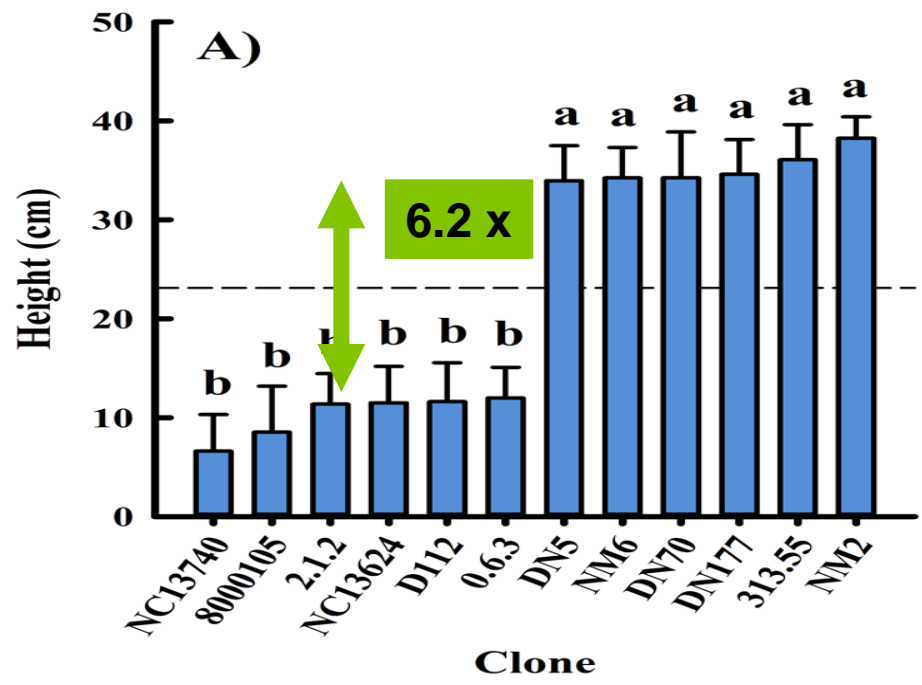


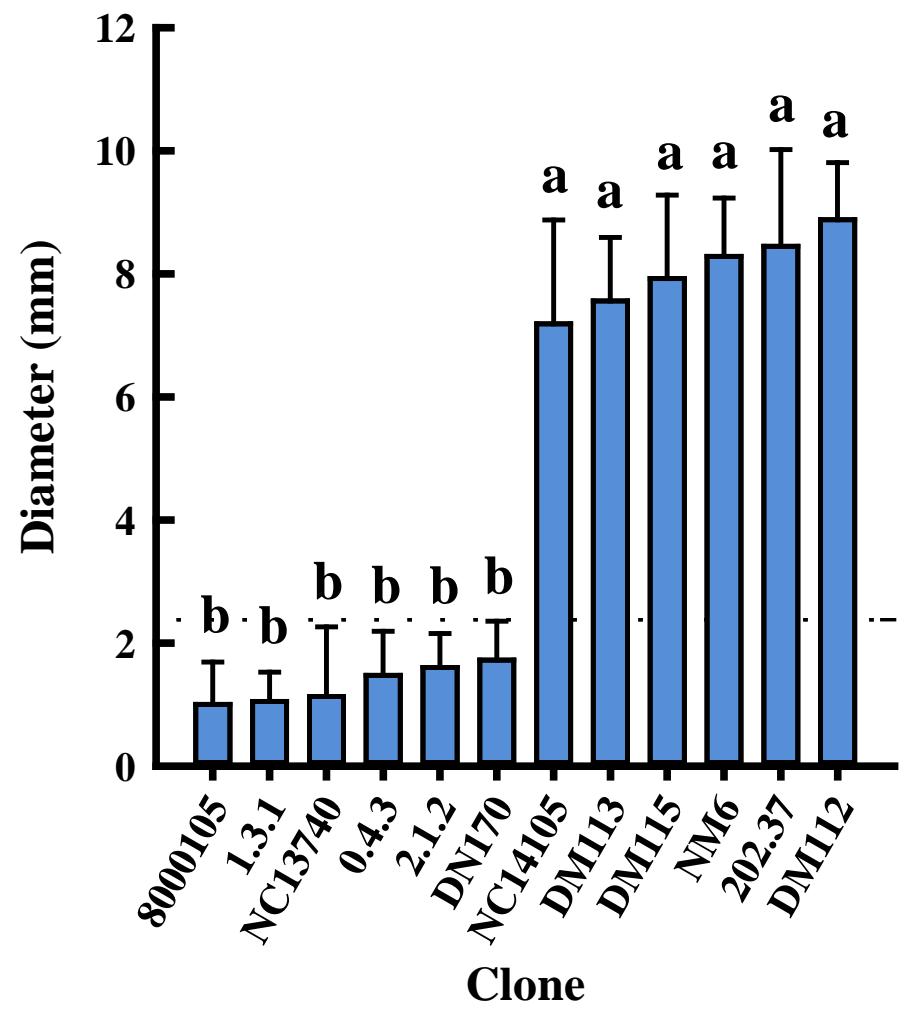
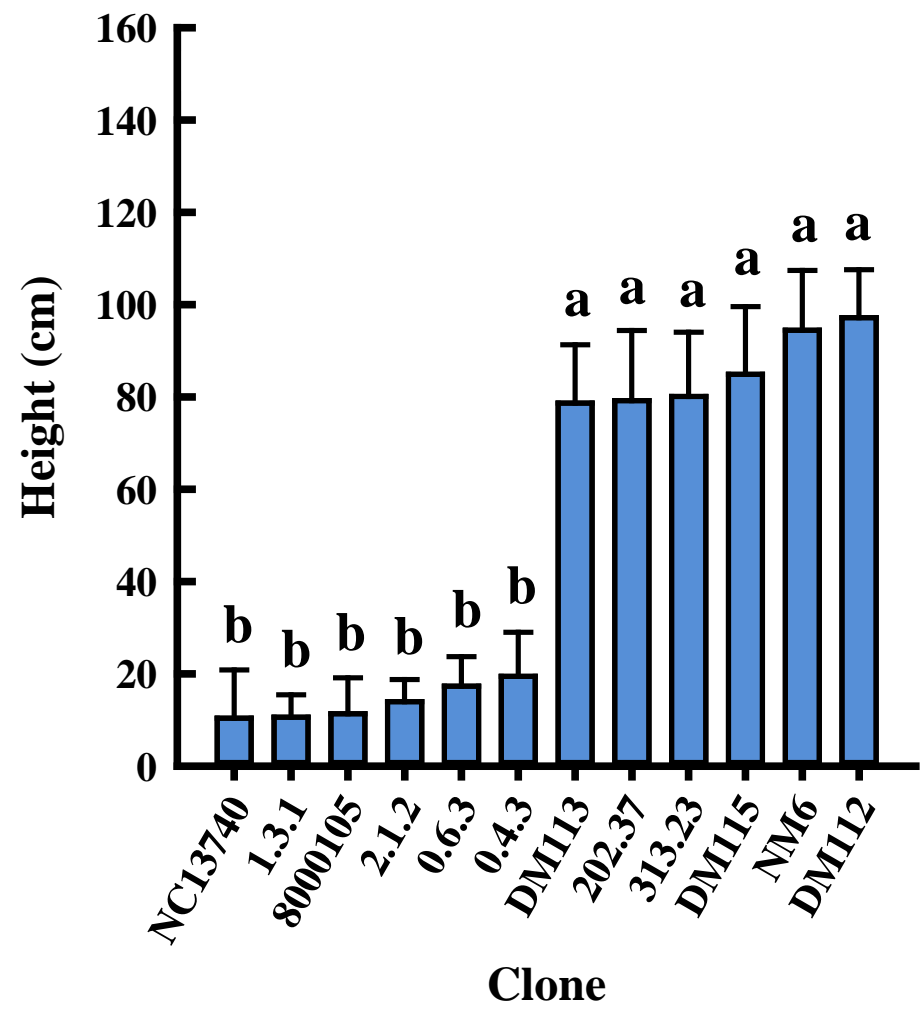
Cycle 1 Probability Values

Source of variation	Height	Diameter	Biomass	Health
Soil Amendment	0.4749	0.7067	0.9130	0.5402
Clone	<0.0001	<0.0001	<0.0001	<0.0001
Soil Treatment × Clone	0.6148	0.4565	0.8831	0.2175
Soil Amendment	0.3755	0.5484	0.9823	0.6691
Genomic Group	<0.0001	<0.0001	<0.0001	<0.0001
Soil Treatment × Genomic Group	0.4080	0.5404	0.2768	0.4137









Rank	Clone	Genomic Group	Index Value	Rank	Clone	Genomic Group	Index Value
----- Clones selected for cycle 2 -----				----- Clones not selected for cycle 2 -----			
1	NM2	NM	8.89	32	1.3.1	T	5.27
2	313.55	DM	8.53	33	0.4.3	T	5.27
3	DN177	DN	8.13	34 ^a	NC13728	TDD	5.11
4	NC14105	DM	8.11	35	D121	D	4.97
5	DN70	DN	8.03	36	D117	D	4.80
6	DN5	DN	7.86	37	1.2.5	T	4.78
7	NM6	NM	7.65	38	DN34	DN	4.69
8	NC14103	DM	7.56	39	5.2.2	T	4.67
9	DM112	DM	7.55	40	3.2.2	T	4.64
10	DM113	DM	7.23	41	113.64	DM	4.63
11	DM114	DM	7.17	42	NC13536	TDD	4.58
12	202.37	DM	7.05	43	D109	D	4.57
13	NC14104	DM	6.98	44	D110	D	4.48
14	NC14106	DM	6.69	45	D105	D	4.28
15	313.23	DM	6.67	46	4.1.3	T	3.88
16	DN17	DN	6.61	47	0.20.3	T	3.87
17	DN170	DN	6.60	48	119.16	DD	3.84
18	412.52	DM	6.55	49	D134	D	3.81
19	DM117	DM	6.53	50	D125	D	3.71
20	DM108	DM	6.42	51	0.7.5	T	3.56
21	NC13820	TDD	6.25	52	D133	D	3.40
22	C910809	D	6.01	53	D118	D	3.33
23	DN154	DN	5.90	54	1.1.5	T	2.83
24	NE264	DN	5.88	55	D112	D	2.74
25	DM115	DM	5.81	56	0.6.3	T	2.69
26	NC14107	DM	5.80	57	NC13624	TDD	2.58
27	DN164	DN	5.77	58	2.1.2	T	2.45
28	DM111	DM	5.75	59	8000105	D	1.83
29	DN182	DN	5.60	60	NC13740	TDD	1.58
30	DM105	DM	5.53				
31	NC13555	TDD	5.41				

^a Despite not being ranked in the top 32 genotypes, clone NC13728 was advanced to cycle 2 in order to get adequate representation of the TDD clones for the genomic group analyses.

$$IV = 0.3*MASS + 0.15*HEIGHT + 0.15*DIAMETER + 0.2*SURVIVAL + 0.2*HEALTH$$

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6	DN5	DN	7.86	37	1.2.5	T	4.78
7	NM6	NM	7.65	38	DN34	DN	4.69
8	NC14105	DM	7.56	39	3.2.2	T	4.67
9	DM112	DM	7.55	40	3.2.2	T	4.64
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25	DM115	DM	5.81	56	0.6.3	T	2.69
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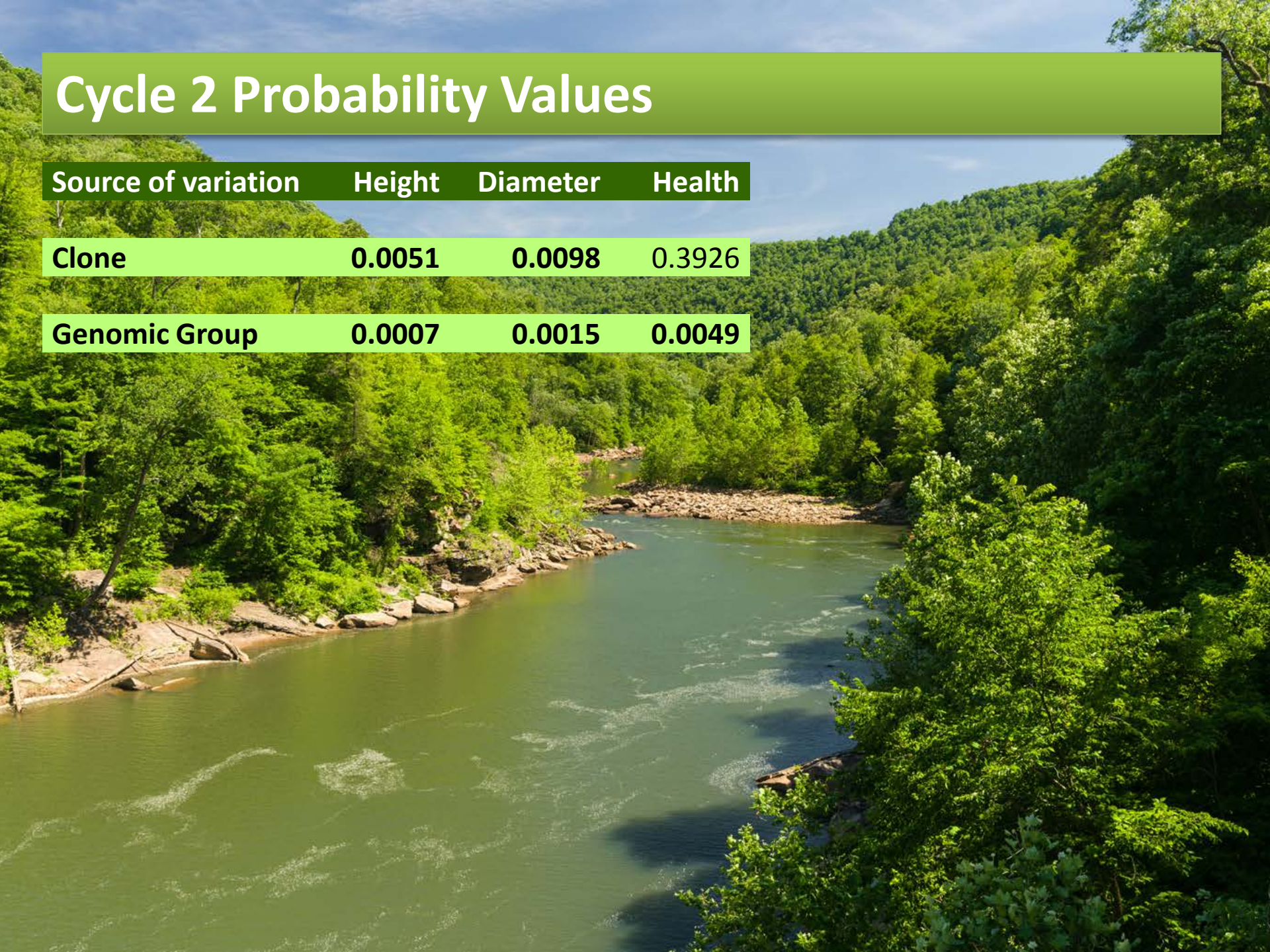
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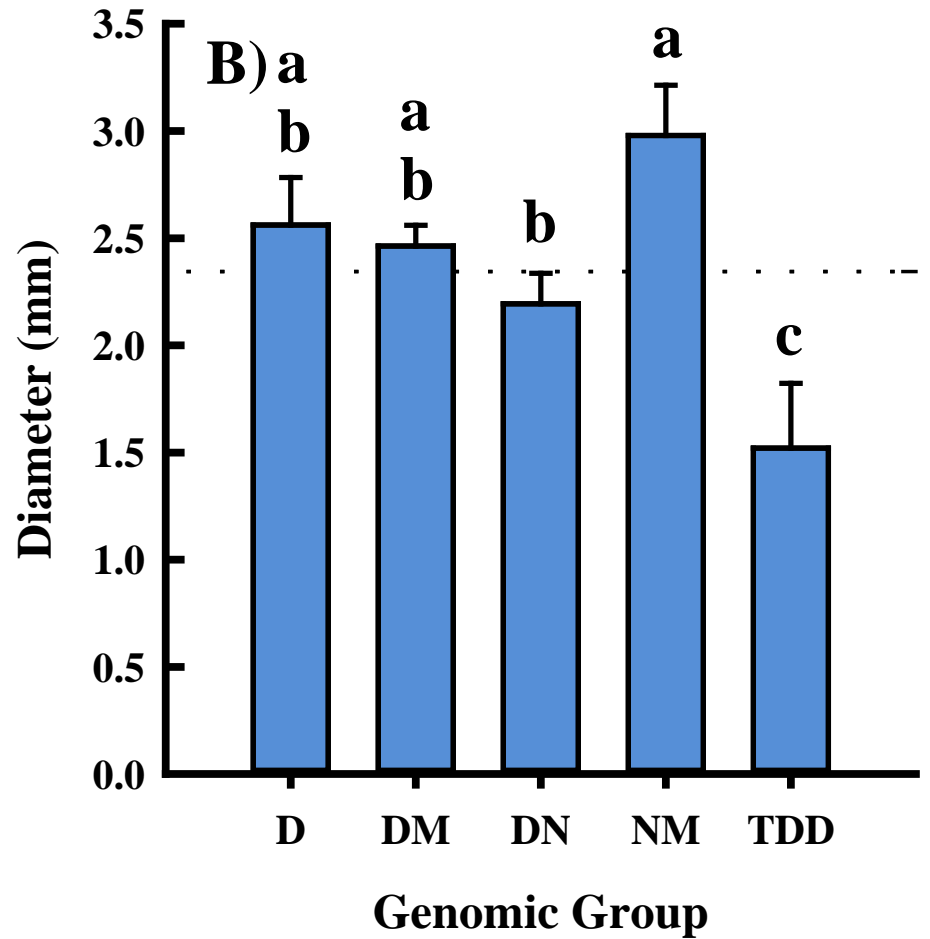
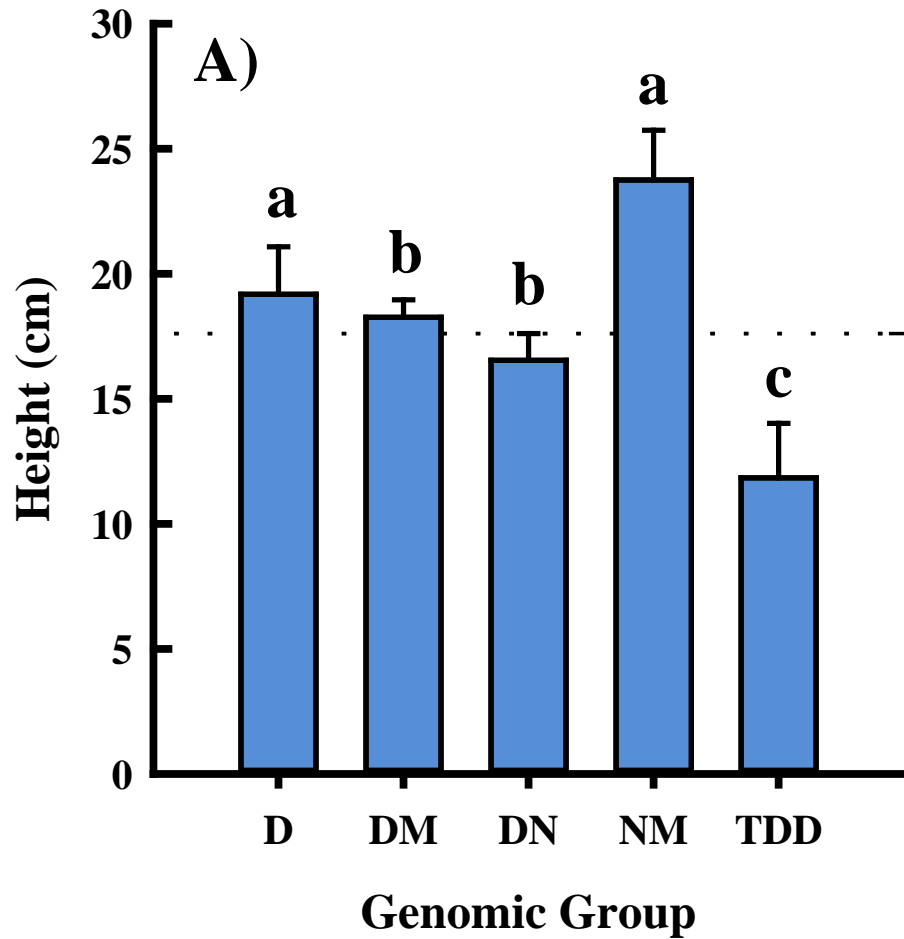


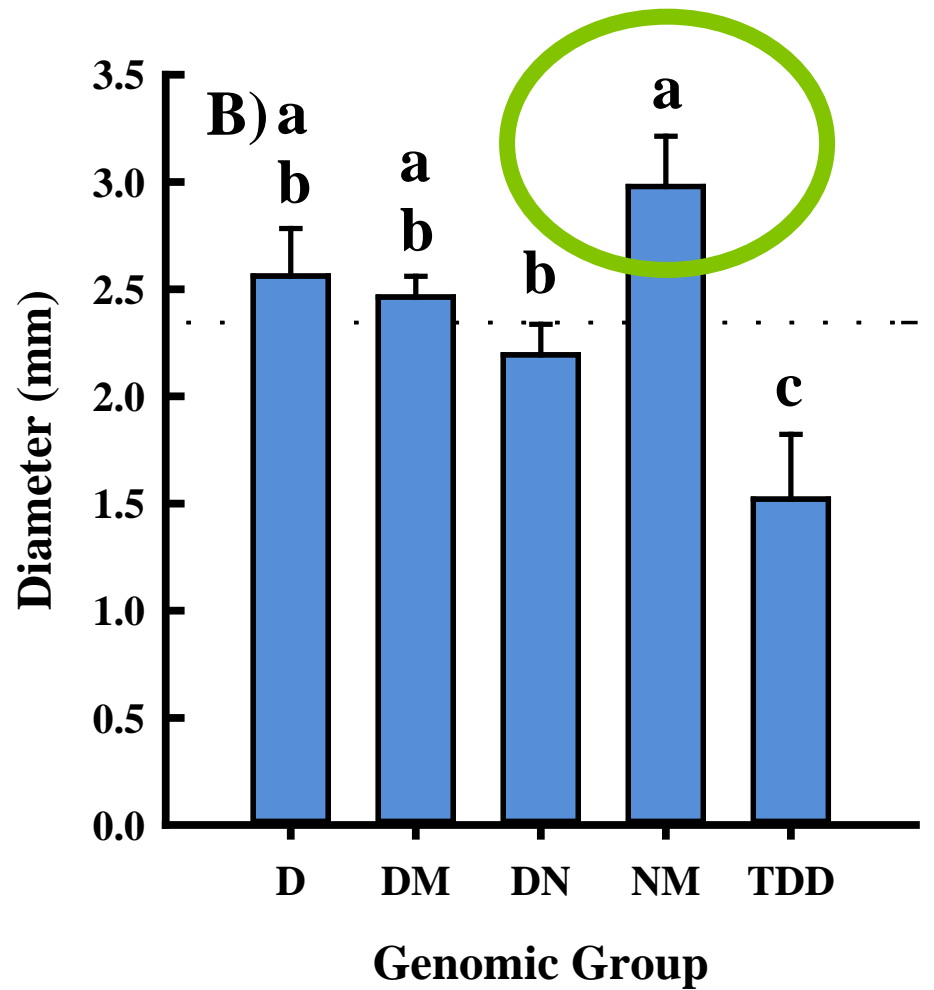
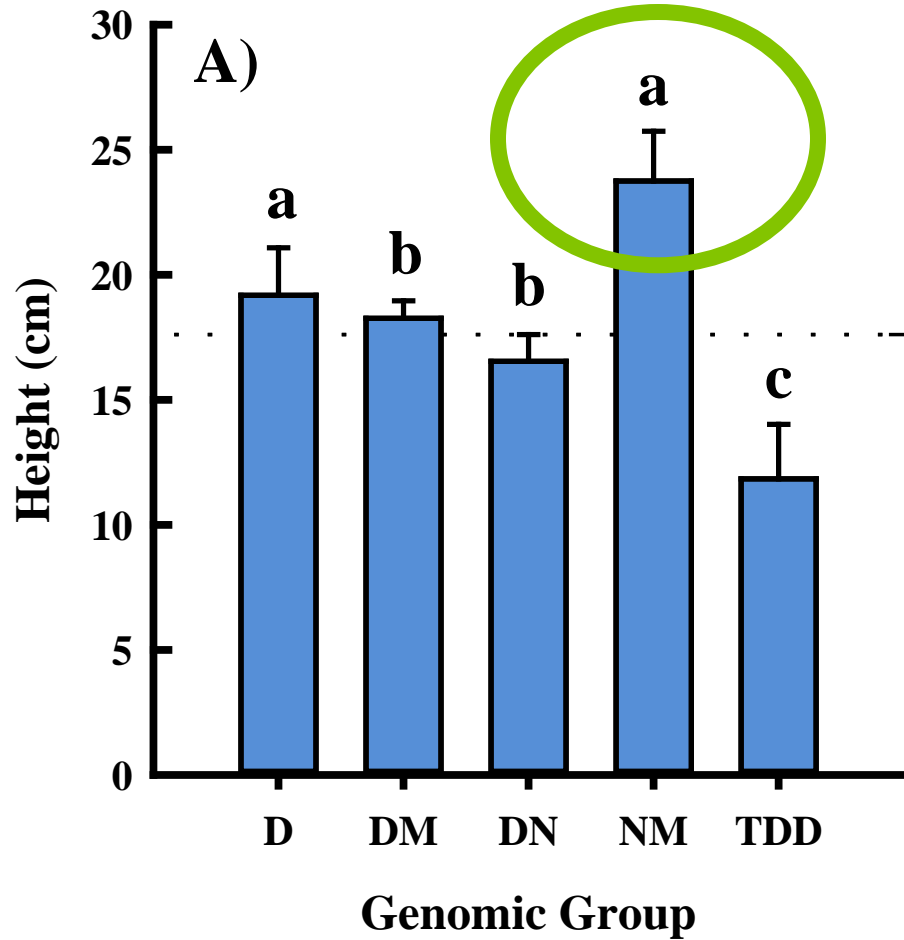
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Cycle 2 Probability Values

Source of variation	Height	Diameter	Health
Clone	0.0051	0.0098	0.3926
Genomic Group	0.0007	0.0015	0.0049







Conclusions

❑ SOIL AMENDMENT PRACTICES

- ❑ Biochar treatment DID NOT significantly increase tree survival & growth relative to standard soil ripping techniques without amendments

❑ PLANT SELECTION

- ❑ Phyto-recurrent selection is a viable tool for selecting superior genotypes
- ❑ *P. nigra* × *P. suaveolens* subsp. *maximowiczii* 'NM' performances were superior to all other genomic groups



Funding

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Marshall University's Center for Environmental, Geotechnical & Applied Sciences (CEGAS)

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West Virginia Division of Energy Office of Coalfield Community Development
&
Appalachian Regional Commission

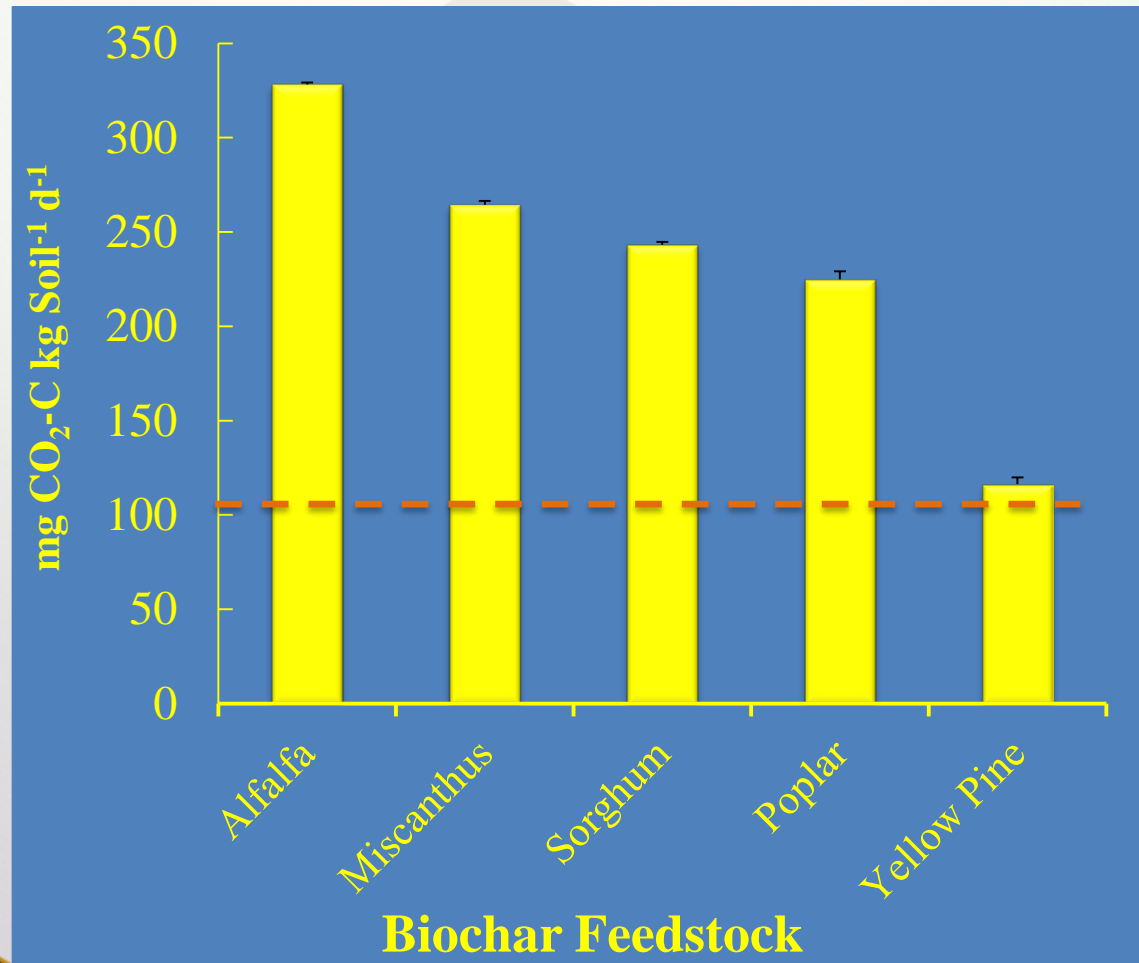


Soil Treatment - Biochar

<i>Biochar</i>	pH	EC (mS/cm)	Ash Content	P	S	C	N	O	C/N Ratio
				----- % weight -----					
<i>Alfalfa</i>	12	11.2	28.5	0.85	0.14	57	1.54	23	37
<i>Miscanthus</i>	8.3	7.08	27.3	1.22	0.19	54	1.04	25	52
<i>Poplar</i>	9.2	0.72	26.2	-	-	61	0.67	22	91
<i>Sorghum</i>	10	4.32	35.5	0.48	0.19	50	0.61	26	82
<i>Yellow Pine</i>	10	9.59	-	-	-	75	0.32	20	238

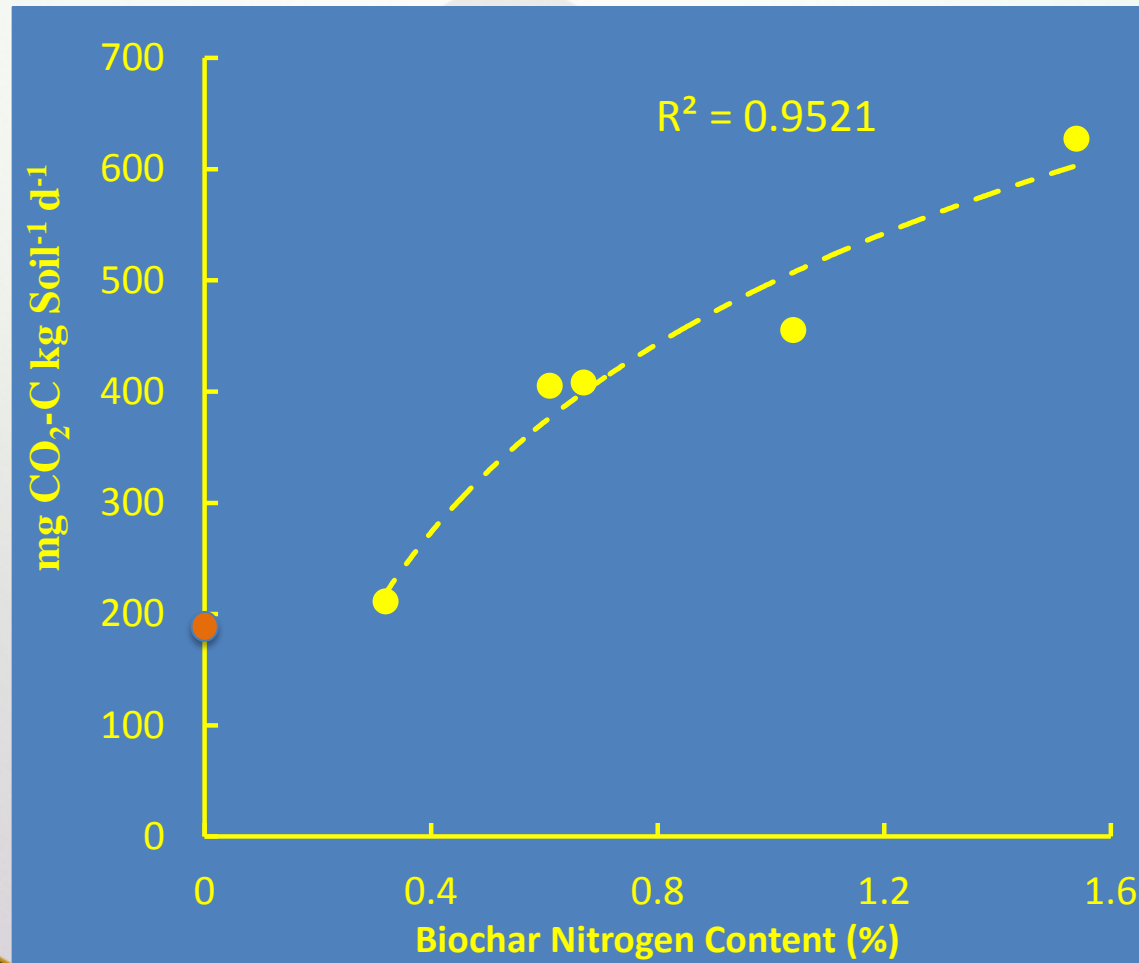
Soil Treatment - Biochar

CO₂ Release From Biochar Amended Soil – Feedstock Effect



Soil Treatment - Biochar

CO₂ Release From Biochar Amended Soil – Char TN Content



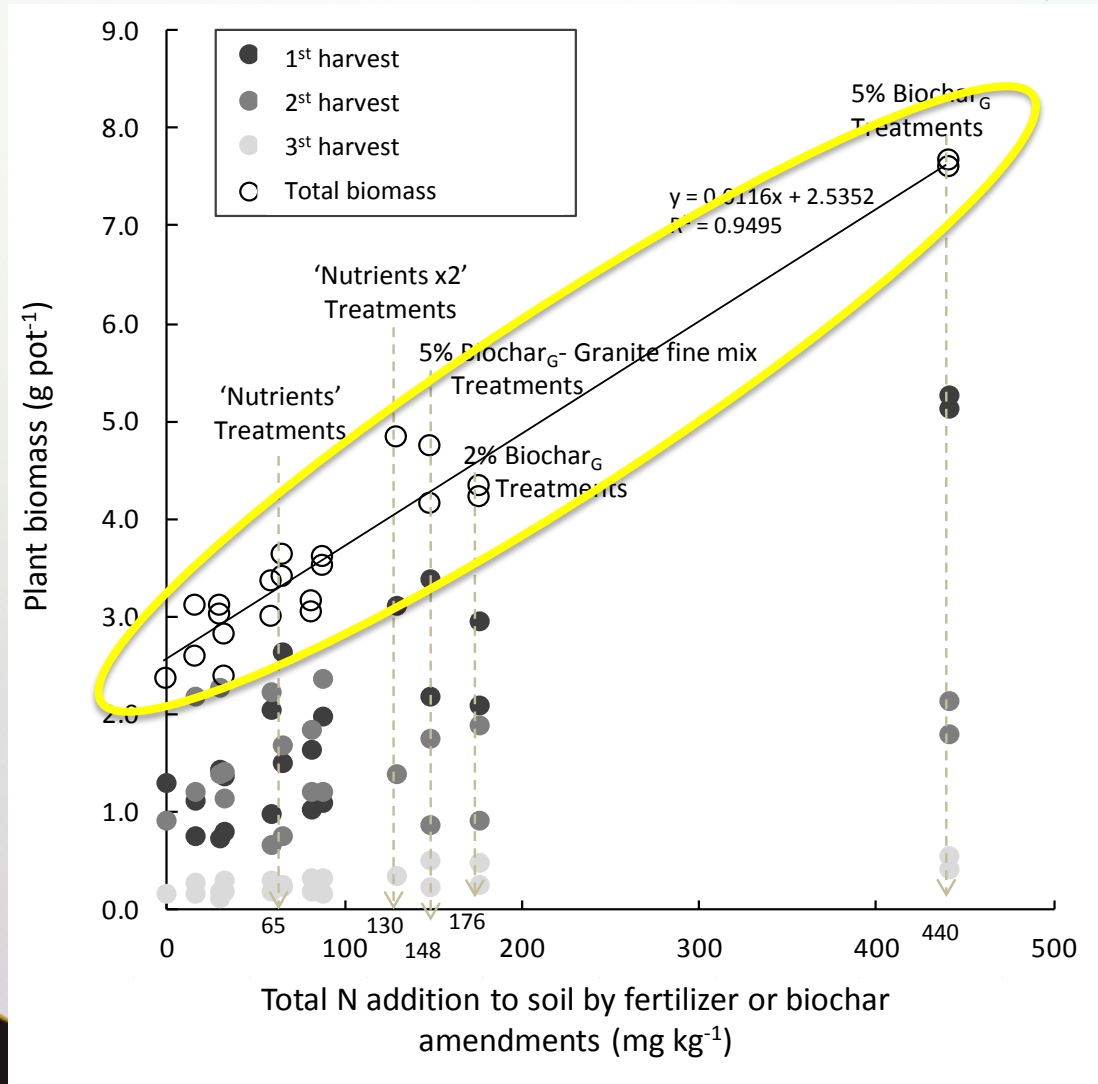
Soil Treatment - Biochar

Biochar Amended Soil – Foxtail Millet Growth Response

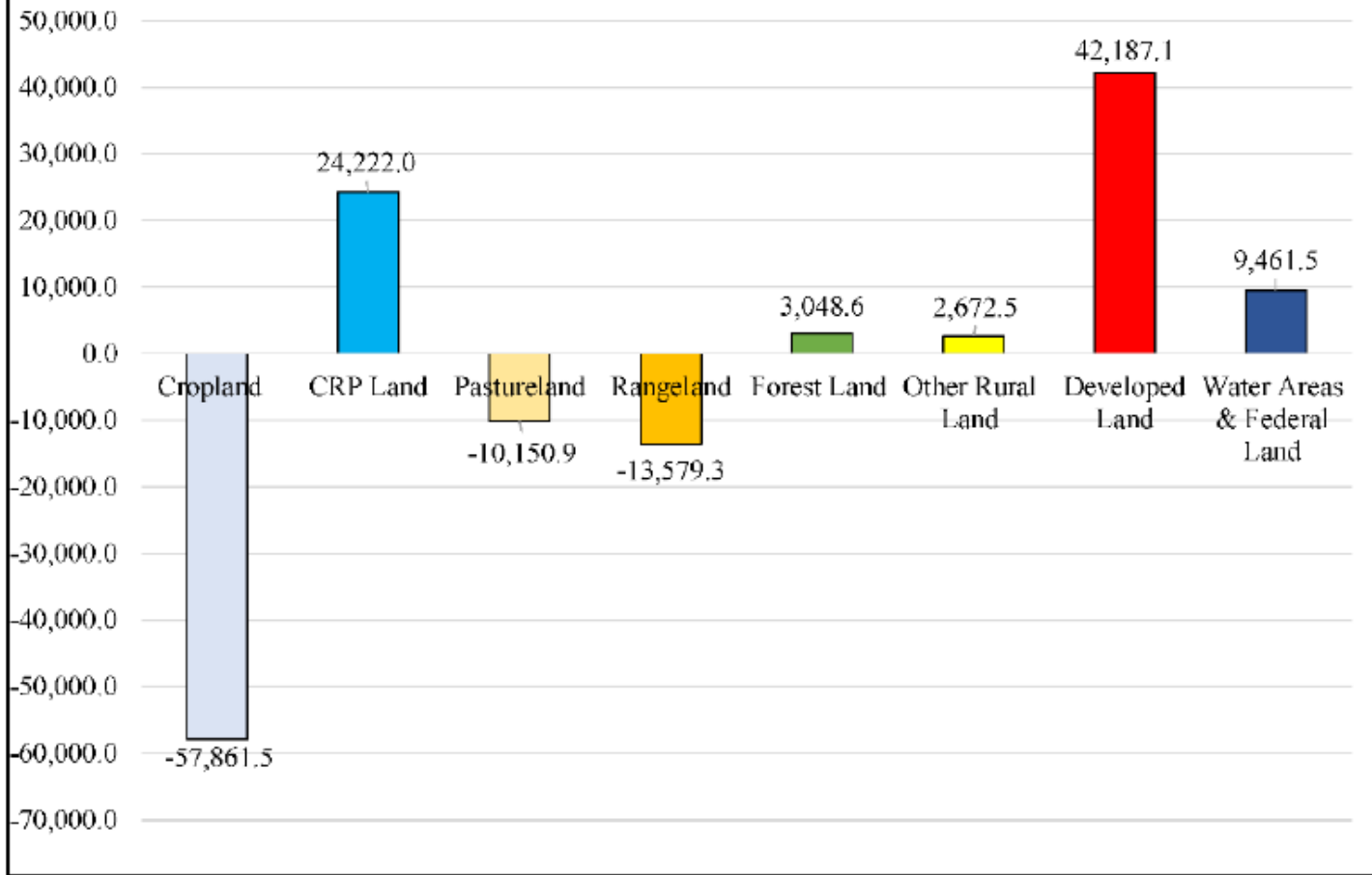


Soil Treatment - Biochar

Biochar Amended Soil – Foxtail Millet Growth Response

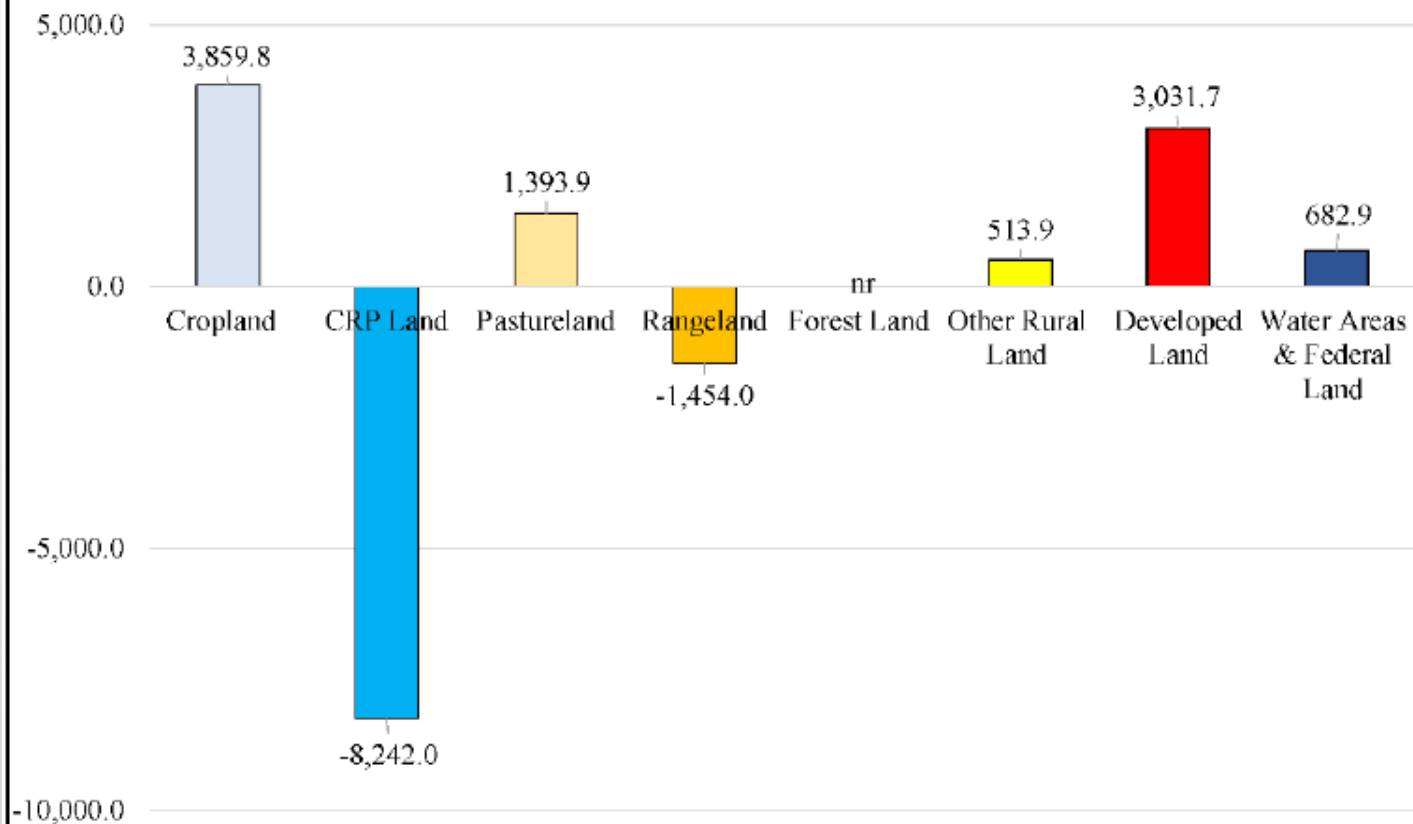


Net Change in Land Cover/Use between 1982 and 2012 Thousands of Acres



USDA 2015. 2012 National Resources Inventory

Net Change in Land Cover/Use between 2007 and 2012 Thousands of Acres



The net change estimate for forest land is not reliable (nr) as the margin of error is greater than the estimate.

