Select results

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Introduction

- Presented here are the results of single-moderator and model selection analysis for the exotic and native datasets.
- All models have undergone sensitivity analysis and outliers have been removed.
- Results are summarized in the form of forest plots, and heterogeneity statistics are presented in a single table.
- Code is omitted to conserve space. See other documents for code and more details.
- Calculating the probability based on the prediction interval has been updated since the 11/2021 results
 document.

Summary

Summary effect

- Overall, C addition treatment significantly decreases exotic plant abundance but does not affect native plant abundance.
 - When predicting the effect of C addition on a random population, we would expect exotic plant abundance to decrease 80% of the time, and native plant abundance to increase 41% of the time.

Single moderator models

- C addition treatment has been most successful in southeast Australia (New South Wales) and the western United States, especially in the Southwest and Pacific Northwest.
- C addition treatment steadily becomes less effective over time (effect size decreases as duration since first and last C application increases).
- Sucrose alone is more effective than sawdust alone; but both had a significant effect when used alone, but not in combination.
 - However, this could be an artifact of sample size/number of comparisons. Only sucrose, sucrose AND sawdust, and sawdust have comparable n values (number of studies/comparisons).
- Increasing C rate decreases exotic plant abundance and increases native plant abundance.
 - Although these trends are statistically significant, practically speaking, the increase in effect likely
 does not compensate for the higher cost.
- The number and timing of C applications show inconsistent trends, but studies that applied C only once saw a simultaneous decrease in exotic plant abundance and increase in native plant abundance.

- Studies that applied C 3-6 times reported the greatest decrease in exotic plant abundance; however, native plant abundance also decreased.
- C application treatment was more effective on annual exotic plants than perennial, but perennial graminoids were still significantly affected.
- In the subset of annual exotic plants, forbs decreased slightly more than graminoids, but in the overall exotic dataset, graminoids decreased more than forbs, and to a larger degree.
 - C treatment had no significant effect on perennial native plant abundance.

Model selection

- Candidate model included region, dlc (duration since last C application), C_type, cratc (C rate as categorical), plotc (plot size), capt (months applying C), and plant_apgfs (plant lifeform).
- dlc, cratc, region, plotc, and plant_apgfs were important moderators for the exotic dataset (importance value > 0.8).
- dlc, cratc, region, C_type, and plant_apgfs were important moderators for the native dataset (importance value > 0.8).

No moderators (summary effect)

Exotic

Model summary

```
summary(ex.mv.o)
```

```
##
## Multivariate Meta-Analysis Model (k = 486; method: REML)
##
##
      logLik
               Deviance
                               AIC
                                          BIC
                                                     AICc
   -690.7916
             1381.5831
                        1387.5831
                                    1400.1355
                                               1387.6330
##
##
## Variance Components:
##
##
               estim
                        sqrt nlvls
                                     fixed
                                                    factor
## sigma^2.1
              0.3457
                      0.5880
                                 50
                                                    exp ID
                                        no
                                            exp_ID/obs_ID
##
  sigma^2.2
              0.1554
                      0.3942
                                486
                                        no
##
## Test for Heterogeneity:
## Q(df = 485) = 1012.8128, p-val < .0001
##
## Model Results:
##
## estimate
                                         ci.lb
                        zval
                                pval
                                                  ci.ub
                 se
##
   -0.5768 0.1009 -5.7168
                             <.0001
                                      -0.7746
                                               -0.3791
##
##
## Signif. codes: 0 '*** 0.001 '** 0.01 '* 0.05 '.' 0.1 ' 1
```

• C treatment causes significant decrease in exotic plant abundance.

Prediction interval

```
predict(ex.mv.o)
```

```
## pred se ci.lb ci.ub pi.lb pi.ub
## -0.5768 0.1009 -0.7746 -0.3791 -1.9782 0.8246
```

- The prediction interval (pi.lb and pi.ub) measures the distribution of true effects. A prediction of the effect size for any given population would fall within the prediction interval 95% of the time (Borenstein 2019, Research Synthesis Methods).
- The prediction interval can be used to calculate the probability the effect of a new study will be below or above a certain threshold using the left-tail cumulative t-distribution with k-1 degrees of freedom (IntHout 2016, BMJ Open).
 - See Supplementary Appendix Formula 2: we are finding the probability that t-value \leq T, where $T = \frac{D-M}{SD_{PI}}$, and D is the threshold (0), M is the summary effect, and SD_{PI} is the standard deviation of the prediction interval.
 - I am calculating SD_{PI} from the definition of the prediction interval, $PI = Mean \pm 2SD$ (Borenstein 2019, Research Synthesis Methods).
 - After finding T, calculate the cumulative distribution function with k-1 degrees of freedom.
 - * Use the calculator at https://www.danielsoper.com/statcalc/calculator.aspx?id=41 or the function pt().

```
SD_PI <- abs(predict(ex.mv.o)$pi.lb - predict(ex.mv.o)$pred) / 2
t <- (0 - predict(ex.mv.o)$pred) / SD_PI
pt(t, ex.mv.o$k - 1, lower.tail = TRUE)</pre>
```

```
## [1] 0.7946017
```

• There is an 80% chance a new study/randomly selected exotic plant population will decrease in response to C addition.

I^2 statistic

```
## [1] I^2 = 56.4347952512755
```

- I^2 is a measure of relative heterogeneity. It is the proportion of variance in observed effects that is due to variance in true effects (effects in an infinitely large population with no sampling error), or, in other words, the ratio of excess dispersion to total dispersion ($\frac{V_{TRUE}}{V_{OBS}}$, expressed here as a percentage; Borenstein 2009).
- I^2 can be thought of as the inconsistency across studies; an I^2 of 0% means there is no observed variance that is actually due to true variance, and therefore no variance to explain, while an I^2 of 100% means that all the variance could be explained by something besides sampling error, such as the effect of moderators (Borenstein 2019, Research Synthesis Methods).

QM/QT

```
## [1] QM/QT = 0.0312595156674703
```

• QM/QT is the proportion of heterogeneity explained by the model $(\frac{Q_M}{Q_E+Q_M})$, and is "functionally equivalent" to the R^2 of a linear model (Koricheva 2013).

Native

Model summary

```
summary(nt.mv)
##
## Multivariate Meta-Analysis Model (k = 655; method: REML)
##
      logLik
              Deviance
                               AIC
                                          BIC
                                                    AICc
##
##
  -804.1182
            1608.2365
                        1614.2365
                                   1627.6858
                                              1614.2734
##
## Variance Components:
##
                        sqrt nlvls fixed
##
                                                   factor
               estim
## sigma^2.1
             0.1155
                     0.3398
                                 53
                                        nο
                                                   exp_ID
## sigma^2.2
             0.0000 0.0001
                                655
                                        no
                                            exp_ID/obs_ID
##
## Test for Heterogeneity:
## Q(df = 654) = 925.7573, p-val < .0001
##
## Model Results:
##
                                        ci.lb
## estimate
                                                ci.ub
                 se
                        zval
                                pval
##
   -0.0742 0.0592 -1.2543 0.2097
                                     -0.1902 0.0417
##
## ---
## Signif. codes: 0 '*** 0.001 '** 0.01 '* 0.05 '.' 0.1 ' 1
```

• C treatment does not cause a significant change in native plant abundance.

Prediction interval

```
predict(nt.mv)

##

## pred se ci.lb ci.ub pi.lb pi.ub

## -0.0742 0.0592 -0.1902 0.0417 -0.7502 0.6018

SD_PI <- (abs(predict(nt.mv)$pi.lb) - abs(predict(nt.mv)$pred)) / 2

t <- (0 - predict(nt.mv)$pred) / SD_PI

pt(t, nt.mv$k - 1, lower.tail = TRUE)

## [1] 0.5868547

1 - pt(t, nt.mv$k - 1, lower.tail = TRUE)</pre>
```

[1] 0.4131453

• There is a 59% chance a new study or randomly selected population of native plants will decrease in response to C addition (or a 41% chance they will increase).

I^2 statistic

[1] $I^2 = 20.5377724218779$

• A low I^2 indicates most of the variance is due to randomness and cannot be explained with moderators. Therefore, the following native single moderator models should be interpreted with caution.

QM/QT

[1] QM/QT = 0.00169661166339359

Single moderator models

Duration since first C app (dfc)

Duration since first (exotic) Estimate [95% CI] Overall (n = 486)-0.58 [-0.77, -0.38] central Canada (n = 14) -0.06 [-0.67, 0.54] coastal CA, USA (n = 55) -0.14 [-0.58, 0.30] Great Basin, USA (n = 101) -0.39 [-0.85, 0.08] Great Plains, USA (n = 36) -0.41 [-0.91, 0.09] Midwest, USA (n = 136)-0.58 [-1.17, 0.01] northeast USA (n = 18)0.25 [-0.74, 1.24] Pacific Northwest, USA (n = 48) -0.74 [-1.13, -0.35] Rocky Mountains, USA (n = 9)-0.47 [-1.06, 0.12] South Africa (n = 2)0.17 [-1.23, 1.56] southeast Australia (n = 55) -1.39 [-1.87, -0.91] southwest USA (n = 12)-1.47 [-2.24, -0.70] -3 -2 -1 0 2 Observed Outcome

Duration since first (native) Estimate [95% CI] Overall (n = 655)-0.07 [-0.19, 0.04] 3 (n = 5)0.10 [-0.59, 0.78] 5-6 (n = 5)-0.53 [-1.11, 0.04] 7-12 (n = 67)0.12 [-0.24, 0.48] 13-18 (n = 99)-0.27 [-0.58, 0.03] 19-24 (n = 67)0.29 [-0.06, 0.64] 25-36 (n = 73)-0.19 [-0.47, 0.08] 37-50 (n = 143)0.00 [-0.26, 0.26] 100-200 (n = 9)0.06 [-0.75, 0.88] >200 (n = 187)-0.21 [-0.43, 0.01] -1.5 -1 -0.50 0.5 1 **Observed Outcome**

Duration since last C app (dlc)

Duration since last (native)

Duration (exotic) Estimate [95% CI] Overall (n = 487)-0.60 [-0.80, -0.40] 3 (n = 88)-1.47 [-1.95, -1.00] -0.27 [-0.97, 0.43] 5-6 (n = 11)7-12 (n = 127)-0.72 [-1.26, -0.18] 13-18 (n = 104)-0.85 [-1.29, -0.42] 19-24 (n = 22)-0.33 [-0.86, 0.19] 25-36 (n = 11)-1.01 [-1.75, -0.26] 37-50 (n = 101)-0.28 [-0.72, 0.15] 100-200 (n = 6)0.01 [-0.99, 1.01] >200 (n = 17)-0.56 [-1.20, 0.08] -2 0 1 2 -1

Observed Outcome

Estimate [95% CI]

Overall (n = 655)-0.07 [-0.19, 0.04] 0-1.5 (n = 127) -0.40 [-0.65, -0.16] 2 (n = 78)0.14 [-0.15, 0.42] 3-3.5 (n = 10)-0.06 [-0.54, 0.41] 4-6 (n = 35)-0.40 [-0.73, -0.08] 7-12 (n = 14)-0.19 [-0.57, 0.19] 13-18 (n = 294)-0.01 [-0.23, 0.21] 19-24 (n = 64)0.33 [-0.04, 0.70] 36-49 (n = 20)0.22 [-0.24, 0.68] >100 (n = 13) 0.17 [-0.45, 0.79]

0

Observed Outcome

0.5

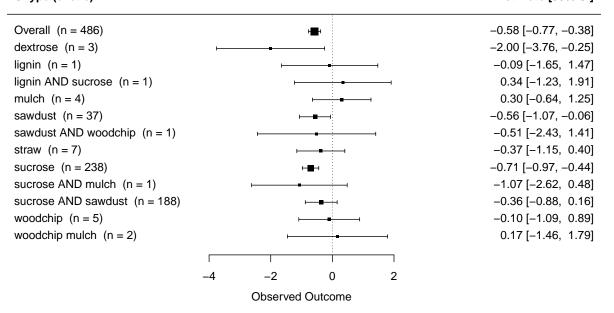
-0.5

-1

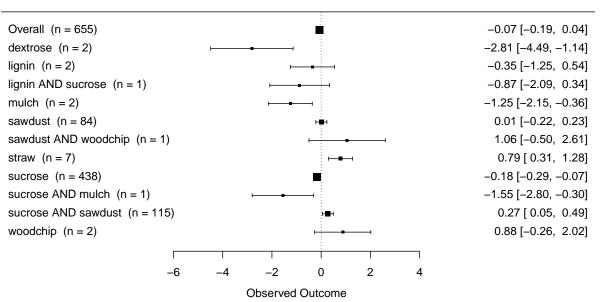
C type

C type (exotic)

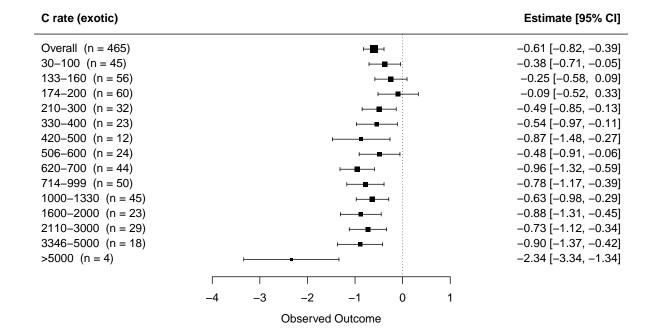
Estimate [95% CI]

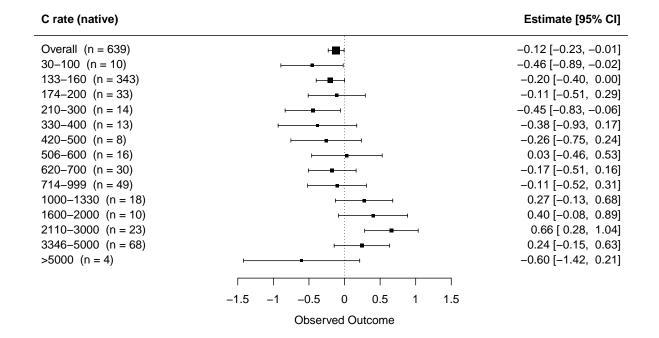


C type (native)



C rate (cratc)

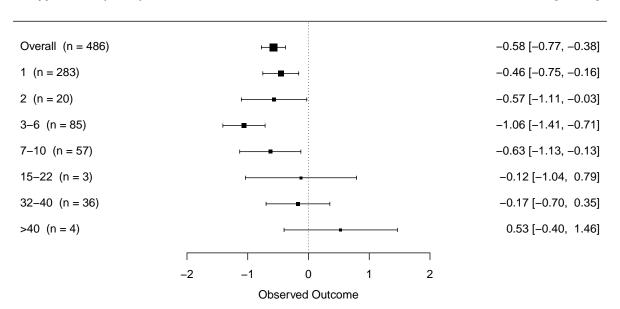




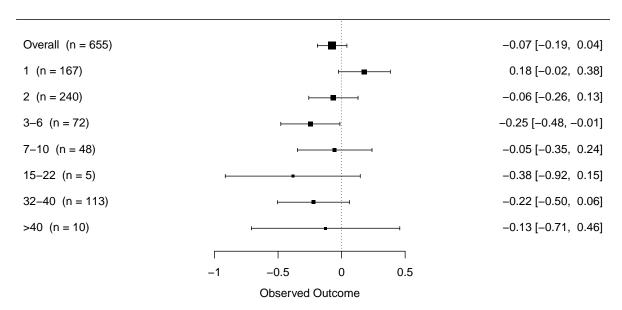
Total C applications

C applications (exotic)

Estimate [95% CI]



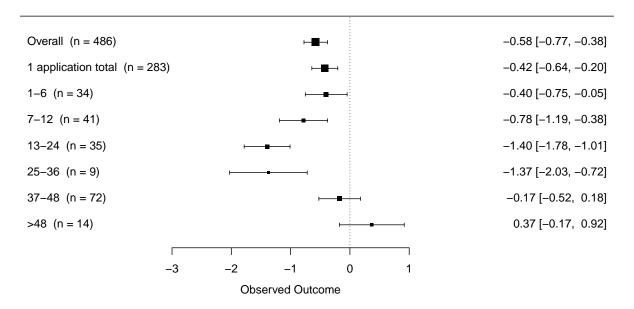
C applications (native)



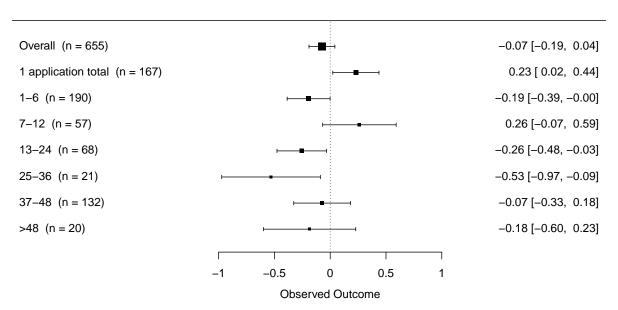
Months applying C (capt)

Months applying C (exotic)

Estimate [95% CI]



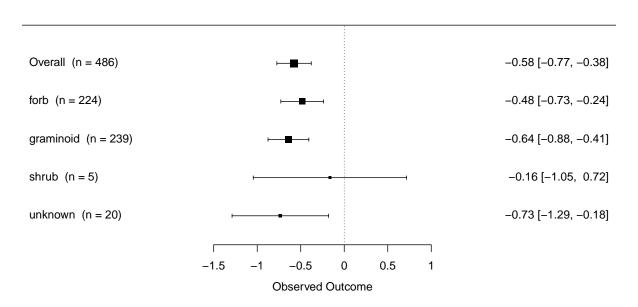
Months applying C (native)



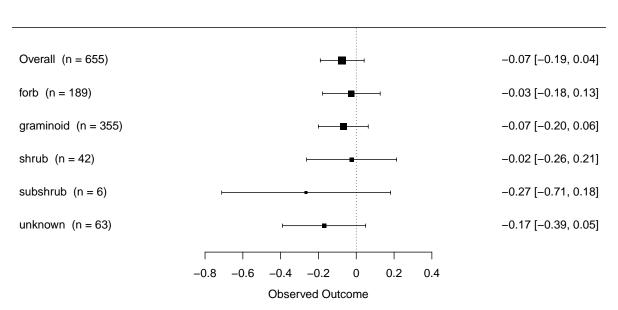
Grass/forb/shrub

Grass/forb/shrub (exotic)

Estimate [95% CI]



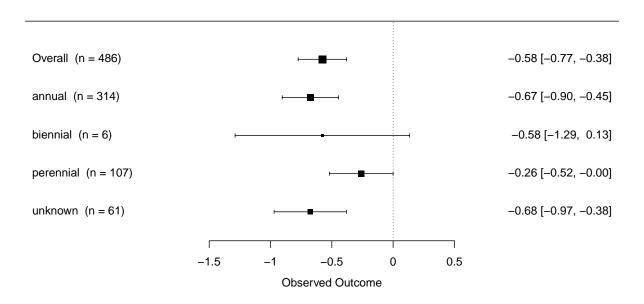
Grass/forb/shrub (native)



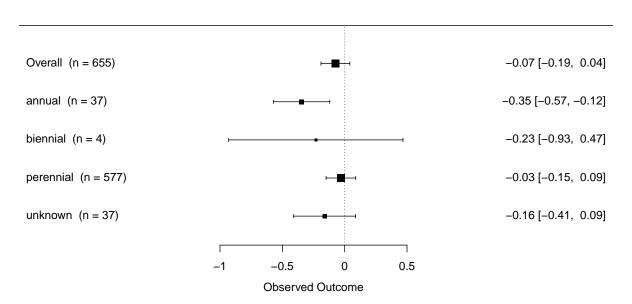
Annual/perennial

Annual/perennial (exotic)

Estimate [95% CI]



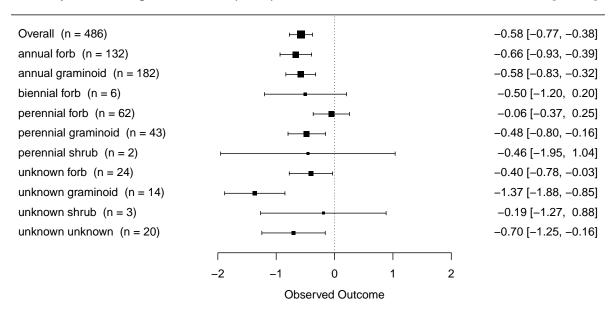
Annual/perennial (native)



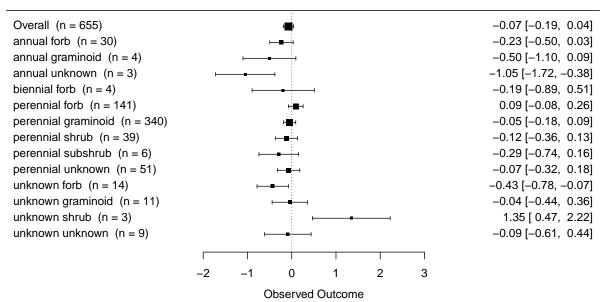
Annual/perennial and grass/forb/shrub

Annual/perennial and grass/forb/shrub (exotic)

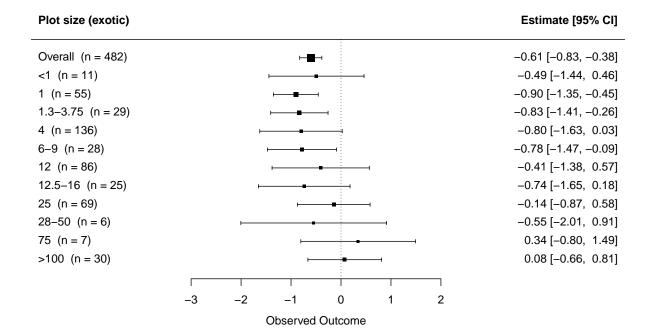
Estimate [95% CI]

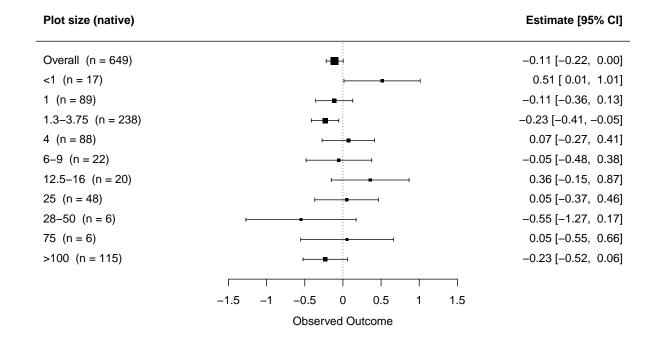


Annual/perennial and grass/forb/shrub (native)



Plot size

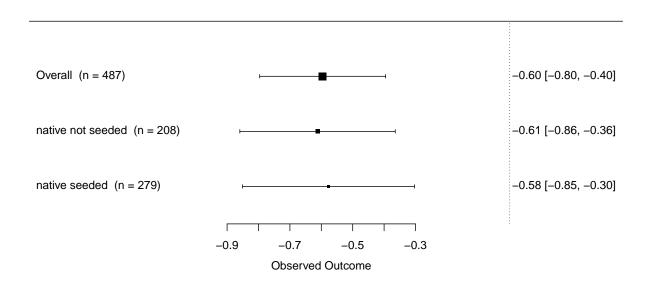




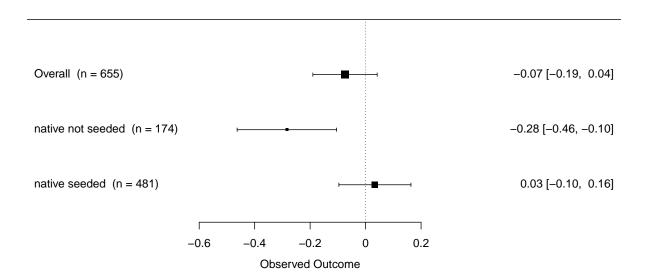
Seeding of native

Seeding of native (exotic)

Estimate [95% CI]



Seeding of native (native)



Heterogeneity stats

Model	QE	QE_df	QE_p	QM	QM_df	QM_p	QM_QT
Summary							
Exotic	1013.0	485	0.0e + 00	32.680	1	0.0000000	0.031260
Native	925.8	654	0.0e + 00	1.573	1	0.2097000	0.001697
Region							
Exotic	871.4	485	0.0e + 00	71.990	11	0.0000000	0.076320
Native	841.8	654	3.0e-07	32.480	10	0.0003331	0.037150
Duration since first C app							
Exotic	959.4	486	0.0e + 00	60.610	9	0.0000000	0.059420
Native	884.8	654	0.0e + 00	13.620	9	0.1364000	0.015160
Duration					-		
Exotic	954.6	487	0.0e + 00	64.110	9	0.0000000	0.062940
Native	883.2	654	0.0e + 00	23.030	9	0.0061220	0.025420
	000.2	001	0.00 00	20.000		0.0001220	0.020120
C type Exotic	948.4	487	0.0e + 00	42.870	12	0.0000238	0.043240
Native	814.0	654	5.6e-06	56.400	11	0.0000238	0.043240 0.064800
	014.0	004	5.06-00	30.400	11	0.0000000	0.004800
Crate	070.0	404	0.0 +00	70.000	1.4	0.0000000	0.076540
Exotic	879.8	464 638	0.0e+00	72.930	14	0.0000000	0.076540
Native	799.4		2.6e-06	41.870	14	0.0001297	0.049770
Total C applications							
Exotic	956.7	487	0.0e + 00	49.720	7	0.0000000	0.049400
Native	875.6	654	0.0e + 00	12.450	7	0.0867700	0.014020
Months applying C							
Exotic	890.9	487	0.0e + 00	98.500	7	0.0000000	0.099550
Native	866.5	654	0.0e+00	19.860	7	0.0058790	0.022410
Grass/forb/shrub							
Exotic	1003.0	487	0.0e + 00	48.880	4	0.0000000	0.046450
Native	909.1	654	0.0e + 00	10.630	4	0.0310600	0.011560
Annual/perennial							
Exotic	1034.0	487	0.0e + 00	33.070	4	0.0000012	0.030980
Native	918.8	654	0.0e + 00	4.090	5	0.5365000	0.004432
Annual/perennial and grass/forb/shrub							
Exotic	973.9	487	0.0e+00	68.770	10	0.0000000	0.065950
Native	881.2	654	0.0e + 00	37.230	13	0.0003817	0.040530
Plot size							
Exotic	937.3	481	0.0e + 00	37.840	11	0.0000833	0.038800
Native	824.1	648	9.0e-07	18.570	10	0.0461400	0.022030
	-		0.00 01	10.010	10	5.0 151 100	0.022000
Seeding of Exotic	or native 1005.0	486	0.0e+00	33.420	2	0.0000001	0.032190
Native	894.7	$\frac{480}{654}$	0.0e+00 0.0e+00	10.240	2	0.0000001 0.0059620	0.032190
native	034.1	004	0.06+00	10.240		0.00038020	0.011320

Model selection

Exotic

```
## model aicc weights ## 1 yi ~ 1 + dlc + plotc + region + plant_apgfs 1286.618 0.9369876
```

• The top model has a weight of 94%.

Native

```
## model aicc weights ## 1 yi ~ 1 + dlc + cratc + region + C_type + plant_apgfs 1503.092 0.9702438
```

• The top model has a weight of 97%.