

# Regression Models - Morrison Bridge NAWQA

*Mark Roes*

*June 3, 2017*

```
data.tsi <- read.csv(series, stringsAsFactors = FALSE)
data.di <- read.csv(discrete, stringsAsFactors = FALSE)
data.all <- merge(data.tsi, data.di, by="DateTime", all.y=TRUE)
#data.all <- with(data.all, rm("DateTime", "Date", "date", "time", "sample_dt", "sample_tm"))
# manage dates
data.all$DateTime <- chon(dates=data.all$Date.chon, times=data.all$Time.chon,
                          format=c("m/d/y", "h:m:s"))
```

Some info of Metolachlor:

Metolachlor is an herbicide used in agriculture to control grasses. Its use is controversial, the chemical is approved in the US for use, but NOT in the EU. There are tolerance levels for this chemical in agriculture, but no maximum allowable level in drinking water – only an advisory level. According to GHS (Globally Harmonized System for Hazard Communication), an international accord on chemical hazard, Metolachlor is:

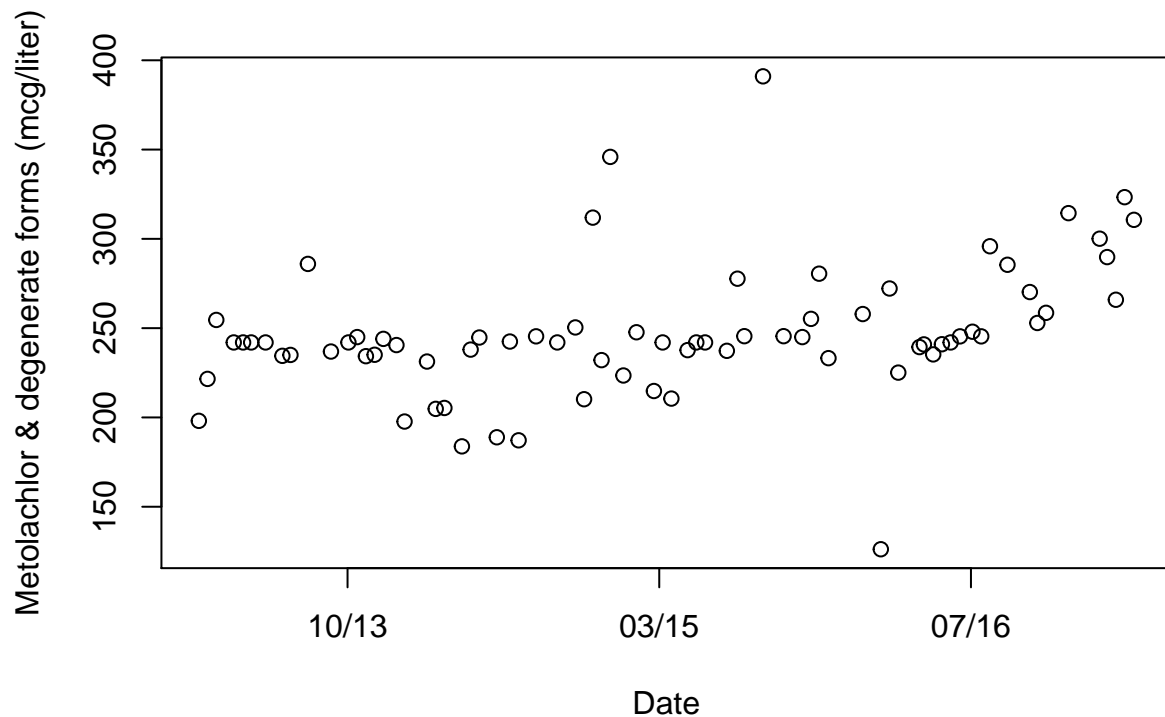
May cause an allergic skin reaction [Warning Sensitization, Skin - Category 1]

Fatal if inhaled [Danger Acute toxicity, inhalation - Category 1, 2]

Very toxic to aquatic life [Warning Hazardous to the aquatic environment, acute hazard - Category 1]

Very toxic to aquatic life with long lasting effects [Warning Hazardous to the aquatic environment, long-term hazard - Category 1]

```
plot(data.all$DateTime, data.all$metoforms, xlab = "Date", ylab = "Metolachlor & degenerate forms (mcg/
```



*#lets try a simple general additive mixed model!*

```
gam.meto<-gam(metoforms~Discharge+StreamVelocity+GageHeight+SensorDepth+WaterTemp+DissolvedO2+O2Saturat.
summary(gam.meto)
```

```
##
## Family: gaussian
## Link function: identity
##
## Formula:
## metoforms ~ Discharge + StreamVelocity + GageHeight + SensorDepth +
##      WaterTemp + DissolvedO2 + O2Saturation + pH + Conductance +
##      Turbidity + SecchiDepth + Chlorophyll + Phycocyanin + fDOM +
##      Nitrate
##
## Parametric coefficients:
##              Estimate Std. Error t value Pr(>|t|)
## (Intercept)  -735.5961   453.0750  -1.624   0.1099
## Discharge      -0.5350     3.0007  -0.178   0.8591
## StreamVelocity  15.6134   116.5071   0.134   0.8939
## GageHeight     -5.0014     9.4019  -0.532   0.5968
## SensorDepth     5.7730     9.0040   0.641   0.5239
## WaterTemp      10.9407     6.9564   1.573   0.1212
## DissolvedO2     37.1763    28.6895   1.296   0.2002
## O2Saturation   -1.7410     2.6390  -0.660   0.5120
## pH              67.8281    52.0944   1.302   0.1981
## Conductance     1.5553     0.9301   1.672   0.0999 .
```

```
## Turbidity      1.7402      1.5456      1.126      0.2648
## SecchiDepth    -3.8917      4.4564     -0.873      0.3861
## Chlorophyll    -9.2617      4.1103     -2.253      0.0280 *
## Phycocyanin    37.4974     33.4809      1.120      0.2673
## fDOM           -0.4162      1.8511     -0.225      0.8229
## Nitrate       -26.3492     40.4253     -0.652      0.5171
## ---
## Signif. codes:  0 '***' 0.001 '**' 0.01 '*' 0.05 '.' 0.1 ' ' 1
##
##
## R-sq.(adj) =  0.0658   Deviance explained = 25.8%
## GCV =    1710   Scale est. = 1340.3      n = 74
```

```
#reduced gam model, did a stepwise removal of factors with highest p-values
#If i remove any more than this, the % deviance explained drops drastically
reduced.gam.meto<-gam(metoforms~WaterTemp+DissolvedO2+pH+Conductance+Turbidity+Chlorophyll+Phycocyanin,
summary(reduced.gam.meto)
```

```
##
## Family: gaussian
## Link function: identity
##
## Formula:
## metoforms ~ WaterTemp + DissolvedO2 + pH + Conductance + Turbidity +
##      Chlorophyll + Phycocyanin
##
## Parametric coefficients:
##              Estimate Std. Error t value Pr(>|t|)
## (Intercept) -590.2417   367.3369  -1.607   0.1129
## WaterTemp     7.3345     3.5943   2.041   0.0453 *
## DissolvedO2   20.6528    12.3621   1.671   0.0995 .
## pH            57.2329    39.2944   1.457   0.1500
## Conductance    1.1694     0.6158   1.899   0.0619 .
## Turbidity      1.4945     0.9118   1.639   0.1060
## Chlorophyll   -7.9208     3.5383  -2.239   0.0286 *
## Phycocyanin   31.1671    24.9213   1.251   0.2155
## ---
## Signif. codes:  0 '***' 0.001 '**' 0.01 '*' 0.05 '.' 0.1 ' ' 1
##
##
## R-sq.(adj) =  0.146   Deviance explained = 22.8%
## GCV = 1374.2   Scale est. = 1225.7      n = 74
```

```
#linear model
lm.meto<-lm(metoforms~Discharge+StreamVelocity+GageHeight+SensorDepth+WaterTemp+DissolvedO2+O2Saturation+
summary(lm.meto)
```

```
##
## Call:
## lm(formula = metoforms ~ Discharge + StreamVelocity + GageHeight +
##      SensorDepth + WaterTemp + DissolvedO2 + O2Saturation + pH +
##      Conductance + Turbidity + SecchiDepth + Chlorophyll + Phycocyanin +
##      fDOM + Nitrate, data = data.all)
##
## Residuals:
##      Min       1Q   Median       3Q      Max
```

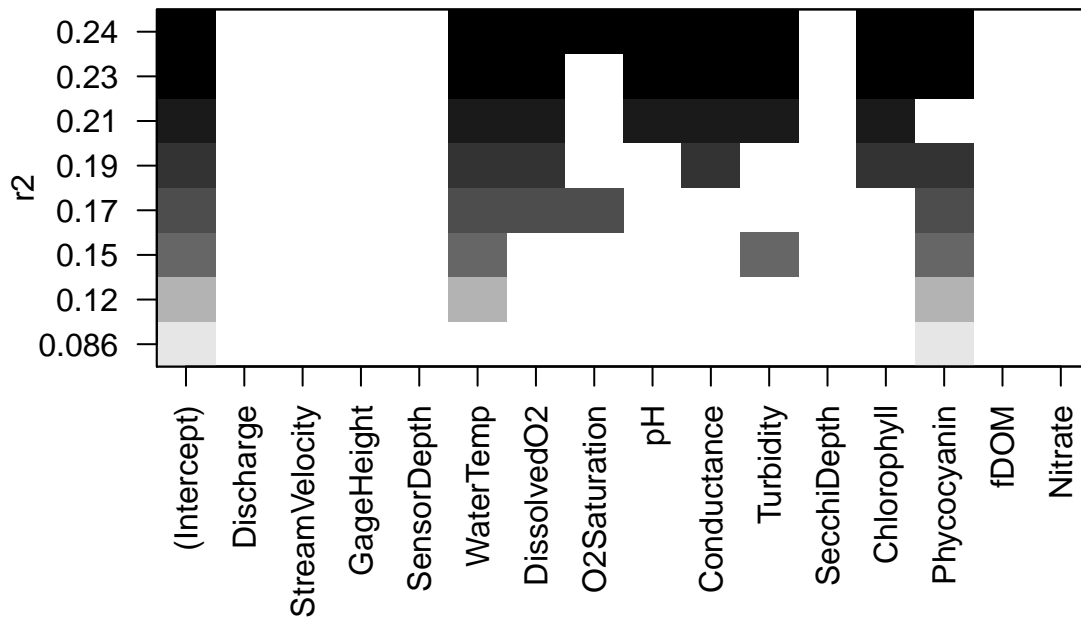
```
## -134.846 -14.244 2.444 10.521 108.683
##
## Coefficients:
## Estimate Std. Error t value Pr(>|t|)
## (Intercept) -735.5961 453.0750 -1.624 0.1099
## Discharge -0.5350 3.0007 -0.178 0.8591
## StreamVelocity 15.6134 116.5071 0.134 0.8939
## GageHeight -5.0014 9.4019 -0.532 0.5968
## SensorDepth 5.7730 9.0040 0.641 0.5239
## WaterTemp 10.9407 6.9564 1.573 0.1212
## DissolvedO2 37.1763 28.6895 1.296 0.2002
## O2Saturation -1.7410 2.6390 -0.660 0.5120
## pH 67.8281 52.0944 1.302 0.1981
## Conductance 1.5553 0.9301 1.672 0.0999 .
## Turbidity 1.7402 1.5456 1.126 0.2648
## SecchiDepth -3.8917 4.4564 -0.873 0.3861
## Chlorophyll -9.2617 4.1103 -2.253 0.0280 *
## Phycocyanin 37.4974 33.4809 1.120 0.2673
## fDOM -0.4162 1.8511 -0.225 0.8229
## Nitrate -26.3492 40.4253 -0.652 0.5171
## ---
## Signif. codes: 0 '***' 0.001 '**' 0.01 '*' 0.05 '.' 0.1 ' ' 1
##
## Residual standard error: 36.61 on 58 degrees of freedom
## Multiple R-squared: 0.2577, Adjusted R-squared: 0.06576
## F-statistic: 1.343 on 15 and 58 DF, p-value: 0.2078
```

```
reduced.lm<-lm(metoforms~WaterTemp+DissolvedO2+pH+Conductance+Turbidity+Chlorophyll+Phycocyanin, data =
summary(reduced.lm)
```

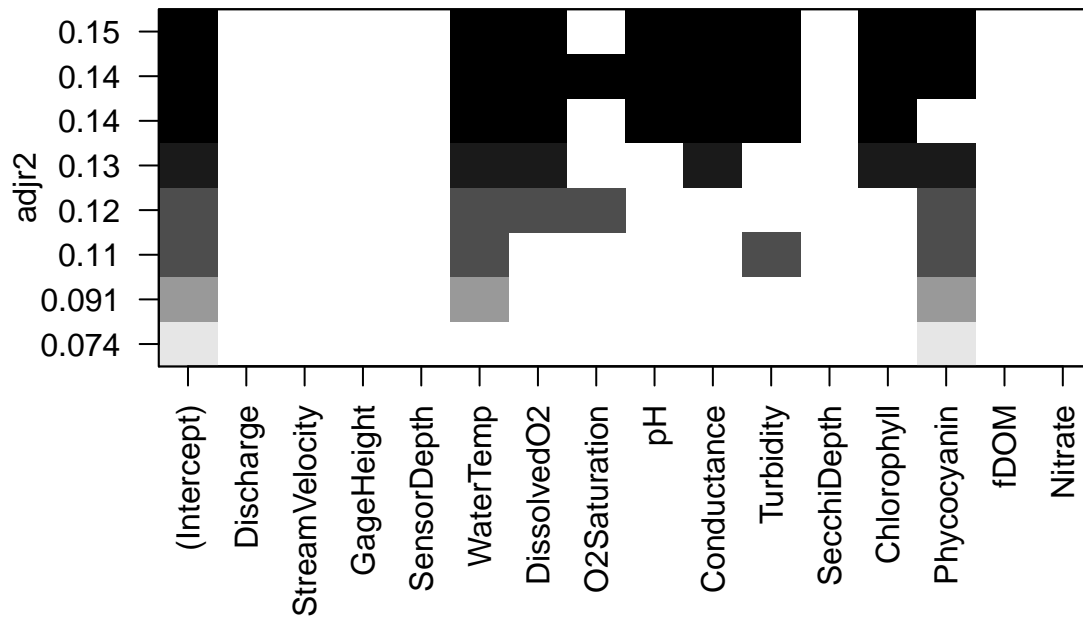
```
##
## Call:
## lm(formula = metoforms ~ WaterTemp + DissolvedO2 + pH + Conductance +
## Turbidity + Chlorophyll + Phycocyanin, data = data.all)
##
## Residuals:
## Min 1Q Median 3Q Max
## -131.063 -12.714 -1.717 13.474 104.427
##
## Coefficients:
## Estimate Std. Error t value Pr(>|t|)
## (Intercept) -590.2417 367.3369 -1.607 0.1129
## WaterTemp 7.3345 3.5943 2.041 0.0453 *
## DissolvedO2 20.6528 12.3621 1.671 0.0995 .
## pH 57.2329 39.2944 1.457 0.1500
## Conductance 1.1694 0.6158 1.899 0.0619 .
## Turbidity 1.4945 0.9118 1.639 0.1060
## Chlorophyll -7.9208 3.5383 -2.239 0.0286 *
## Phycocyanin 31.1671 24.9213 1.251 0.2155
## ---
## Signif. codes: 0 '***' 0.001 '**' 0.01 '*' 0.05 '.' 0.1 ' ' 1
##
## Residual standard error: 35.01 on 66 degrees of freedom
## Multiple R-squared: 0.2276, Adjusted R-squared: 0.1456
## F-statistic: 2.778 on 7 and 66 DF, p-value: 0.01357
```

The linear model with all factors has only 'chlorophyll' as significant. Using the leaps package to test the optimal set of factors.

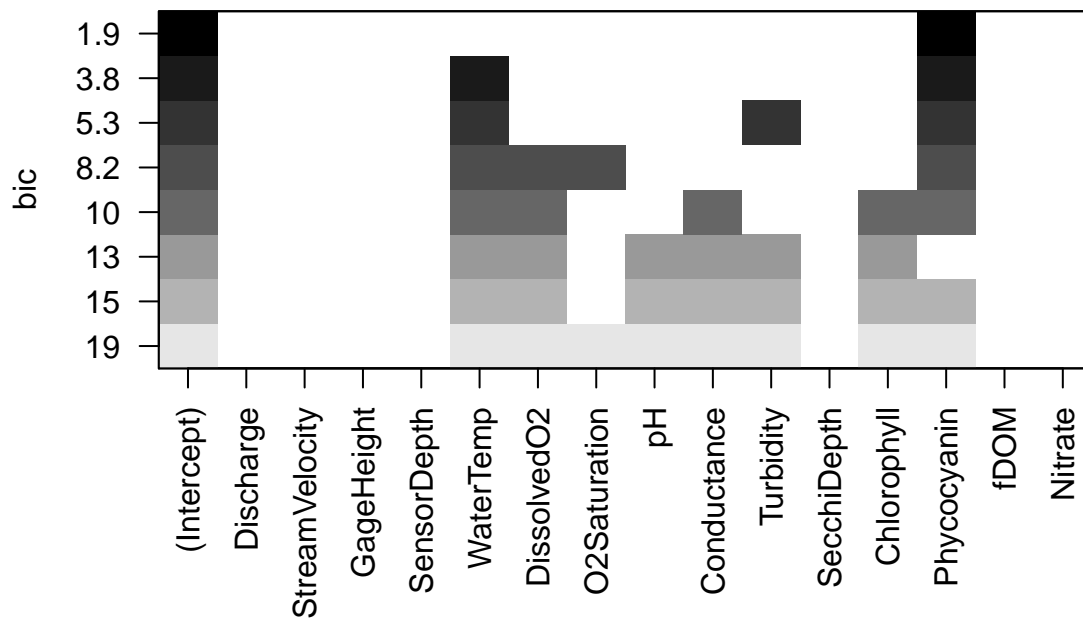
```
leaps.meto<-regsubsets(x = data.all[,4:18], y = data.all$metoforms)
plot(leaps.meto, scale = "r2")
```



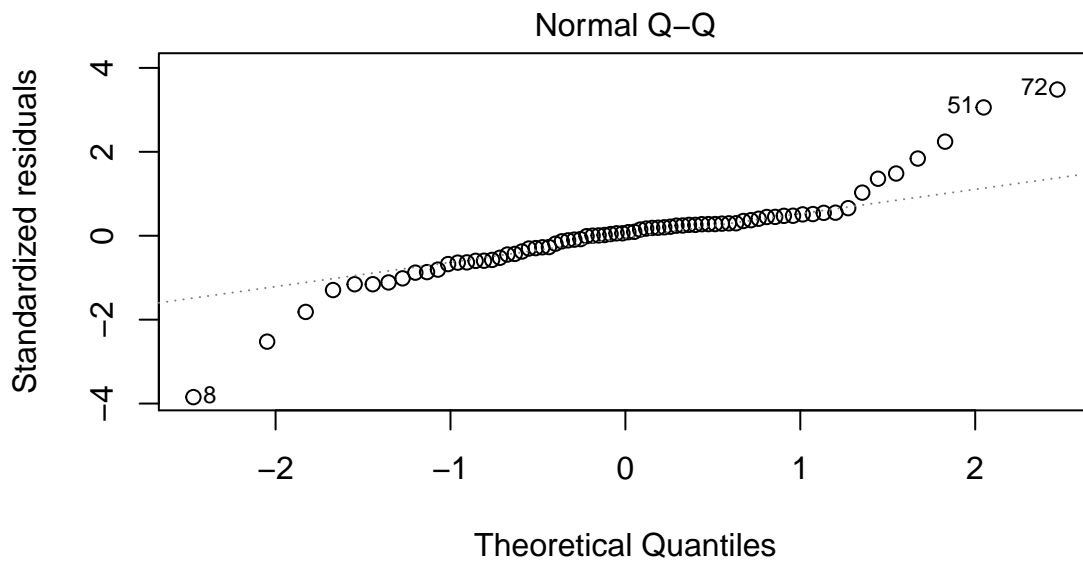
```
plot(leaps.meto, scale = "adjr2")
```



```
plot(leaps.meto, scale = "bic")
```



```
#Now test for fit with these 3 models created by leaps
#full model
plot(lm(metoforms~Discharge+StreamVelocity+GageHeight+SensorDepth+WaterTemp+DissolvedO2+O2Saturation+pH
```

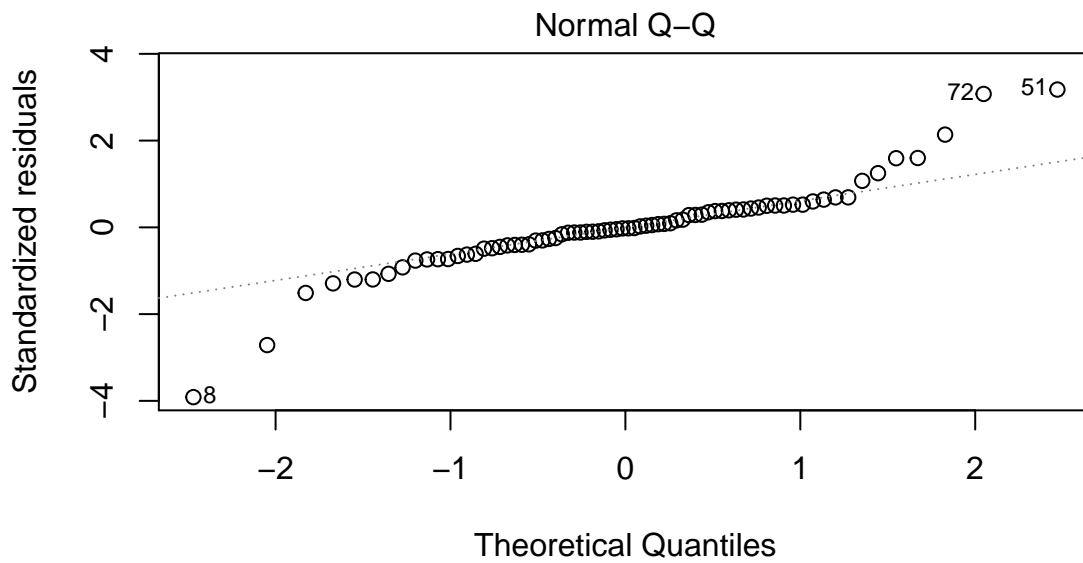


Im(metoforms ~ Discharge + StreamVelocity + GageHeight + SensorDepth + Wate .

*#highest r2*

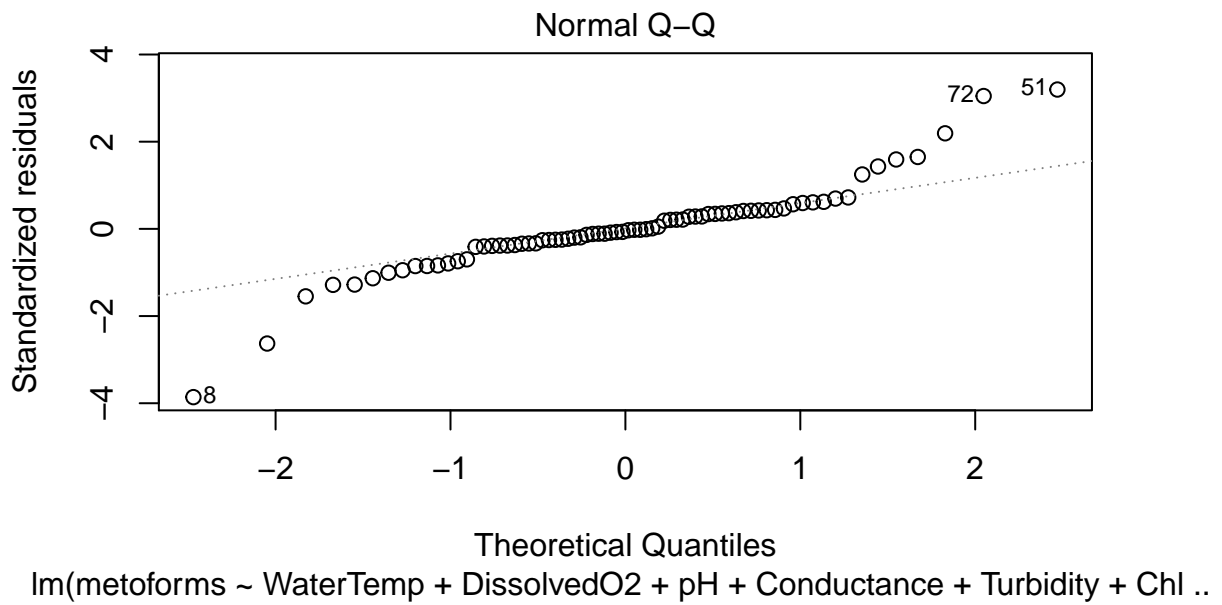
`plot(lm(metoforms~WaterTemp+DissolvedO2+O2Saturation+pH+Conductance+Turbidity+Chlorophyll+Phycocyanin, c`



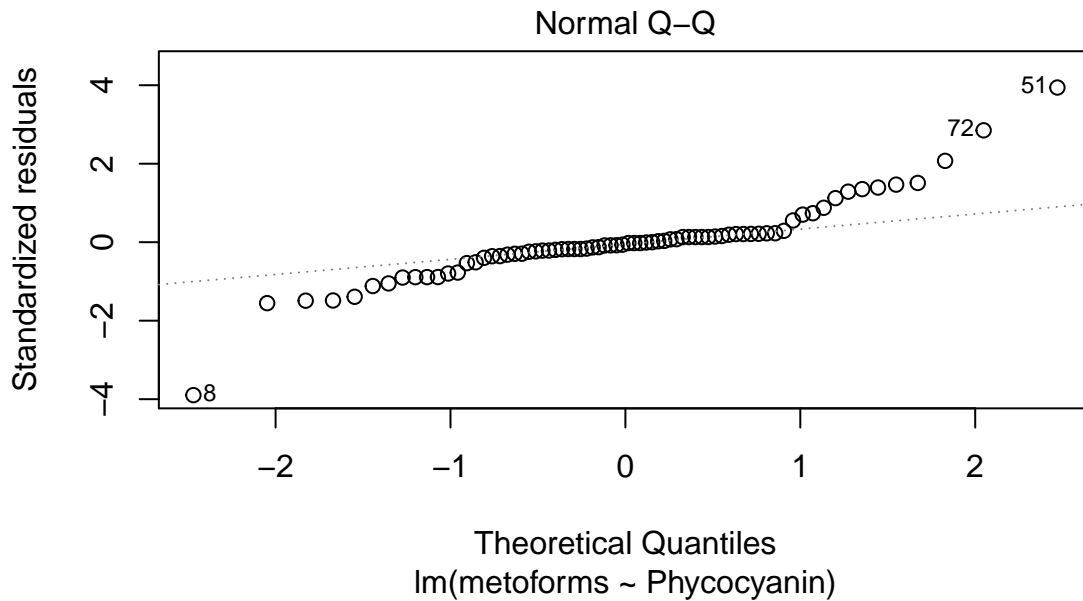


Im(metoforms ~ WaterTemp + DissolvedO2 + O2Saturation + pH + Conductance + .

```
#highest adjr2
plot(lm(metoforms~WaterTemp+DissolvedO2+pH+Conductance+Turbidity+Chlorophyll+Phycocyanin, data = data.a
```



```
#lowest BIC
plot(lm(metoforms~Phycocyanin, data = data.all), which = 2)
```



```
#quantile regression for meto with tau = .5
rq.meto<-rq(metoforms~WaterTemp+DissolvedO2+pH+Conductance+Turbidity+Chlorophyll+Phycocyanin, data = da
summary(rq.meto)
```

```
##
## Call: rq(formula = metoforms ~ WaterTemp + DissolvedO2 + pH + Conductance +
##       Turbidity + Chlorophyll + Phycocyanin, data = data.all)
##
## tau: [1] 0.5
##
## Coefficients:
##           coefficients lower bd   upper bd
## (Intercept) -285.12336   -983.40177   323.35978
## WaterTemp      1.33968     -2.17665    9.43410
## DissolvedO2     0.77693    -11.62171   20.75141
## pH             60.86486     -8.13841  110.82329
## Conductance     0.63455     -0.05009    1.03935
## Turbidity       1.70864      0.03051    2.38374
## Chlorophyll    -4.68017     -6.84272    0.48336
## Phycocyanin     35.00731      1.45508   85.75796
```

```
#Let's compare the AIC values for all of these models:
#AIC for quantile regression
AIC.rq(rq.meto)
```

```
## [1] 713.1188
## attr(,"edf")
```

```
## [1] 8
#AIC for full LM model
AIC(lm.meto)

## [1] 758.8205
#AIC for best linear model by BIC
AIC(lm(metoforms~Phycocyanin, data = data.all))

## [1] 746.2035
#AIC for best linear model by r2
AIC(lm(metoforms~WaterTemp+DissolvedO2+O2Saturation+pH+Conductance+Turbidity+Chlorophyll+Phycocyanin, data = data.all))

## [1] 746.8268
#AIC for best linear model by adjr2
AIC(lm(metoforms~WaterTemp+DissolvedO2+pH+Conductance+Turbidity+Chlorophyll+Phycocyanin, data = data.all))

## [1] 745.7691
#AIC for the two general additive mixed models
llgam<-logLik.gam(gam.meto)
AIC(llgam)

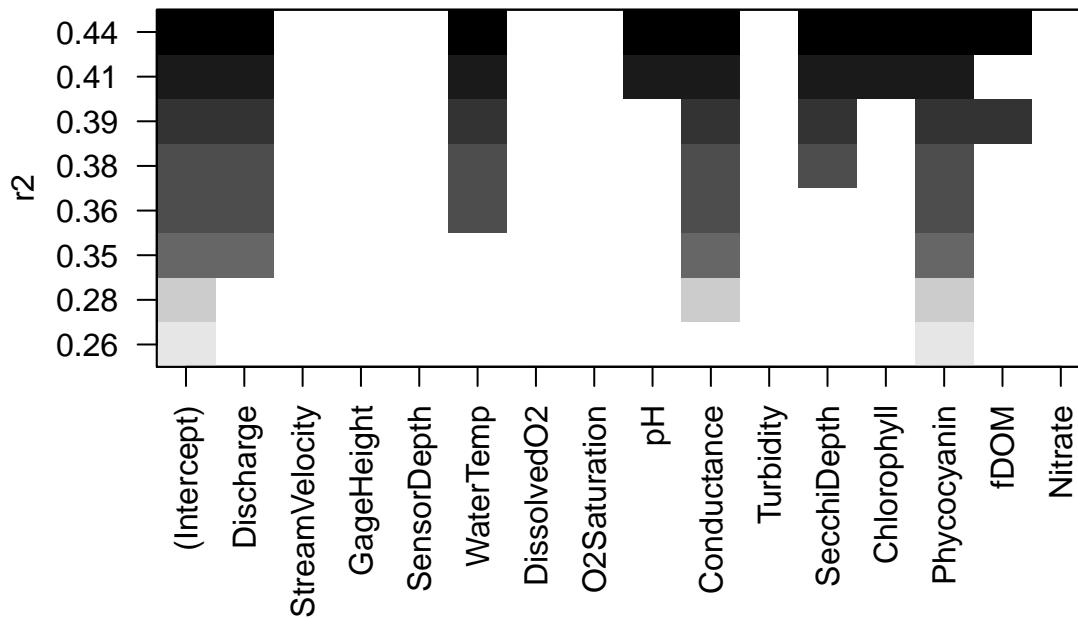
## [1] 758.8205
red.llgam<-logLik.gam(reduced.gam.meto)
AIC(red.llgam)

## [1] 745.7691
#nonlinear quantile regression estimate, havent got to work yet
#nlrq.meto<-nlrq(metoforms~WaterTemp+DissolvedO2+pH+Conductance+Turbidity+Chlorophyll+Phycocyanin, data = data.all)

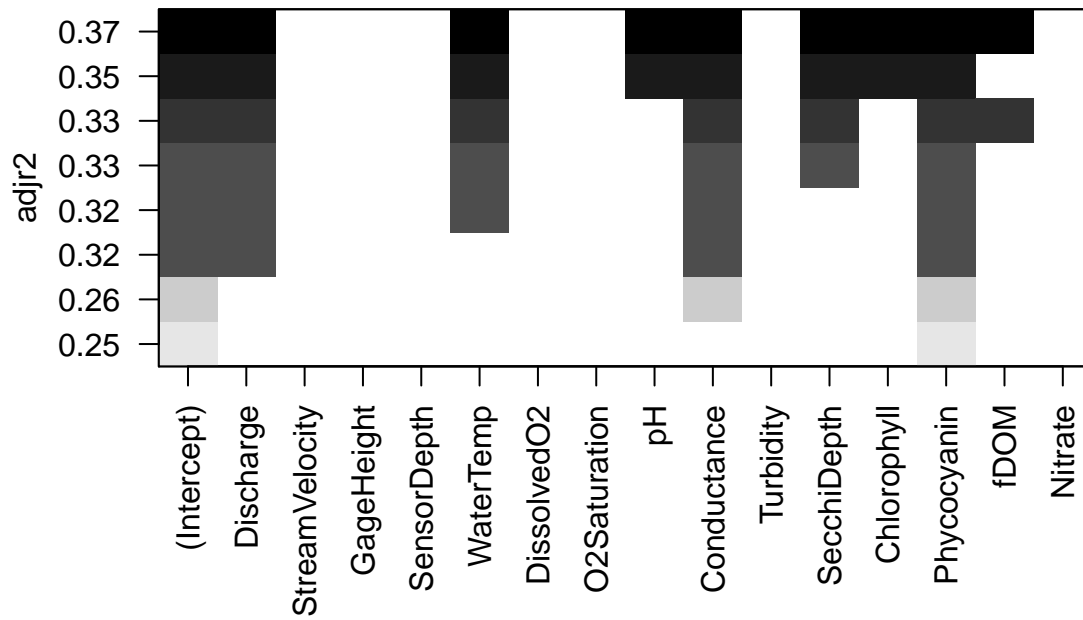
total<-data.all$diazforms+data.all$metoforms+data.all$hexaforms+data.all$fipforms
lm.all<-lm(total~Discharge+StreamVelocity+GageHeight+SensorDepth+WaterTemp+DissolvedO2+O2Saturation+pH+
summary(lm.all)

##
## Call:
## lm(formula = total ~ Discharge + StreamVelocity + GageHeight +
##      SensorDepth + WaterTemp + DissolvedO2 + O2Saturation + pH +
##      Conductance + Turbidity + SecchiDepth + Chlorophyll + Phycocyanin +
##      fDOM + Nitrate, data = data.all)
##
## Residuals:
##      Min       1Q   Median       3Q      Max
## -136.00  -34.32  -10.10   27.45  246.89
##
## Coefficients:
##              Estimate Std. Error t value Pr(>|t|)
## (Intercept)  -1853.5910   896.8906  -2.067  0.0432 *
## Discharge       2.9646     5.9401   0.499  0.6196
## StreamVelocity -69.9011   230.6331  -0.303  0.7629
## GageHeight    -14.3647    18.6117  -0.772  0.4434
## SensorDepth    19.0579    17.8240   1.069  0.2894
## WaterTemp     16.6452    13.7707   1.209  0.2317
## DissolvedO2    45.9892    56.7927   0.810  0.4214
```

```
## O2Saturation      -3.7784      5.2240     -0.723     0.4724
## pH                258.2146    103.1242      2.504     0.0151 *
## Conductance       4.1235      1.8411      2.240     0.0290 *
## Turbidity        -0.9972      3.0596     -0.326     0.7456
## SecchiDepth      -18.4593      8.8217     -2.092     0.0408 *
## Chlorophyll      -18.6846      8.1367     -2.296     0.0253 *
## Phycocyanin      145.6791     66.2776      2.198     0.0320 *
## fDOM              -6.1385      3.6645     -1.675     0.0993 .
## Nitrate          89.3141     80.0245      1.116     0.2690
## ---
## Signif. codes:  0 '***' 0.001 '**' 0.01 '*' 0.05 '.' 0.1 ' ' 1
##
## Residual standard error: 72.47 on 58 degrees of freedom
## Multiple R-squared:  0.483, Adjusted R-squared:  0.3493
## F-statistic: 3.612 on 15 and 58 DF, p-value: 0.0002078
subset.all<-regsubsets(x = data.all[,4:18], y = total)
plot(subset.all, scale = "r2")
```



```
plot(subset.all, scale = "adjr2")
```



```
plot(subset.all, scale = "bic")
```

