# Temp-Nut Plots

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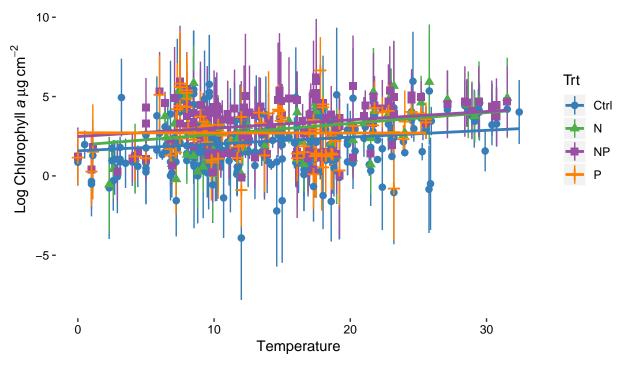
1. ANCOVAs testing the effect of temperature on nutrient treatment (raw chlorophyll a) and nutrient LRRs

Lsmeans tests for significant differences among treatments, averaging over temperature

Lstrends tests for differences among the relationships between treatments and temperature

a. Temperature and raw Chl-a

## Warning: Removed 556 rows containing missing values (geom\_pointrange).



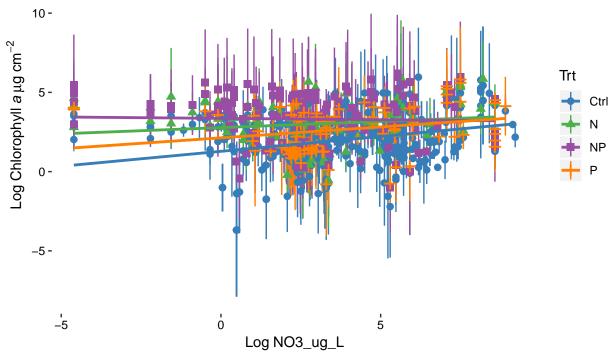
```
#Chla vs. Temp ANCOVAs
options(contrasts=c("contr.sum", "contr.poly"))
Model=aov(log(Chla_mean+.01)~Trt*Temp, data=Data)
Anova(Model, type="III")
## Anova Table (Type III tests)
##
## Response: log(Chla_mean + 0.01)
                Sum Sq Df F value
                                       Pr(>F)
## (Intercept)
               449.19
                         1 193.7091 < 2.2e-16 ***
                 26.10
## Trt
                             3.7519
                                      0.01082 *
                 38.90
                         1 16.7768 4.696e-05 ***
## Temp
## Trt:Temp
                 10.40
                         3
                             1.4954
                                      0.21453
## Residuals
               1623.22 700
## Signif. codes: 0 '***' 0.001 '**' 0.05 '.' 0.1 ' ' 1
#Log meets assumptions
\#par(mfrow=c(2,2))
#plot(Model)
#Lsmeans (pairwise trt differences, averaging over temperature)
lsmeans(Model, pairwise~Trt)
```

```
## NOTE: Results may be misleading due to involvement in interactions
## $1smeans
## Trt
                         SE df lower.CL upper.CL
          lsmean
## Ctrl 2.190645 0.08051639 700 2.032562 2.348727
## N
        2.907072 0.15261647 700 2.607431 3.206713
        3.227157 0.12174516 700 2.988127 3.466186
## NP
## P
        2.707094 0.15826899 700 2.396355 3.017833
##
## Results are given on the log (not the response) scale.
## Confidence level used: 0.95
##
## $contrasts
## contrast
               estimate
                               SE df t.ratio p.value
## Ctrl - N -0.7164270 0.1725534 700
                                       -4.152 0.0002
                                      -7.101 <.0001
## Ctrl - NP -1.0365117 0.1459616 700
## Ctrl - P -0.5164492 0.1775724 700
                                      -2.908 0.0196
## N - NP
             -0.3200846 0.1952272 700
                                      -1.640 0.3571
## N - P
              0.1999779 0.2198655 700
                                       0.910 0.7998
## NP - P
              0.5200625 0.1996771 700
                                        2.605 0.0463
## Results are given on the log (not the response) scale.
## P value adjustment: tukey method for comparing a family of 4 estimates
#Lstrends (pairwise trt differences in the way chl-a responds to temperature)
comp=lstrends(Model, ~ Trt, var = "Temp")
cld(comp)
## Trt
           Temp.trend
                              SE df
                                        lower.CL
                                                   upper.CL .group
        -0.0005966077 0.02617721 700 -0.05199186 0.05079864 1
## Ctrl 0.0431581478 0.01168599 700 0.02021436 0.06610193 1
## NP
         0.0524759991 0.01826102 700 0.01662306 0.08832894 1
         0.0685480396 0.02097145 700 0.02737357 0.10972251 1
## N
##
## Trends are based on the log (transformed) scale
## Confidence level used: 0.95
## P value adjustment: tukey method for comparing a family of 4 estimates
## significance level used: alpha = 0.05
```

### b.) Temperature and Nutrient LRRs

```
## Warning: Removed 914 rows containing non-finite values (stat_smooth).
## Warning: Removed 914 rows containing missing values (geom_pointrange).
    5.0 -
    2.5 -
                                                                                 Trt
                                                                                     Ctrl
                                                                                     NP
   -2.5 -
   -5.0 -
          ó
                              10
                                                                     30
                                                 20
                                     Temperature
#Chla vs. Temp ANCOVAs
options(contrasts=c("contr.sum", "contr.poly"))
Model=aov(LRR_Mean~Trt*Temp, data=Data)
Anova(Model, type="III")
## Anova Table (Type III tests)
## Response: LRR_Mean
                Sum Sq Df F value
                                       Pr(>F)
                         1 138.7054 < 2.2e-16 ***
## (Intercept)
               76.597
## Trt
                 4.697
                             4.2525 0.014983 *
                 4.953
                             8.9687
                                    0.002945 **
## Temp
                         1
## Trt:Temp
                 0.614
                         2
                             0.5561 0.573974
## Residuals
               189.966 344
## Signif. codes: 0 '***' 0.001 '**' 0.05 '.' 0.1 ' ' 1
#Doesn't quite meet assumptions
\#par(mfrow=c(2,2))
#plot(Model)
#Lsmeans (pairwise trt differences, averaging over temperature)
```

```
lsmeans(Model, pairwise~Trt)
## NOTE: Results may be misleading due to involvement in interactions
## $1smeans
## Trt
          lsmean
                         SE df lower.CL upper.CL
       0.7389219 0.07436612 344 0.5926523 0.8851914
## NP 1.1303873 0.05933107 344 1.0136900 1.2470846
        0.7687583 0.07745849 344 0.6164064 0.9211101
## P
##
## Confidence level used: 0.95
##
## $contrasts
## contrast
                estimate
                                 SE df t.ratio p.value
## N - NP
           -0.39146542 0.09513409 344 -4.115 0.0001
            -0.02983642 0.10737847 344 -0.278 0.9583
## N - P
## NP - P
             0.36162900 0.09757045 344
                                         3.706 0.0007
##
## P value adjustment: tukey method for comparing a family of 3 estimates
#Lstrends (pairwise trt differences in the way chl-a responds to temperature)
comp=lstrends(Model, ~ Trt, var = "Temp")
cld(comp)
## Trt Temp.trend
                             SE df
                                      lower.CL
                                                   upper.CL .group
## NP -0.02641861 0.008911366 344 -0.04394624 -0.008890989 1
        -0.01690078 0.012774458 344 -0.04202666 0.008225095
## P
## N
        -0.01249417 0.010234051 344 -0.03262336 0.007635018 1
##
## Confidence level used: 0.95
## P value adjustment: tukey method for comparing a family of 3 estimates
## significance level used: alpha = 0.05
c.) NO3_ug_L and raw Chl-a
\#Chla\ vs.\ NO3\_uq\_L
ggplot(Data, aes(x=log(NO3_ug_L+.01), y=log(Chla_mean+.01), color=Trt)) +
  geom_pointrange(aes(ymin=log(Chla_mean+.01)-log(Chla_SE+.01), ymax=log(Chla_mean+.01)+ log(Chla_SE+.0
  geom_smooth(method=lm,
               se=FALSE)+
    \#scale\ colour\ hue(l=50)\ +
  theme(axis.text.x = element_text(colour="black"), panel.grid.minor = element_blank(), panel.grid.majo
  ylab(bquote(Log~Chlorophyll~italic(a)~mu*g*~cm^-2))+
  xlab(bquote(Log~NO3_ug_L))+
  scale colour manual(values=c("#377eb8", "#4daf4a", "#984ea3", "#ff7f00"))
## Warning: Removed 443 rows containing non-finite values (stat_smooth).
## Warning: Removed 443 rows containing missing values (geom_pointrange).
```



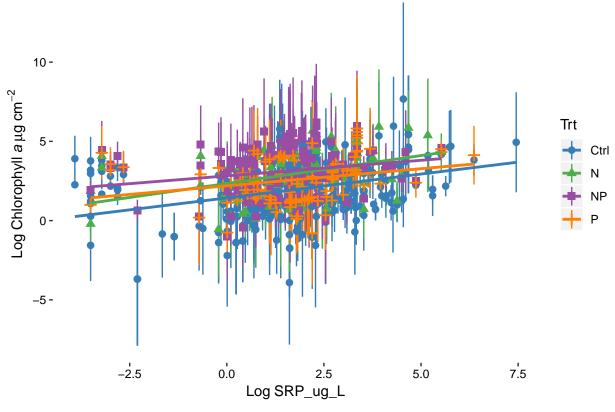
```
#Chla vs. NO3_ug_L ANCOVAs
options(contrasts=c("contr.sum", "contr.poly"))
Model=aov(log(Chla_mean+.01)~Trt*log(NO3_ug_L+.01), data=Data)
Anova(Model, type="III")
## Anova Table (Type III tests)
##
## Response: log(Chla_mean + 0.01)
                             Sum Sq Df F value
                                                   Pr(>F)
## (Intercept)
                            1274.42
                                     1 547.823 < 2.2e-16 ***
## Trt
                             241.74
                                      3 34.638 < 2.2e-16 ***
## log(NO3_ug_L + 0.01)
                              40.57
                                        17.440 3.285e-05 ***
## Trt:log(NO3_ug_L + 0.01)
                              39.77
                                      3
                                          5.699 0.0007305 ***
## Residuals
                            1891.31 813
## ---
## Signif. codes: 0 '***' 0.001 '**' 0.05 '.' 0.1 ' ' 1
#Log meets assumptions pretty well
\#par(mfrow=c(2,2))
#plot(Model)
#Lsmeans (pairwise trt differences, averaging over NO3_ug_Lerature)
lsmeans(Model, pairwise~Trt)
```

```
## NOTE: Results may be misleading due to involvement in interactions
## $1smeans
## Trt
                         SE df lower.CL upper.CL
          lsmean
## Ctrl 2.440884 0.09955109 813 2.245477 2.636292
## N
        3.258676 0.20588527 813 2.854547 3.662805
        3.251149 0.17893846 813 2.899913 3.602385
## NP
##
  Р
        2.976425 0.20153330 813 2.580838 3.372012
##
## Results are given on the log (not the response) scale.
## Confidence level used: 0.95
##
## $contrasts
## contrast
                 estimate
                                 SE df t.ratio p.value
## Ctrl - N -0.817791937 0.2286901 813
                                        -3.576 0.0021
                                        -3.957 0.0005
## Ctrl - NP -0.810264889 0.2047667 813
## Ctrl - P -0.535541310 0.2247801 813
                                        -2.383 0.0812
## N - NP
              0.007527048 0.2727778 813
                                          0.028 1.0000
## N - P
              0.282250628 0.2881049 813
                                          0.980 0.7611
## NP - P
              0.274723580 0.2695081 813
                                          1.019 0.7382
## Results are given on the log (not the response) scale.
## P value adjustment: tukey method for comparing a family of 4 estimates
#Lstrends (pairwise trt differences in the way chl-a responds to NO3_ug_Lerature)
comp=lstrends(Model, ~ Trt, var = "NO3_ug_L")
cld(comp)
   Trt NO3_ug_L.trend
##
                                 SE df
                                             lower.CL
                                                          upper.CL .group
## NP
         -3.375578e-05 8.180243e-05 813 -1.943246e-04 0.0001268131 1
## N
          1.511512e-04 9.443989e-05 813 -3.422353e-05 0.0003365260 12
          2.651905e-04 1.153585e-04 813 3.875493e-05 0.0004916262 12
## P
          3.638642e-04 5.504633e-05 813 2.558145e-04 0.0004719138
## Ctrl
##
## Trends are based on the log (transformed) scale
## Confidence level used: 0.95
## P value adjustment: tukey method for comparing a family of 4 estimates
## significance level used: alpha = 0.05
```

### d.) NO3\_ug\_L and Nutrient LRRs

```
## Warning: Removed 866 rows containing non-finite values (stat_smooth).
## Warning: Removed 867 rows containing missing values (geom_pointrange).
    5.0 -
    2.5 -
                                                                                Trt
                                                                                    Ctrl
O.0 -
   -2.5 -
   -5.0 -
        -5
                                log(NO3 ug L+0.01)
#Chla vs. NO3_ug_L ANCOVAs
options(contrasts=c("contr.sum", "contr.poly"))
Model=aov(LRR_Mean~Trt*NO3_ug_L, data=Data)
Anova(Model, type="III")
## Anova Table (Type III tests)
## Response: LRR_Mean
                 Sum Sq Df F value
                                       Pr(>F)
                        1 642.8127 < 2.2e-16 ***
## (Intercept) 279.550
                             21.1878 1.83e-09 ***
## Trt
                 18.429
                 5.028
## NO3_ug_L
                          1 11.5615 0.0007425 ***
## Trt:NO3_ug_L
                 1.216
                         2
                              1.3976 0.2484117
## Residuals
               170.475 392
## Signif. codes: 0 '***' 0.001 '**' 0.05 '.' 0.1 ' ' 1
#Doesn't really meet assumptions- outliers?
\#par(mfrow=c(2,2))
#plot(Model)
#Lsmeans (pairwise trt differences, averaging over NO3_uq_Lerature)
```

```
lsmeans(Model, pairwise~Trt)
## NOTE: Results may be misleading due to involvement in interactions
## $1smeans
## Trt
          lsmean
                         SE df lower.CL upper.CL
##
       0.7767926 0.06019105 392 0.6584549 0.8951302
## NP 1.1714382 0.04966521 392 1.0737947 1.2690817
       0.6967082 0.06665349 392 0.5656652 0.8277512
## P
##
## Confidence level used: 0.95
##
## $contrasts
## contrast
                estimate
                                 SE df t.ratio p.value
## N - NP
           -0.39464563 0.07803586 392 -5.057 <.0001
             0.08008437 0.08980897 392
## N - P
                                          0.892 0.6459
## NP - P
             0.47473000 0.08312232 392
                                         5.711 <.0001
##
## P value adjustment: tukey method for comparing a family of 3 estimates
#Lstrends (pairwise trt differences in the way chl-a responds to NO3_ug_Lerature)
comp=lstrends(Model, ~ Trt, var = "NO3_ug_L")
cld(comp)
## Trt NO3 ug L.trend
                                 SE df
                                            lower.CL
                                                           upper.CL .group
        -1.455899e-04 4.649224e-05 392 -0.0002369953 -5.418462e-05 1
         -1.301904e-04 6.747370e-05 392 -0.0002628460 2.465163e-06 1
## N
## P
         -4.230766e-05 4.513564e-05 392 -0.0001310459 4.643055e-05 1
##
## Confidence level used: 0.95
## P value adjustment: tukey method for comparing a family of 3 estimates
## significance level used: alpha = 0.05
e.) SRP_ug_L and raw Chl-a
#Chla vs. SRP_uq_L
ggplot(Data, aes(x=log(SRP_ug_L+.01), y=log(Chla_mean+.01), color=Trt)) +
  geom_pointrange(aes(ymin=log(Chla_mean+.01)-log(Chla_SE+.01), ymax=log(Chla_mean+.01)+ log(Chla_SE+.0
  geom_smooth(method=lm,
               se=FALSE)+
    \#scale\ colour\ hue(l=50)\ +
  theme(axis.text.x = element_text(colour="black"), panel.grid.minor = element_blank(), panel.grid.majo
  ylab(bquote(Log~Chlorophyll~italic(a)~mu*g*~cm^-2))+
  xlab(bquote(Log~SRP_ug_L))+
  scale colour manual(values=c("#377eb8", "#4daf4a", "#984ea3", "#ff7f00"))
## Warning: Removed 421 rows containing non-finite values (stat_smooth).
## Warning: Removed 421 rows containing missing values (geom_pointrange).
```



```
#Chla vs. SRP_ug_L ANCOVAs
options(contrasts=c("contr.sum", "contr.poly"))
Model=aov(log(Chla_mean+.01)~Trt*log(SRP_ug_L+.01), data=Data)
Anova(Model, type="III")
## Anova Table (Type III tests)
##
## Response: log(Chla_mean + 0.01)
                             Sum Sq Df F value
                                                    Pr(>F)
## (Intercept)
                            1419.27
                                      1 582.8538 < 2.2e-16 ***
## Trt
                             139.42
                                         19.0853 5.610e-12 ***
## log(SRP_ug_L + 0.01)
                             114.88
                                         47.1774 1.267e-11 ***
                                      1
## Trt:log(SRP_ug_L + 0.01)
                               6.08
                                      3
                                          0.8325
                                                    0.4761
## Residuals
                            2033.26 835
## ---
## Signif. codes: 0 '***' 0.001 '**' 0.05 '.' 0.1 ' ' 1
#Log meets assumptions
\#par(mfrow=c(2,2))
#plot(Model)
#Lsmeans (pairwise trt differences, averaging over SRP_ug_Lerature)
lsmeans(Model, pairwise~Trt)
```

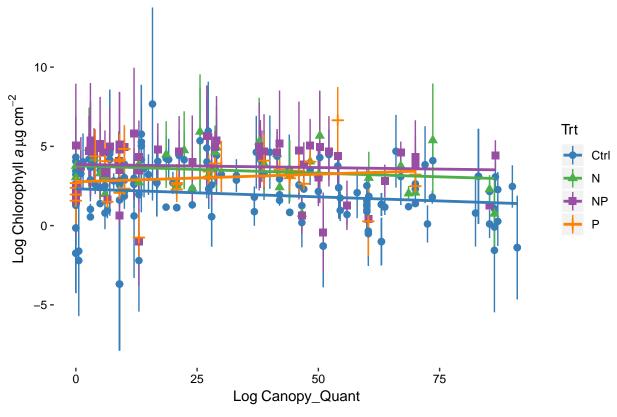
```
## NOTE: Results may be misleading due to involvement in interactions
## $1smeans
## Trt
                         SE df lower.CL upper.CL
          lsmean
## Ctrl 2.337342 0.09151571 835 2.157715 2.516970
## N
        3.401943 0.16910997 835 3.070012 3.733873
        3.413292 0.17219052 835 3.075315 3.751269
## NP
## P
        2.844303 0.19081731 835 2.469765 3.218841
##
## Results are given on the log (not the response) scale.
## Confidence level used: 0.95
##
## $contrasts
## contrast
                estimate
                                SE df t.ratio p.value
## Ctrl - N -1.06460025 0.1922844 835
                                       -5.537 <.0001
## Ctrl - NP -1.07594948 0.1949992 835
                                        -5.518 <.0001
                                        -2.396 0.0787
## Ctrl - P -0.50696019 0.2116279 835
## N - NP
             -0.01134923 0.2413457 835
                                        -0.047
                                               1.0000
## N - P
              0.55764006 0.2549695 835
                                         2.187 0.1276
## NP - P
              0.56898930 0.2570230 835
                                         2.214 0.1203
## Results are given on the log (not the response) scale.
## P value adjustment: tukey method for comparing a family of 4 estimates
#Lstrends (pairwise trt differences in the way chl-a responds to SRP_ug_Lerature)
comp=lstrends(Model, ~ Trt, var = "SRP_ug_L")
cld(comp)
  Trt SRP_ug_L.trend
##
                                SE df
                                          lower.CL
                                                     upper.CL .group
## NP
           0.007035297 0.002946108 835 0.001252649 0.01281794
## P
           0.007739924 0.003267081 835 0.001327266 0.01415258 1
           0.010779502 0.001447751 835 0.007937844 0.01362116 1
## Ctrl
           0.012622325 0.003073171 835 0.006590277 0.01865437 1
## N
##
## Trends are based on the log (transformed) scale
## Confidence level used: 0.95
## P value adjustment: tukey method for comparing a family of 4 estimates
## significance level used: alpha = 0.05
```

### f.) SRP\_ug\_L and Nutrient LRRs

```
## Warning: Removed 855 rows containing non-finite values (stat_smooth).
## Warning: Removed 856 rows containing missing values (geom_pointrange).
    5.0 -
    2.5 -
                                                                                  Trt
                                                                                      Ctrl
- 0.0 -
                                                                                      Ν
                                                                                      NP
   -2.5 -
   -5.0 -
                 -2.5
                                0.0
                                              2.5
                                                            5.0
                                                                          7.5
                                log(SRP_ug_L + 0.01)
#Chla vs. SRP_ug_L ANCOVAs
options(contrasts=c("contr.sum", "contr.poly"))
Model=aov(LRR_Mean~Trt*log(SRP_ug_L+.01), data=Data)
Anova(Model, type="III")
## Anova Table (Type III tests)
## Response: LRR_Mean
                             Sum Sq Df F value
                                                    Pr(>F)
## (Intercept)
                                      1 249.6849 < 2.2e-16 ***
                            180.877
## Trt
                             10.990
                                      2
                                          7.5853 0.0005834 ***
## log(SRP_ug_L + 0.01)
                              0.290
                                          0.4003 0.5272902
                                      1
## Trt:log(SRP_ug_L + 0.01)
                              0.982
                                      2
                                          0.6781 0.5081671
## Residuals
                            293.390 405
## Signif. codes: 0 '***' 0.001 '**' 0.05 '.' 0.1 ' ' 1
#Doesn't really meet assumptions- outliers?
\#par(mfrow=c(2,2))
#plot(Model)
#Lsmeans (pairwise trt differences, averaging over SRP_ug_Lerature)
```

```
lsmeans(Model, pairwise~Trt)
## NOTE: Results may be misleading due to involvement in interactions
## $1smeans
## Trt
          lsmean
                         SE df lower.CL upper.CL
       0.8957124 0.08653870 405 0.7255913 1.065834
## NP 1.1603460 0.08678302 405 0.9897446 1.330947
       0.8123530 0.09788279 405 0.6199312 1.004775
## P
##
## Confidence level used: 0.95
##
## $contrasts
## contrast
                estimate
                                SE df t.ratio p.value
## N - NP
           -0.26463361 0.1225571 405 -2.159 0.0796
             0.08335938 0.1306522 405
## N - P
                                       0.638 0.7992
## NP - P
             0.34799299 0.1308141 405
                                       2.660 0.0221
##
## P value adjustment: tukey method for comparing a family of 3 estimates
#Lstrends (pairwise trt differences in the way chl-a responds to SRP_ug_Lerature)
comp=lstrends(Model, ~ Trt, var = "SRP_ug_L")
cld(comp)
## Trt SRP ug L.trend
                                SE df
                                          lower.CL
                                                       upper.CL .group
         -0.0031507109 0.002374742 405 -0.007819071 0.001517649 1
         -0.0002610640 0.002633466 405 -0.005438034 0.004915906
## P
## N
         0.0006749867 0.002477163 405 -0.004194716 0.005544689 1
##
## Confidence level used: 0.95
## P value adjustment: tukey method for comparing a family of 3 estimates
## significance level used: alpha = 0.05
g.) Canopy_Quant and raw Chl-a
#Chla vs. Canopy_Quant
ggplot(Data, aes(x=Canopy_Quant, y=log(Chla_mean+.01), color=Trt)) +
  geom_pointrange(aes(ymin=log(Chla_mean+.01)-log(Chla_SE+.01), ymax=log(Chla_mean+.01)+ log(Chla_SE+.0
  geom_smooth(method=lm,
               se=FALSE)+
    \#scale\ colour\ hue(l=50)\ +
  theme(axis.text.x = element_text(colour="black"), panel.grid.minor = element_blank(), panel.grid.majo
  ylab(bquote(Log~Chlorophyll~italic(a)~mu*g*~cm^-2))+
  xlab(bquote(Log~Canopy_Quant))+
  scale colour manual(values=c("#377eb8", "#4daf4a", "#984ea3", "#ff7f00"))
```

## Warning: Removed 944 rows containing non-finite values (stat\_smooth).
## Warning: Removed 944 rows containing missing values (geom\_pointrange).



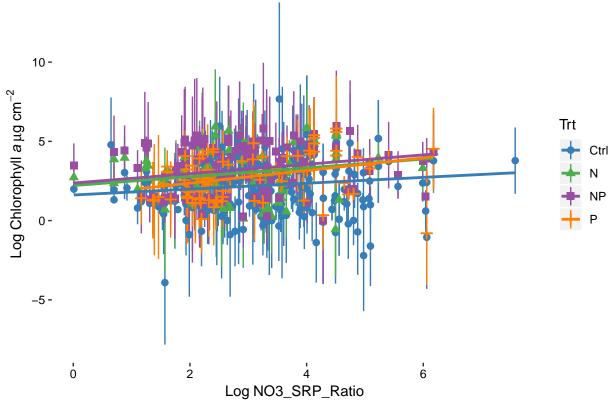
```
#Chla vs. Canopy_Quant ANCOVAs
options(contrasts=c("contr.sum", "contr.poly"))
Model=aov(log(Chla_mean+.01)~Trt*Canopy_Quant, data=Data)
Anova(Model, type="III")
## Anova Table (Type III tests)
##
## Response: log(Chla_mean + 0.01)
                     Sum Sq Df F value
                                           Pr(>F)
## (Intercept)
                     791.80
                              1 246.8310 < 2.2e-16 ***
                                  6.2805 0.0003772 ***
## Trt
                     60.44
                       1.29
                                  0.4022 0.5264201
## Canopy_Quant
                             1
## Trt:Canopy_Quant
                       5.75
                                  0.5978 0.6168920
## Residuals
                    1000.85 312
## Signif. codes: 0 '***' 0.001 '**' 0.05 '.' 0.1 ' ' 1
#Log meets assumptions
\#par(mfrow=c(2,2))
#plot(Model)
#Lsmeans (pairwise trt differences, averaging over Canopy_Quanterature)
lsmeans(Model, pairwise~Trt)
```

```
## NOTE: Results may be misleading due to involvement in interactions
## $1smeans
## Trt
          lsmean
                        SE df lower.CL upper.CL
## Ctrl 1.998297 0.1319976 312 1.738579 2.258015
## N
        3.480799 0.2769059 312 2.935960 4.025638
        3.734079 0.2347833 312 3.272120 4.196038
## NP
##
  Р
        3.070523 0.3203330 312 2.440237 3.700809
##
## Results are given on the log (not the response) scale.
## Confidence level used: 0.95
##
## $contrasts
## contrast
               estimate
                               SE df t.ratio p.value
## Ctrl - N -1.4825024 0.3067576 312
                                       -4.833 <.0001
                                       -6.444 <.0001
## Ctrl - NP -1.7357820 0.2693447 312
## Ctrl - P -1.0722257 0.3464630 312
                                       -3.095 0.0115
## N - NP
             -0.2532796 0.3630428 312 -0.698 0.8979
## N - P
              0.4102767 0.4234266 312
                                       0.969 0.7672
## NP - P
              0.6635563 0.3971605 312
                                        1.671 0.3409
## Results are given on the log (not the response) scale.
## P value adjustment: tukey method for comparing a family of 4 estimates
#Lstrends (pairwise trt differences in the way chl-a responds to Canopy_Quanterature)
comp=lstrends(Model, ~ Trt, var = "Canopy_Quant")
cld(comp)
  Trt Canopy_Quant.trend
                                    SE df
                                              lower.CL
                                                           upper.CL .group
## Ctrl
              -0.010215161 0.004665081 312 -0.01939416 -0.001036163
## N
              -0.009310378 0.011924572 312 -0.03277312 0.014152367
## NP
              -0.003907969 0.009315450 312 -0.02223702 0.014421078 1
               0.009581223 0.015045086 312 -0.02002144 0.039183883 1
## P
##
## Trends are based on the log (transformed) scale
## Confidence level used: 0.95
## P value adjustment: tukey method for comparing a family of 4 estimates
## significance level used: alpha = 0.05
```

### h.) Canopy\_Quant and Nutrient LRRs

```
## Warning: Removed 1129 rows containing non-finite values (stat_smooth).
## Warning: Removed 1130 rows containing missing values (geom_pointrange).
    5.0 -
    2.5 -
                                                                                 Trt
                                                                                      Ctrl
O.0 -
                                                                                      Ν
                                                                                      NP
   -2.5 -
   -5.0 -
          Ó
                           25
                                             50
                                                               75
                              log(Canopy_Quant + 0.01)
#Chla vs. Canopy_Quant ANCOVAs
options(contrasts=c("contr.sum", "contr.poly"))
Model=aov(LRR_Mean~Trt*Canopy_Quant, data=Data)
Anova(Model, type="III")
## Anova Table (Type III tests)
## Response: LRR_Mean
                    Sum Sq Df F value
                                         Pr(>F)
## (Intercept)
                    48.874
                             1 81.5275 2.12e-15 ***
## Trt
                     2.459
                             2 2.0511
                                         0.1328
## Canopy_Quant
                     0.291
                             1 0.4853
                                         0.4873
## Trt:Canopy_Quant 0.548
                             2
                                0.4572
                                         0.6341
## Residuals
                    77.333 129
## Signif. codes: 0 '***' 0.001 '**' 0.05 '.' 0.1 ' ' 1
#Doesn't really meet assumptions- a few outliers
\#par(mfrow=c(2,2))
#plot(Model)
#Lsmeans (pairwise trt differences, averaging over Canopy_Quanterature)
```

```
lsmeans(Model, pairwise~Trt)
## NOTE: Results may be misleading due to involvement in interactions
## $1smeans
## Trt
          lsmean
                        SE df lower.CL upper.CL
       0.9163666 0.1213602 129 0.6762525 1.156481
## NP 1.2624673 0.1001872 129 1.0642445 1.460690
        1.0663704 0.1355395 129 0.7982022 1.334539
## P
##
## Confidence level used: 0.95
##
## $contrasts
## contrast
              estimate
                               SE df t.ratio p.value
## N - NP
           -0.3461008 0.1573714 129 -2.199 0.0751
            -0.1500038 0.1819320 129 -0.825 0.6885
## N - P
## NP - P
             0.1960969 0.1685480 129
                                      1.163 0.4772
##
## P value adjustment: tukey method for comparing a family of 3 estimates
#Lstrends (pairwise trt differences in the way chl-a responds to Canopy_Quanterature)
comp=lstrends(Model, ~ Trt, var = "Canopy_Quant")
cld(comp)
## Trt Canopy_Quant.trend
                                   SE df
                                              lower.CL
                                                           upper.CL .group
             -0.001434254 0.004027030 129 -0.009401832 0.006533324 1
              0.002350881 0.005154942 129 -0.007848299 0.012550061
## N
## P
              0.005509649 0.006503928 129 -0.007358531 0.018377830 1
##
## Confidence level used: 0.95
## P value adjustment: tukey method for comparing a family of 3 estimates
## significance level used: alpha = 0.05
i.) NO3_SRP_Ratio and raw Chl-a
#Chla vs. NO3_SRP_Ratio
ggplot(Data, aes(x=log(NO3_SRP_Ratio+.01), y=log(Chla_mean+.01), color=Trt)) +
  geom_pointrange(aes(ymin=log(Chla_mean+.01)-log(Chla_SE+.01), ymax=log(Chla_mean+.01)+ log(Chla_SE+.0
  geom_smooth(method=lm,
               se=FALSE)+
    \#scale\ colour\ hue(l=50)\ +
  theme(axis.text.x = element_text(colour="black"), panel.grid.minor = element_blank(), panel.grid.majo
  ylab(bquote(Log~Chlorophyll~italic(a)~mu*g*~cm^-2))+
  xlab(bquote(Log~NO3_SRP_Ratio))+
  scale colour manual(values=c("#377eb8", "#4daf4a", "#984ea3", "#ff7f00"))
## Warning: Removed 621 rows containing non-finite values (stat_smooth).
## Warning: Removed 621 rows containing missing values (geom_pointrange).
```



```
#Chla vs. NO3_SRP_Ratio ANCOVAs
options(contrasts=c("contr.sum", "contr.poly"))
Model=aov(log(Chla_mean+.01)~Trt*log(NO3_SRP_Ratio+.01), data=Data)
Anova(Model, type="III")
## Anova Table (Type III tests)
##
## Response: log(Chla_mean + 0.01)
                                  Sum Sq Df F value
                                                         Pr(>F)
## (Intercept)
                                  258.85
                                           1 106.3428 < 2.2e-16 ***
## Trt
                                    9.85
                                               1.3493
                                                         0.2574
## log(NO3_SRP_Ratio + 0.01)
                                              23.6746 1.441e-06 ***
                                   57.63
                                           1
## Trt:log(NO3_SRP_Ratio + 0.01)
                                    5.07
                                           3
                                               0.6942
                                                         0.5558
## Residuals
                                 1545.67 635
## ---
## Signif. codes: 0 '***' 0.001 '**' 0.05 '.' 0.1 ' ' 1
#Log generally meets assumptions
\#par(mfrow=c(2,2))
#plot(Model)
#Lsmeans (pairwise trt differences, averaging over NO3_SRP_Ratioerature)
lsmeans(Model, pairwise~Trt)
```

```
## NOTE: Results may be misleading due to involvement in interactions
## $1smeans
## Trt
                        SE df lower.CL upper.CL
          lsmean
## Ctrl 2.325592 0.1025302 635 2.124253 2.526932
## N
        3.278474 0.2057745 635 2.874393 3.682555
        3.503343 0.1660907 635 3.177189 3.829496
## NP
##
  Р
        3.093732 0.1981893 635 2.704547 3.482918
##
## Results are given on the log (not the response) scale.
## Confidence level used: 0.95
##
## $contrasts
## contrast
               estimate
                               SE df t.ratio p.value
## Ctrl - N -0.9528813 0.2299034 635
                                       -4.145 0.0002
                                       -6.034 <.0001
## Ctrl - NP -1.1777503 0.1951886 635
## Ctrl - P -0.7681399 0.2231399 635
                                       -3.442 0.0034
## N - NP
             -0.2248690 0.2644414 635
                                       -0.850 0.8302
## N - P
              0.1847414 0.2856959 635
                                       0.647 0.9167
## NP - P
              0.4096104 0.2585829 635
                                        1.584 0.3885
## Results are given on the log (not the response) scale.
## P value adjustment: tukey method for comparing a family of 4 estimates
#Lstrends (pairwise trt differences in the way chl-a responds to NO3_SRP_Ratioerature)
comp=lstrends(Model, ~ Trt, var = "NO3_SRP_Ratio")
cld(comp)
  Trt NO3_SRP_Ratio.trend
                                     SE df
                                                lower.CL
                                                            upper.CL .group
## Ctrl
                0.003353242 0.001405632 635 5.929938e-04 0.006113491 1
## N
                0.005035482 0.002519954 635 8.703013e-05 0.009983934
                0.005355182 0.002042649 635 1.344017e-03 0.009366346 1
## NP
                0.007219915 0.002462726 635 2.383843e-03 0.012055987 1
## P
##
## Trends are based on the log (transformed) scale
## Confidence level used: 0.95
## P value adjustment: tukey method for comparing a family of 4 estimates
## significance level used: alpha = 0.05
```

### j.) NO3\_SRP\_Ratio and Nutrient LRRs

```
## Warning: Removed 927 rows containing non-finite values (stat_smooth).
## Warning: Removed 927 rows containing missing values (geom_pointrange).
    5.0 -
    2.5 -
                                                                                  Trt
                                                                                      Ctrl
B 0.0 -
                                                                                      Ν
                                                                                      NP
   -2.5 -
   -5.0 -
          Ó
                                                             6
                             log(NO3 SRP Ratio + 0.01)
#Chla vs. NO3_SRP_Ratio ANCOVAs
options(contrasts=c("contr.sum", "contr.poly"))
Model=aov(LRR_Mean~Trt*log(NO3_SRP_Ratio+.01), data=Data)
Anova(Model, type="III")
## Anova Table (Type III tests)
## Response: LRR_Mean
                                  Sum Sq Df F value
                                                         Pr(>F)
## (Intercept)
                                  59.204
                                           1 74.2606 2.802e-16 ***
## Trt
                                   3.426
                                             2.1484
                                                         0.1183
## log(NO3_SRP_Ratio + 0.01)
                                                         0.1414
                                   1.733
                                            1 2.1735
## Trt:log(NO3_SRP_Ratio + 0.01)
                                   2.143
                                           2
                                              1.3439
                                                         0.2622
## Residuals
                                 265.481 333
## Signif. codes: 0 '***' 0.001 '**' 0.05 '.' 0.1 ' ' 1
\#Doesn't\ quite\ meet\ assumptions-\ outliers?
\#par(mfrow=c(2,2))
#plot(Model)
#Lsmeans (pairwise trt differences, averaging over NO3_SRP_Ratioerature)
```

```
lsmeans(Model, pairwise~Trt)
## NOTE: Results may be misleading due to involvement in interactions
## $1smeans
          lsmean
## Trt
                         SE df lower.CL upper.CL
       0.7162073 0.11128448 333 0.4972981 0.9351165
## NP 1.1571047 0.09023207 333 0.9796080 1.3346014
       0.9040719 0.10810498 333 0.6914171 1.1167266
##
## Confidence level used: 0.95
##
## $contrasts
## contrast
              estimate
                              SE df t.ratio p.value
## N - NP
           -0.4408975 0.1432692 333 -3.077 0.0064
            -0.1878646 0.1551481 333 -1.211 0.4476
## N - P
## NP - P
             0.2530328 0.1408138 333
                                      1.797 0.1722
##
## P value adjustment: tukey method for comparing a family of 3 estimates
#Lstrends (pairwise trt differences in the way chl-a responds to NO3_SRP_Ratioerature)
comp=lstrends(Model, ~ Trt, var = "NO3_SRP_Ratio")
cld(comp)
## Trt NO3 SRP Ratio.trend
                                    SE df
                                               lower.CL
                                                            upper.CL .group
             -0.0034091951 0.001854773 333 -0.007057744 0.0002393541 1
             -0.0017956850 0.001503460 333 -0.004753162 0.0011617917
## NP
              0.0007854361 0.001812651 333 -0.002780254 0.0043511263 1
## P
##
## Confidence level used: 0.95
## P value adjustment: tukey method for comparing a family of 3 estimates
## significance level used: alpha = 0.05
k.) NO3_ug_L and Nutrient LRRs
#LRRs vs. NO3_uq_L
ggplot(Data, aes(x=log(NO3_ug_L+.01), y=LRR_Mean, color=Trt)) +
 geom_pointrange(aes(ymin=LRR_Mean-LRR_Var, ymax=LRR_Mean + LRR_Var, shape=Trt, color=Trt))+
 geom_smooth(method=lm,
               se=FALSE)+
```

```
## Warning: Removed 866 rows containing non-finite values (stat_smooth).
## Warning: Removed 867 rows containing missing values (geom_pointrange).
```

scale colour manual(values=c("#377eb8", "#4daf4a", "#984ea3", "#ff7f00"))

 $\#scale\ colour\ hue(l=50)\ +$ 

xlab(bquote(log(NO3\_ug\_L+.01)))+

ylab(bquote(LRR))+

ylim(-5,5)+

theme(axis.text.x = element\_text(colour="black"), panel.grid.minor = element\_blank(), panel.grid.majo

#plot(Model)

lsmeans(Model, pairwise~Trt)

```
2.5 -
                                                                                 Trt
                                                                                     Ctrl
R 0.0-
                                                                                     Ν
                                                                                     NP
   -2.5 -
   -5.0 -
        -5
                                log(NO3\_ug\_L + 0.01)
#Chla vs. NO3_ug_L ANCOVAs
options(contrasts=c("contr.sum", "contr.poly"))
Model=aov(LRR_Mean~Trt*log(NO3_ug_L+.01), data=Data)
Anova(Model, type="III")
## Anova Table (Type III tests)
##
## Response: LRR_Mean
                             Sum Sq Df F value
                                                    Pr(>F)
## (Intercept)
                            123.947
                                      1 286.3828 < 2.2e-16 ***
                             17.441
                                      2 20.1493 4.682e-09 ***
## log(NO3_ug_L + 0.01)
                              1.234
                                          2.8513 0.092095 .
                                      1
## Trt:log(NO3_ug_L + 0.01)
                              4.094
                                      2
                                          4.7301 0.009336 **
## Residuals
                            169.658 392
## Signif. codes: 0 '***' 0.001 '**' 0.05 '.' 0.1 ' ' 1
#Meets assumptions
\#par(mfrow=c(2,2))
```

#Lsmeans (pairwise trt differences, averaging over NO3\_ug\_Lerature)

```
## NOTE: Results may be misleading due to involvement in interactions
## $1smeans
   Trt
                         SE df lower.CL upper.CL
          lsmean
       0.7136920 0.08700477 392 0.5426377 0.8847464
##
   N
       0.9688516 0.07556051 392 0.8202971 1.1174062
       0.7467103 0.08499691 392 0.5796035 0.9138171
##
##
## Confidence level used: 0.95
##
## $contrasts
## contrast estimate
                              SE df t.ratio p.value
   N - NP -0.2551596 0.1152355 392 -2.214 0.0700
## N - P
            -0.0330183 0.1216318 392 -0.271 0.9602
  NP - P
             0.2221413 0.1137271 392
                                       1.953 0.1254
## P value adjustment: tukey method for comparing a family of 3 estimates
#Lstrends (pairwise trt differences in the way chl-a responds to NO3_uq_Lerature)
comp=lstrends(Model, ~ Trt, var = "NO3_ug_L")
cld(comp)
                                SE df
##
  Trt NO3_ug_L.trend
                                            lower.CL
                                                          upper.CL .group
        -1.496627e-04 4.027377e-05 392 -2.288423e-04 -7.048307e-05
        -5.405734e-05 4.649557e-05 392 -1.454692e-04 3.735454e-05 12
         6.234816e-05 5.679443e-05 392 -4.931163e-05 1.740079e-04
##
##
## Confidence level used: 0.95
## P value adjustment: tukey method for comparing a family of 3 estimates
## significance level used: alpha = 0.05
```

- 2.) Three-way interaction models examine the effects of in-stream NO3, SRP, and temp on chlorophyll a (broken out by treatment) and LRRs (broken out by treatment)
- a.) Temp-Nut Interactions and raw Chl-a

```
##Controls
Ctrl=subset(Data, Trt=="Ctrl")
#Chla vs. NO3_ug_L, SRP, Temp ANCOVAs
options(contrasts=c("contr.sum", "contr.poly"))
Model=aov(log(Chla_mean+.01)~log(NO3_ug_L+.01)*log(SRP_ug_L)*Temp, data=Ctrl)
Anova(Model, type="III")
## Anova Table (Type III tests)
##
## Response: log(Chla_mean + 0.01)
                                           Sum Sq Df F value
                                                                 Pr(>F)
## (Intercept)
                                            4.51
                                                   1 2.8220 0.094237
## log(NO3_ug_L + 0.01)
                                            11.63
                                                  1 7.2796 0.007451 **
```

```
## log(SRP_ug_L)
                                            4.31
                                                   1 2.6983 0.101720
## Temp
                                           38.07
                                                   1 23.8244 1.887e-06 ***
## log(NO3 ug L + 0.01):log(SRP ug L)
                                           12.45
                                                   1 7.7914 0.005656 **
                                           10.74
                                                   1 6.7196 0.010099 *
## log(NO3_ug_L + 0.01):Temp
                                                  1 0.3445 0.557791
## log(SRP_ug_L):Temp
                                           0.55
## log(NO3_ug_L + 0.01):log(SRP_ug_L):Temp
                                           1.71
                                                   1 1.0678 0.302438
## Residuals
                                          397.94 249
## ---
## Signif. codes: 0 '***' 0.001 '**' 0.05 '.' 0.1 ' ' 1
#Log meets assumptions
\#par(mfrow=c(2,2))
#plot(Model)
##Nitrogens
Nitrogen=subset(Data, Trt=="N")
#Chla vs. NO3_ug_L, SRP, Temp ANCOVAs
options(contrasts=c("contr.sum", "contr.poly"))
Model=aov(log(Chla_mean+.01)~log(NO3_ug_L+.01)*log(SRP_ug_L)*Temp, data=Nitrogen)
Anova(Model, type="III")
## Anova Table (Type III tests)
## Response: log(Chla_mean + 0.01)
                                          Sum Sq Df F value Pr(>F)
##
## (Intercept)
                                           0.119 1 0.0898 0.765270
## log(NO3_ug_L + 0.01)
                                           0.054 1 0.0404 0.841300
## log(SRP_ug_L)
                                           0.228 1 0.1720 0.679623
## Temp
                                          10.556 1 7.9559 0.006231 **
## log(NO3_ug_L + 0.01):log(SRP_ug_L)
                                           3.591 1 2.7062 0.104440
                                           0.094 1 0.0710 0.790706
## log(NO3_ug_L + 0.01):Temp
                                           0.004 1 0.0030 0.956816
## log(SRP_ug_L):Temp
## log(NO3_ug_L + 0.01):log(SRP_ug_L):Temp 1.408 1 1.0610 0.306536
## Residuals
                                          92.873 70
## ---
## Signif. codes: 0 '***' 0.001 '**' 0.05 '.' 0.1 ' ' 1
#Log meets assumptions
\#par(mfrow=c(2,2))
#plot(Model)
##Phosphorus
Phosphorus=subset(Data, Trt=="P")
#Chla vs. NO3_ug_L, SRP, Temp ANCOVAs
options(contrasts=c("contr.sum", "contr.poly"))
```

```
Model=aov(log(Chla_mean+.01)~log(NO3_ug_L+.01)*log(SRP_ug_L)*Temp, data=Phosphorus)
Anova(Model, type="III")
## Anova Table (Type III tests)
##
## Response: log(Chla_mean + 0.01)
##
                                          Sum Sq Df F value Pr(>F)
## (Intercept)
                                           2.210 1 2.1127 0.15104
## log(NO3_ug_L + 0.01)
                                           0.479 1 0.4580 0.50103
## log(SRP_ug_L)
                                           4.290 1 4.1016 0.04708 *
                                           0.726 1 0.6940 0.40794
## Temp
## log(NO3_ug_L + 0.01):log(SRP_ug_L)
                                           6.732 1 6.4365 0.01367 *
## log(NO3_ug_L + 0.01):Temp
                                           0.110 1 0.1051 0.74685
## log(SRP_ug_L):Temp
                                           1.201 1 1.1486 0.28793
## log(NO3_ug_L + 0.01):log(SRP_ug_L):Temp 3.296 1 3.1510 0.08071.
## Residuals
                                          65.895 63
## ---
## Signif. codes: 0 '***' 0.001 '**' 0.05 '.' 0.1 ' ' 1
#Log meets assumptions
\#par(mfrow=c(2,2))
#plot(Model)
##Nitrogen + Phosphorus
NP=subset(Data, Trt=="NP")
#Chla vs. NO3_ug_L, SRP, Temp ANCOVAs
options(contrasts=c("contr.sum", "contr.poly"))
Model=aov(log(Chla_mean+.01)~log(NO3_ug_L+.01)*log(SRP_ug_L)*Temp, data=NP)
Anova(Model, type="III")
## Anova Table (Type III tests)
##
## Response: log(Chla_mean + 0.01)
##
                                           Sum Sq Df F value
                                                                Pr(>F)
## (Intercept)
                                           12.769 1 7.5159 0.007308 **
## log(NO3_ug_L + 0.01)
                                            2.282 1 1.3433 0.249355
## log(SRP_ug_L)
                                           35.479 1 20.8834 1.467e-05 ***
                                            1.132 1 0.6664 0.416369
## Temp
## log(NO3_ug_L + 0.01):log(SRP_ug_L)
                                           40.047 1 23.5720 4.720e-06 ***
## log(NO3_ug_L + 0.01):Temp
                                           2.812 1 1.6549 0.201420
## log(SRP_ug_L):Temp
                                           28.105 1 16.5432 9.815e-05 ***
## log(NO3_ug_L + 0.01):log(SRP_ug_L):Temp 25.954 1 15.2769 0.000174 ***
## Residuals
                                          161.396 95
## ---
## Signif. codes: 0 '***' 0.001 '**' 0.05 '.' 0.1 ' ' 1
#Log meets assumptions
\#par(mfrow=c(2,2))
#plot(Model)
```

## b.) Temp-Nut interactions on LRRs

```
##Nitrogens
Nitrogen=subset(Data, Trt=="N")
#LRR vs. NO3_ug_L, SRP, Temp ANCOVAs
options(contrasts=c("contr.sum", "contr.poly"))
Model=aov(LRR_Mean~log(NO3_ug_L+.01)*log(SRP_ug_L)*Temp, data=Nitrogen)
Anova(Model, type="III")
## Anova Table (Type III tests)
##
## Response: LRR_Mean
##
                                           Sum Sq Df F value Pr(>F)
                                           0.4393 1 1.2828 0.2612
## (Intercept)
## log(NO3_ug_L + 0.01)
                                            0.0070 1 0.0203 0.8871
## log(SRP_ug_L)
                                           0.0299 1 0.0873 0.7685
## Temp
                                            0.0037 1 0.0109 0.9173
                                           0.0027 1 0.0078 0.9297
## log(NO3_ug_L + 0.01):log(SRP_ug_L)
## log(NO3_ug_L + 0.01):Temp
                                            0.0010 1 0.0029 0.9568
## log(SRP_ug_L):Temp
                                           0.0074 1 0.0216 0.8836
## log(NO3_ug_L + 0.01):log(SRP_ug_L):Temp 0.0032 1 0.0094 0.9231
## Residuals
                                           23.9695 70
#Log mostly meets assumptions
\#par(mfrow=c(2,2))
#plot(Model)
##Phosphorus
Phosphorus=subset(Data, Trt=="P")
#Chla vs. NO3_ug_L, SRP, Temp ANCOVAs
options(contrasts=c("contr.sum", "contr.poly"))
Model=aov(LRR_Mean~log(NO3_ug_L+.01)*log(SRP_ug_L)*Temp, data=Phosphorus)
Anova(Model, type="III")
## Anova Table (Type III tests)
##
## Response: LRR_Mean
##
                                            Sum Sq Df F value Pr(>F)
## (Intercept)
                                            0.0580 1 0.1759 0.6763
## log(NO3_ug_L + 0.01)
                                            0.0579 1 0.1754 0.6768
## log(SRP_ug_L)
                                            0.0034 1 0.0104 0.9191
## Temp
                                           0.0020 1 0.0061 0.9380
## log(NO3_ug_L + 0.01):log(SRP_ug_L)
                                           0.0001 1 0.0004 0.9838
## log(NO3_ug_L + 0.01):Temp
                                           0.0122 1 0.0371 0.8479
                                           0.0016 1 0.0047 0.9454
## log(SRP_ug_L):Temp
## log(NO3_ug_L + 0.01):log(SRP_ug_L):Temp 0.0062 1 0.0188 0.8915
```

```
## Residuals
                                       20.7854 63
#Log mostly meets assumptions
\#par(mfrow=c(2,2))
#plot(Model)
##Nitrogen + Phosphorus
NP=subset(Data, Trt=="NP")
#Chla vs. NO3_ug_L, SRP, Temp ANCOVAs
options(contrasts=c("contr.sum", "contr.poly"))
Model=aov(LRR_Mean~log(NO3_ug_L+.01)*log(SRP_ug_L)*Temp, data=NP)
Anova(Model, type="III")
## Anova Table (Type III tests)
## Response: LRR_Mean
##
                                       Sum Sq Df F value
                                                           Pr(>F)
## (Intercept)
                                        4.733 1 12.3177 0.0006878 ***
## log(NO3_ug_L + 0.01)
                                        0.015 1 0.0380 0.8458286
## log(SRP_ug_L)
                                        5.753 1 14.9697 0.0002002 ***
                                        0.298 1 0.7750 0.3809054
## Temp
## log(NO3_ug_L + 0.01):log(SRP_ug_L)
                                        1.084 1 2.8210 0.0963264 .
## log(NO3_ug_L + 0.01):Temp
                                        0.015 1 0.0395 0.8428611
## log(SRP_ug_L):Temp
                                        4.519 1 11.7601 0.0008967 ***
## Residuals
                                       36.507 95
## ---
## Signif. codes: 0 '***' 0.001 '**' 0.05 '.' 0.1 ' ' 1
#Log meets assumptions
\#par(mfrow=c(2,2))
#plot(Model)
```