

W271 Spring 18: Lab 2

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Alcohol Consumption, Self-Esteem and Romantic Interactions

Introduction

The researchers stated the hypothesis as follow: “We hypothesized that negative interactions with romantic partners would be associated with alcohol consumption (and an increased desire to drink). We predicted that people with low trait self-esteem would drink more on days they experienced more negative relationship interactions compared with days during which they experienced fewer negative relationship interactions. The relation between drinking and negative relationship interactions should not be evident for individuals with high trait self-esteem.”

EDA

```
library(car); require(dplyr); library(Hmisc); library(mcprofile); library(ggplot2); library(gridExtra);
dehart <- read.table(file="DeHartSimplified.csv", header=TRUE, sep=",")
describe(dehart) #with a 10-page limit, should we include this type of output?
```

```
## dehart
##
## 13 Variables      623 Observations
## -----
## id
##      n missing distinct    Info    Mean    Gmd      .05      .10
##    623      0      89      1    75.89   56.82    7.0    16.2
##    .25    .50    .75    .90    .95
##   33.0   60.0   123.0  147.2   153.0
##
## lowest :    1    2    4    5    7, highest: 153 154 155 156 160
## -----
## studyday
##      n missing distinct    Info    Mean    Gmd
##    623      0      7    0.98      4    2.289
##
## Value      1      2      3      4      5      6      7
## Frequency    89    89    89    89    89    89    89
## Proportion 0.143 0.143 0.143 0.143 0.143 0.143 0.143
## -----
## dayweek
##      n missing distinct    Info    Mean    Gmd
##    623      0      7    0.98      4    2.289
##
## Value      1      2      3      4      5      6      7
## Frequency    89    89    89    89    89    89    89
## Proportion 0.143 0.143 0.143 0.143 0.143 0.143 0.143
## -----
## numall
```

```

##      n missing distinct      Info      Mean      Gmd      .05      .10
##    622      1      18      0.97      2.524      2.636      0.00      0.00
##      .25      .50      .75      .90      .95
##    1.00      2.00      3.75      6.00      8.00
##
## Value      0      1      2      3      4      5      6      7      8      9
## Frequency  141  112  132   81   49   43   24   6    9    7
## Proportion 0.227 0.180 0.212 0.130 0.079 0.069 0.039 0.010 0.014 0.011
##
## Value      10     11     12     13     14     15     18     21
## Frequency    7      4      2      1      1      1      1      1
## Proportion 0.011 0.006 0.003 0.002 0.002 0.002 0.002 0.002
## -----
## nrel
##      n missing distinct      Info      Mean      Gmd      .05      .10
##    623      0      33      0.551      0.359      0.6252      0      0
##      .25      .50      .75      .90      .95
##      0      0      0      1      2
##
## lowest : 0.0000000 0.2000000 0.2500000 0.3333333 0.4000000
## highest: 5.0000000 5.5000000 5.8333333 6.0000000 9.0000000
## -----
## prel
##      n missing distinct      Info      Mean      Gmd      .05      .10
##    623      0      68      0.982      2.583      2.613      0.0000      0.0000
##      .25      .50      .75      .90      .95
##    0.4167      2.0000      4.0000      6.0000      7.8683
##
## lowest : 0.0000000 0.2000000 0.2500000 0.3333333 0.5000000
## highest: 8.1666667 8.3333333 8.5000000 8.6666667 9.0000000
## -----
## negevent
##      n missing distinct      Info      Mean      Gmd      .05      .10
##    623      0      131      0.996      0.4414      0.4123      0.0000      0.0000
##      .25      .50      .75      .90      .95
##    0.1583      0.3500      0.6292      1.0000      1.1500
##
## lowest : 0.00000000 0.02500000 0.03333333 0.05000000 0.07500000
## highest: 1.70000000 1.93000000 1.95000000 2.01666667 2.37666667
## -----
## posevent
##      n missing distinct      Info      Mean      Gmd      .05      .10
##    623      0      216      1      1.048      0.7077      0.200      0.300
##      .25      .50      .75      .90      .95
##    0.600      0.950      1.378      1.938      2.200
##
## lowest : 0.00000000 0.04000000 0.05000000 0.06666667 0.10000000
## highest: 3.23333333 3.25000000 3.30000000 3.40000000 3.88333333
## -----
## gender
##      n missing distinct      Info      Mean      Gmd
##    623      0      2      0.739      1.562      0.4932
##
## Value      1      2

```

```
## Frequency      273    350
## Proportion 0.438 0.562
## -----
## rosn
##      n missing distinct      Info      Mean      Gmd      .05      .10
##    623      0      17    0.993    3.436    0.4663    2.7    2.9
##    .25    .50    .75    .90    .95
##    3.2    3.5    3.8    3.9    4.0
##
## Value      2.1    2.4    2.5    2.7    2.8    2.9    3.0    3.1    3.2    3.3
## Frequency      7      7    14      7    21    35    42    21    28    42
## Proportion 0.011 0.011 0.022 0.011 0.034 0.056 0.067 0.034 0.045 0.067
##
## Value      3.4    3.5    3.6    3.7    3.8    3.9    4.0
## Frequency    35    84    63    49    63    49    56
## Proportion 0.056 0.135 0.101 0.079 0.101 0.079 0.090
## -----
## age
##      n missing distinct      Info      Mean      Gmd      .05      .10
##    623      0      89      1    34.29    5.18    26.24    27.82
##    .25    .50    .75    .90    .95
##   30.53   34.57   38.19   40.15   40.56
##
## lowest : 24.43258 25.57700 26.05613 26.14100 26.23682
## highest: 40.56400 40.58864 40.68720 40.82957 42.27789
## -----
## desired
##      n missing distinct      Info      Mean      Gmd      .05      .10
##    620      3      22    0.996    4.465    1.921    1.333    2.000
##    .25    .50    .75    .90    .95
##    3.333    4.667    5.667    6.667    7.333
##
## lowest : 1.000000 1.333333 1.666667 2.000000 2.333333
## highest: 6.666667 7.000000 7.333333 7.666667 8.000000
## -----
## state
##      n missing distinct      Info      Mean      Gmd      .05      .10
##    620      3      25    0.993    3.966    0.4894    3.222    3.333
##    .25    .50    .75    .90    .95
##    3.667    4.000    4.222    4.556    4.556
##
## lowest : 2.333333 2.444444 2.555556 2.666667 2.777778
## highest: 4.555556 4.666667 4.777778 4.888889 5.000000
## -----
```

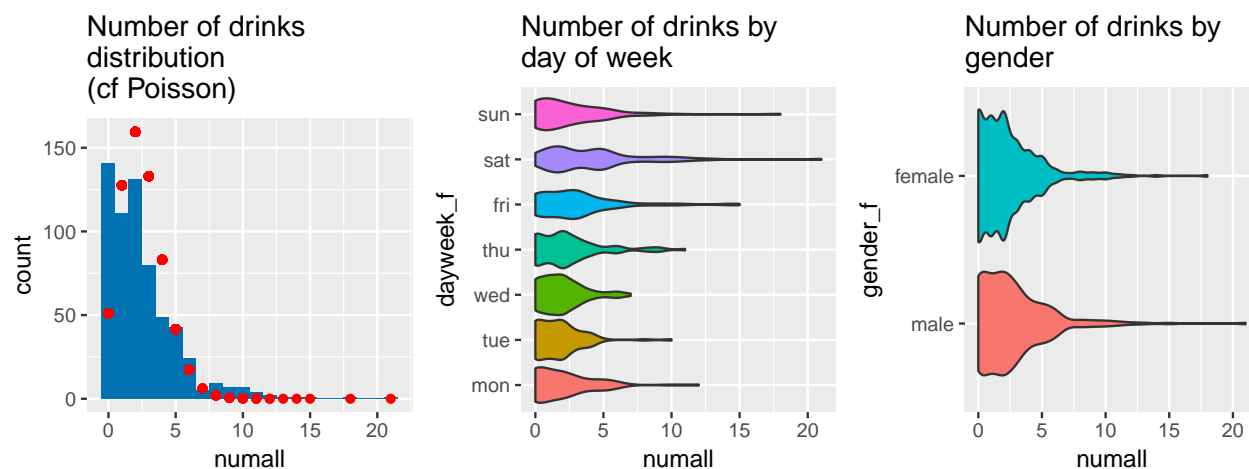
The dataset contains 623 observations of 13 variables representing entries in records kept by study participants. The variable **id** is a numeric identifier for each of the 89 study participants. Each participant recorded entries for seven consecutive days, indexed by the **studyday** variable, with the **dayweek** variable indicating which days of the week these correspond to (Monday = 1). The variable **gender** takes on one of two values according to whether the participant is male (1) or female (2); about 56% of the participants are female.

```
dehart$dayweek_f <- factor(dehart$dayweek); levels(dehart$dayweek_f) = c("mon","tue","wed","thu","fri",
dehart$gender_f <- factor(dehart$gender); levels(dehart$gender_f) = c("male","female")
```

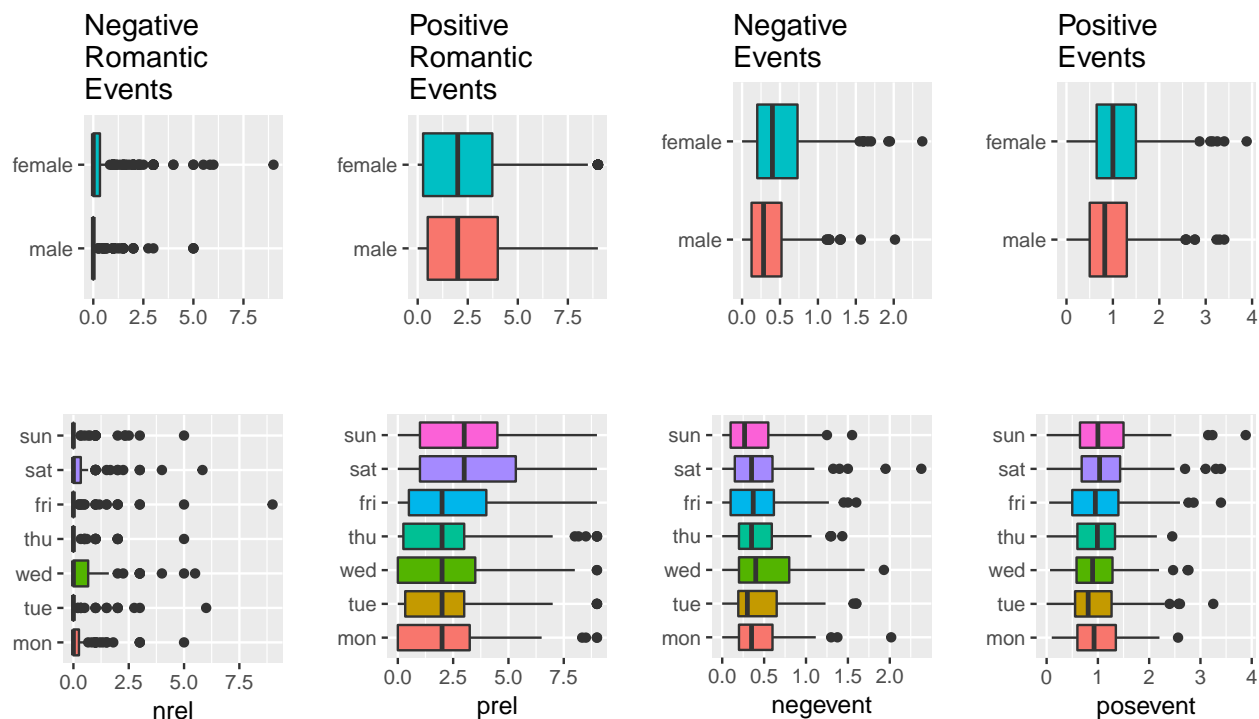
For each of the seven days, participants record the number of drinks consumed with the integer **numall** count variable. There is one missing value. Values range from 0 to 15 with single outliers at 19 and 21. Observations are concentrated in the range 0 to 5. The sample mean and sample variance are 2.52 and 2.66 respectively. The median number of drinks is 3 for Friday, 4 for Saturday and 2 for all other days. There is a pronounced positive skew to the number of drinks for Fridays, Saturdays and Sundays.

The variables **nrel** and **prel** are index measures for the negative and positive romantic-relationship events experienced by the subject each day (ranging from 0 to around 9), while **negevent** and **posevent** are index values combining the total number and intensity of negative and positive events experienced that day (ranging from 0 to around 4). There are no missing values for these variables. Distributions (and scales) for these measurements are very different, although tend to be similar for males and females, and similar across days of the week, although **prel** has significantly higher average (mean and median) values on Saturdays and Sundays. Distributions for all these variables are strongly positively-skewed, but this is particularly extreme for **nrel**, where a small fraction of outlying observations accounts for almost all of the variation.

```
#mean(dehart$numall, na.rm=TRUE); sd(dehart$numall, na.rm=TRUE)
#aggregate(numall ~ dayweek_f, data = dehart, FUN = function(x) c(m = mean(x), n = median(x)))
#The distribution of 622 times a Poisson random variable with parameter lambda is overlaid on the histo
lambda = 2.5
p1 <- ggplot(na.omit(dehart), aes(x = numall)) + geom_histogram(aes(y = ..count..), binwidth = 1, fill=
  geom_point(aes(y = 622*dpois(x = numall,lambda)), color = "red")+ ggtitle("Number of drinks\ndistribu
p2<-ggplot(na.omit(dehart), aes(dayweek_f, numall)) + geom_violin(aes(fill = dayweek_f)) + ggtitle("Num
p3<-ggplot(na.omit(dehart), aes(gender_f, numall)) + geom_violin(aes(fill = gender_f)) + ggtitle("Numbe
grid.arrange(p1, p2, p3, ncol = 3)
```



```
p1a<-ggplot(dehart, aes(gender_f, nrel)) + geom_boxplot(aes(fill = gender_f)) + labs(x = "", y = "") +
p1b<-ggplot(dehart, aes(dayweek_f, nrel)) + geom_boxplot(aes(fill = dayweek_f)) + labs(x = "") + ggtitle
p2a<-ggplot(dehart, aes(gender_f, prel)) + geom_boxplot(aes(fill = gender_f)) + labs(x = "", y = "") +
p2b<-ggplot(dehart, aes(dayweek_f, prel)) + geom_boxplot(aes(fill = dayweek_f)) + labs(x = "") + ggtitle
p3a<-ggplot(dehart, aes(gender_f, negevent)) + geom_boxplot(aes(fill = gender_f)) + labs(x = "", y = "")
p3b<-ggplot(dehart, aes(dayweek_f, negevent)) + geom_boxplot(aes(fill = dayweek_f)) + labs(x = "") + gg
p4a<-ggplot(dehart, aes(gender_f, posevent)) + geom_boxplot(aes(fill = gender_f)) + labs(x = "", y = "")
p4b<-ggplot(dehart, aes(dayweek_f, posevent)) + geom_boxplot(aes(fill = dayweek_f)) + labs(x = "") + gg
grid.arrange(p1a, p2a, p3a, p4a, p1b, p2b, p3b, p4b, ncol = 4)
```



```
aggregate(nrel ~ dayweek_f, data = dehart, FUN = mean)
```

```
##   dayweek_f    nrel
## 1      mon 0.3471910
## 2      tue 0.2882022
## 3      wed 0.5541466
## 4      thu 0.2295880
## 5      fri 0.4014981
## 6      sat 0.4033708
## 7      sun 0.2893258
```

The **roasn** variable measures trait (long-term) self-esteem, a single measurement for each participant taken at the beginning of the study that does not change over the course of the seven days. This measurement ranges between 2 and 4, with a mean value around 3.4. Distributions for males and females differ, with a larger proportion of males recording lower values and a larger proportion of females recording higher values. The **age** variable measures age in years, ranging between 24.4 and 42.3 with a mean value of 34.3 and similar distributions for males and females but with a higher proportion of females recording higher values. Neither **roasn** nor **age** have any missing observations.

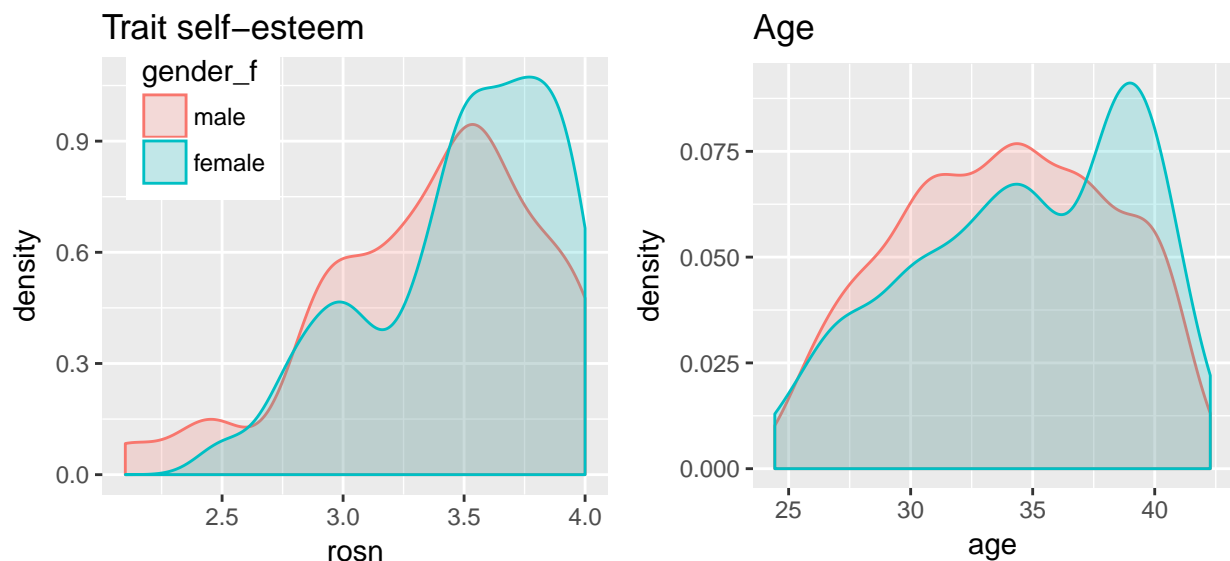
```
quantile(dehart$roasn[dehart$gender == 1])
```

```
##   0%  25%  50%  75% 100%
##  2.1  3.0  3.5  3.7  4.0
```

```
quantile(dehart$roasn[dehart$gender == 2])
```

```
##   0%  25%  50%  75% 100%
##  2.50 3.30 3.55 3.80 4.00
```

```
p1<-ggplot(dehart, aes(x = roasn, fill = gender_f, colour = gender_f)) + geom_density(alpha=0.2)+ ggtitle("Self-Esteem")
p2<-ggplot(dehart, aes(x = age, fill = gender_f, colour = gender_f)) + geom_density(alpha=0.2)+
  ggtitle("Age")+theme(legend.position="none")
grid.arrange(p1, p2, ncol = 2)
```



The **desired** variable is a measure of the participant's recorded desire to drink, with values ranging between 1 and 8, a mean of 4.5 and a fairly symmetric distribution. with a significant share of responses at minimum and maximum values. Average values are slightly higher on average for males than for females and are highest on Friday and Saturday and lowest on Sunday and Monday. The **state** variable is a record of the participant's state (short-term) self-esteem as it varies each day. This ranges between 2 and 5 with a mean of 3.97 and a moderately negative-skewed distribution, little difference between males and females and little difference between days of the week (although with more negative outliers on Mondays). The variables **desired** and **state** both contain three missing values, two of these being from the same observation.

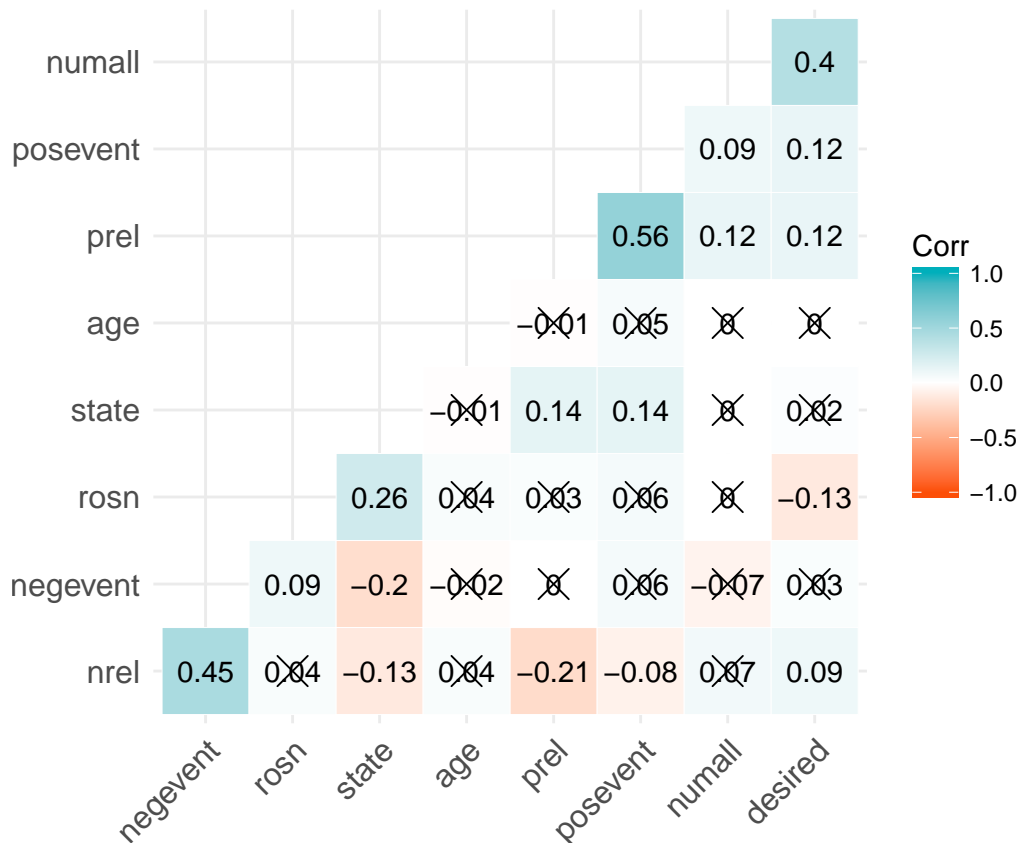
```
# aggregate(desired ~ gender_f, data = dehart, FUN = mean)
# aggregate(desired ~ dayweek_f, data = dehart, FUN = mean)
# aggregate(state ~ gender_f, data = dehart, FUN = mean)
# aggregate(state ~ dayweek_f, data = dehart, FUN = mean)
p1<-ggplot(na.omit(dehart), aes(gender_f, desired)) + geom_boxplot(aes(fill = gender_f)) + labs(x = "")
  ggtitle("Desire to\ndrink") + theme(legend.position="none") + coord_flip()
p2<-ggplot(na.omit(dehart), aes(dayweek_f, desired)) + geom_boxplot(aes(fill = dayweek_f)) + labs(x = "")
  ggtitle("Desire to\ndrink") + theme(legend.position="none") + coord_flip()
p3<-ggplot(na.omit(dehart), aes(gender_f, state)) + geom_boxplot(aes(fill = gender_f)) + labs(x = "") +
  ggtitle("State\nSelf-Esteem") + theme(legend.position="none") + coord_flip()
p4<-ggplot(na.omit(dehart), aes(dayweek_f, state)) + geom_boxplot(aes(fill = dayweek_f)) + labs(x = "")
  ggtitle("State\nSelf-Esteem") + theme(legend.position="none") + coord_flip()
grid.arrange(p1, p2, p3, p4, ncol = 4)
```



Bivariate relationships between these continuous variables are summarized in the below plot of correlation coefficients below. Most of these relationship are weak or insignificant. The strongest positive correlations

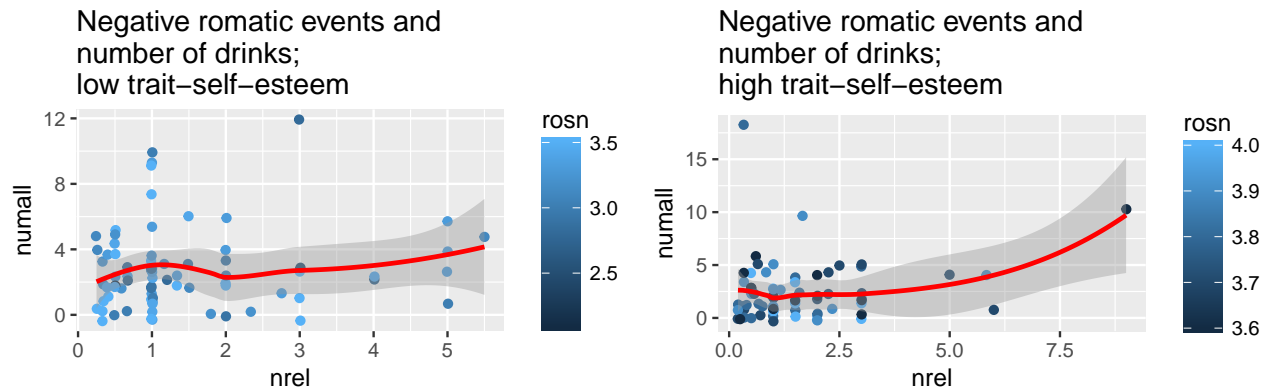
are between **nrel** and **negevent** (negative events and negative romantic events), and between **prel** and **posevent** (positive events and positive romantic events). There is also a moderate positive correlation between the number of drinks and the desire to drink (**numall** and **desired**), and a mild positive correlation between trait and state self-esteem (**rosn** and **state**). State self-esteem has a weak positive correlation to positive event variables and a weak negative correlation to negative event variables, but no correlation with the number of drinks nor the desire to drink. Trait self-esteem has a weak negative correlation with the desire to drink but no correlation with the number of drinks. Age is uncorrelated with any other variable.

```
data <- na.omit(dehart[,c(4,5,6,7,8,10,11,12,13)])
corr <- round(cor(data), 2)
ggcorrplot(corr, p.mat = cor_pmat(data), hc.order = TRUE, type = "lower", color = c("#FC4E07", "white",
```



The heavily skewed distribution for **nrel** indicates that the negative romantic relationship events relevant to the hypothesis are relatively infrequent. A subset of the dataset can be created for participant-days involving a non-zero rating on this variable. The relationship between **nrel** and **numall** can then be compared for participants with below-median trait self-esteem and above-median trait self-esteem. The relationship between negative romantic relationship events and the number of drinks appears to be stronger for individuals with higher trait-self esteem, however the range of this is driven by a relatively small number of individuals recording high **nrel** values that do not appear in the lower trait-self-esteem subset.

```
dehart_nrel = dehart[which(dehart$nrel != 0),]
p1 <- ggplot(na.omit(dehart_nrel[which(dehart_nrel$rosn<=3.5),]), aes(nrel, numall)) + geom_jitter(aes(c
p2 <- ggplot(na.omit(dehart_nrel[which(dehart_nrel$rosn>3.5),]), aes(nrel, numall)) + geom_jitter(aes(c
grid.arrange(p1, p2, ncol = 2)
```



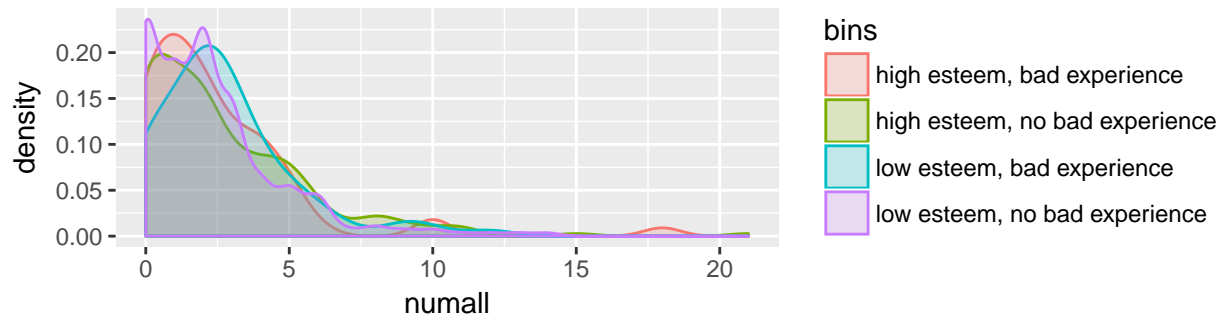
```
dehart$bins <- dehart$rosn
dehart$bins[dehart$rosn <= median(dehart$rosn) & dehart$nrel <= median(dehart$nrel)] = "low esteem, no bad experience"
dehart$bins[dehart$rosn <= median(dehart$rosn) & dehart$nrel > median(dehart$nrel)] = "low esteem, bad experience"
dehart$bins[dehart$rosn > median(dehart$rosn) & dehart$nrel <= median(dehart$nrel)] = "high esteem, no bad experience"
dehart$bins[dehart$rosn > median(dehart$rosn) & dehart$nrel > median(dehart$nrel)] = "high esteem, bad experience"
```

```
p1<- ggplot(na.omit(dehart), aes(numall, fill = bins, colour = bins)) +
  geom_density(alpha=0.2) +
  ggtitle("Self esteem, negative romantic relationship, number of drinks") + ggtitle("Self esteem, negative romantic relationship, number of drinks")

p2<- ggplot(na.omit(dehart), aes(desired, fill = bins, colour = bins)) +
  geom_density(alpha=0.2) +
  ggtitle("Self esteem, negative romantic relationship, number of drinks") + ggtitle("Self esteem, negative romantic relationship, number of drinks")
```

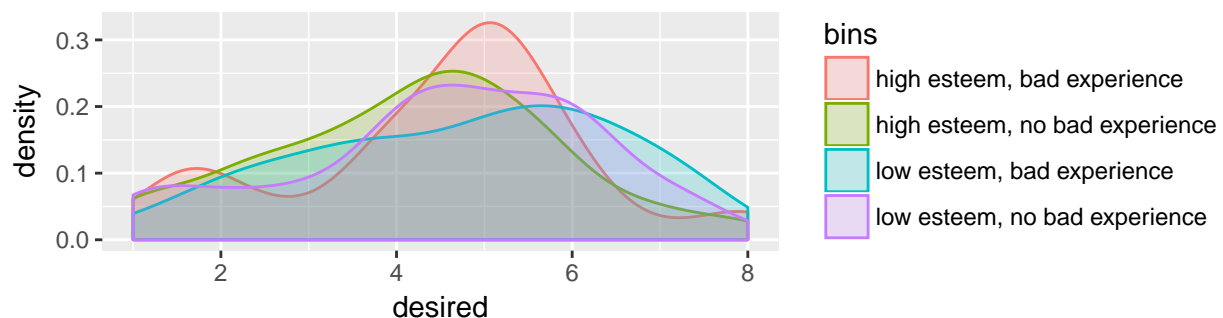
p1

Self esteem, negative romantic relationship, number of drinks



p2

Self esteem, negative romantic relationship, desire to drink




```

# NUMBER OF DRINKS; POISSON MODEL
# significant interaction effect

# BASIC MODEL

N.mod.0 <- glm(formula = numall ~ nrel + rosn, family = poisson(link = "log"), data = na.omit(dehart))
summary(N.mod.0)

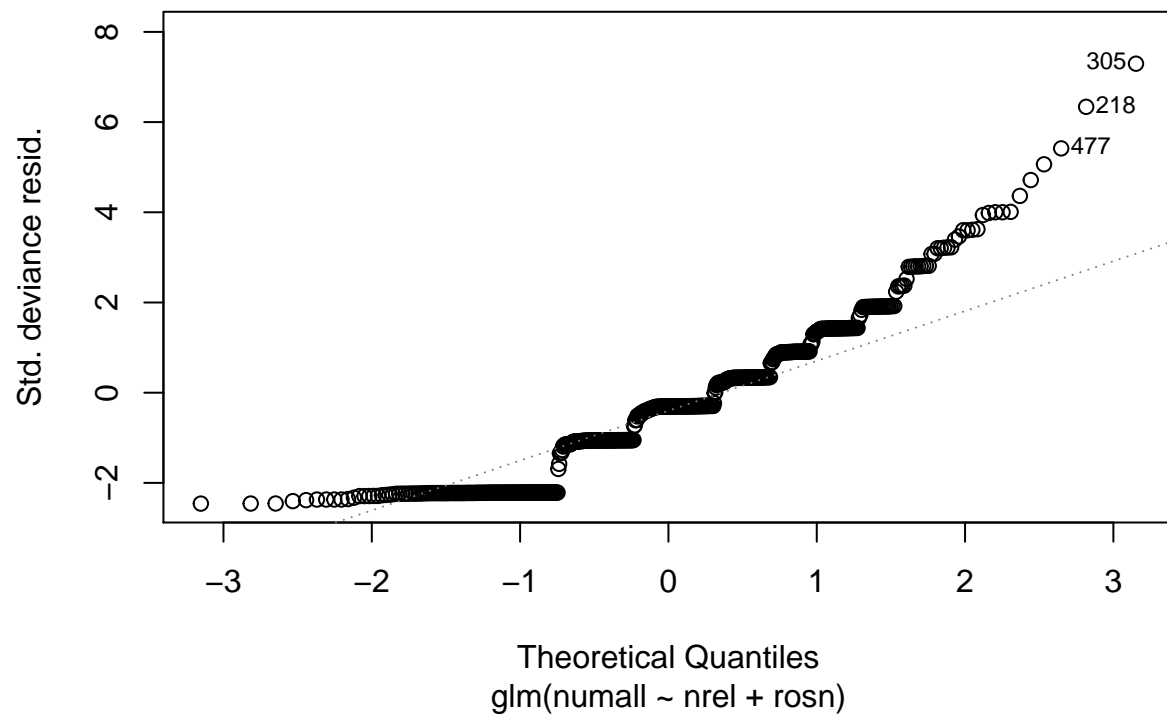
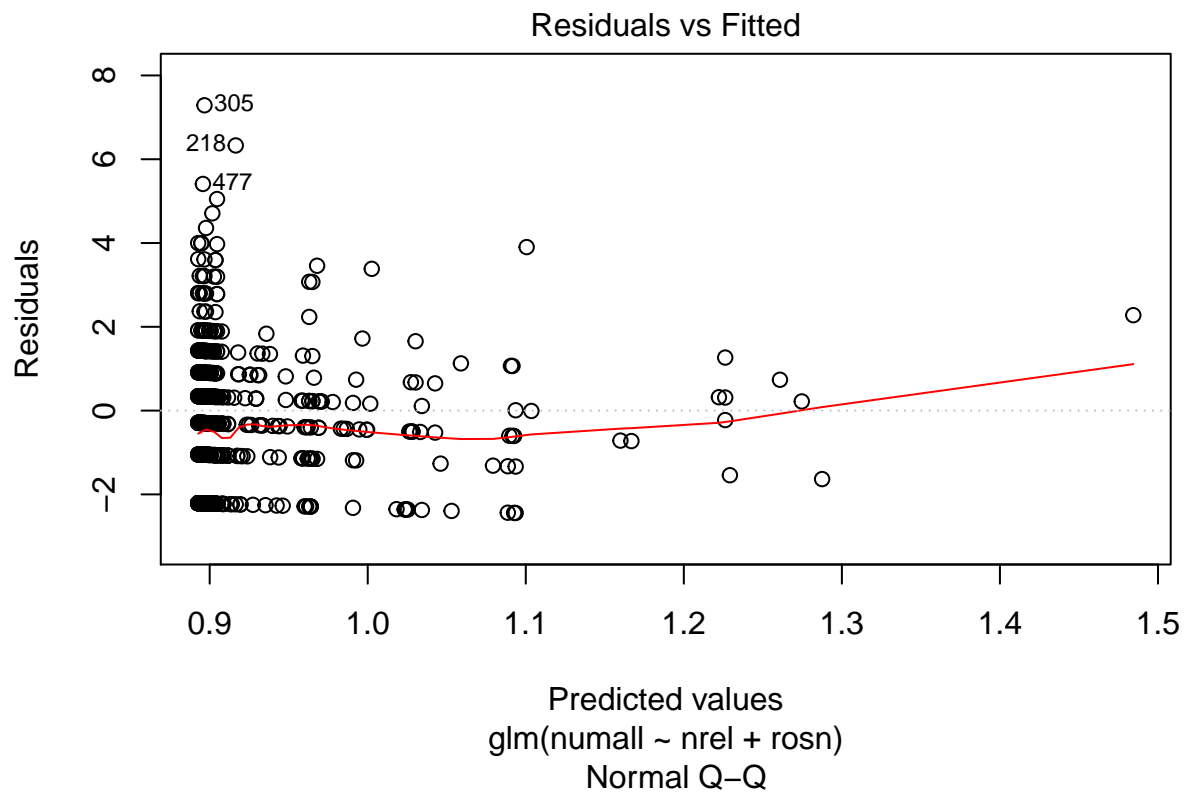
##
## Call:
## glm(formula = numall ~ nrel + rosn, family = poisson(link = "log"),
##      data = na.omit(dehart))
##
## Deviance Residuals:
##      Min       1Q   Median       3Q      Max
## -2.4434  -1.1434  -0.3040   0.3448   7.2878
##
## Coefficients:
##              Estimate Std. Error z value Pr(>|z|)
## (Intercept)  0.932497   0.209136   4.459 8.24e-06 ***
## nrel         0.065298   0.023671   2.759 0.00581 **
## rosn        -0.009943   0.060520  -0.164 0.86951
## ---
## Signif. codes:  0 '***' 0.001 '**' 0.01 '*' 0.05 '.' 0.1 ' ' 1
##
## (Dispersion parameter for poisson family taken to be 1)
##
##      Null deviance: 1583.5  on 617  degrees of freedom
## Residual deviance: 1576.5  on 615  degrees of freedom
## AIC: 2944.5
##
## Number of Fisher Scoring iterations: 5

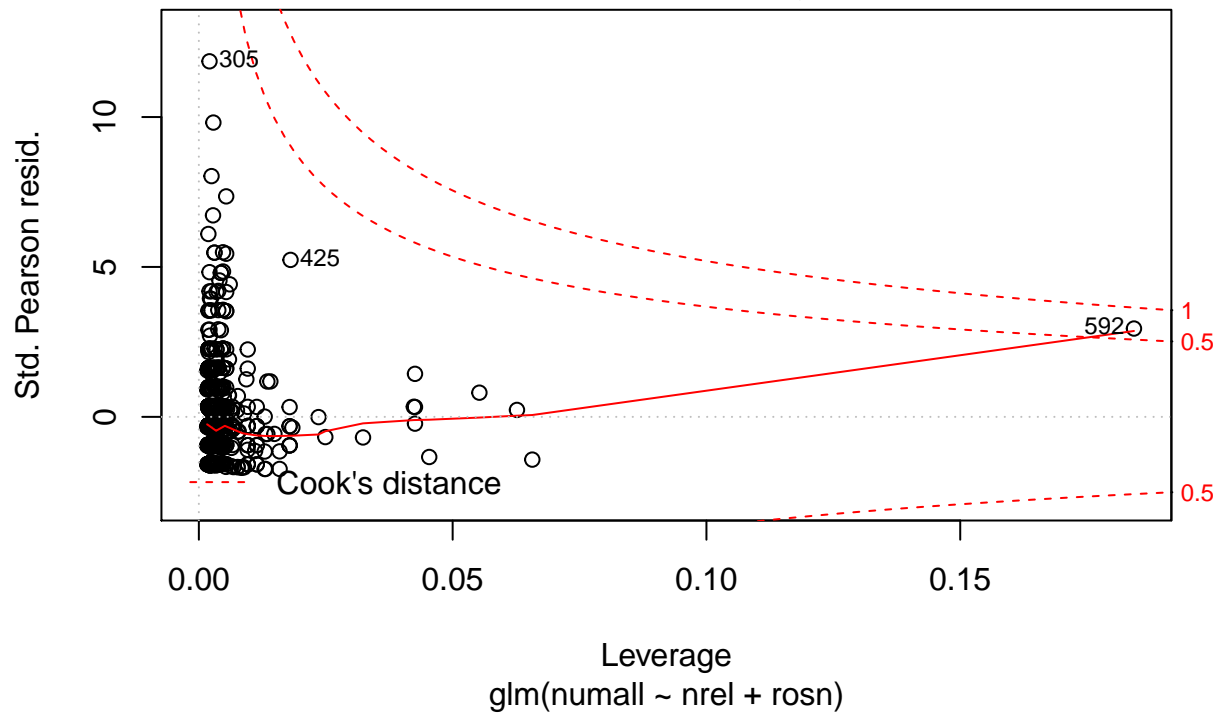
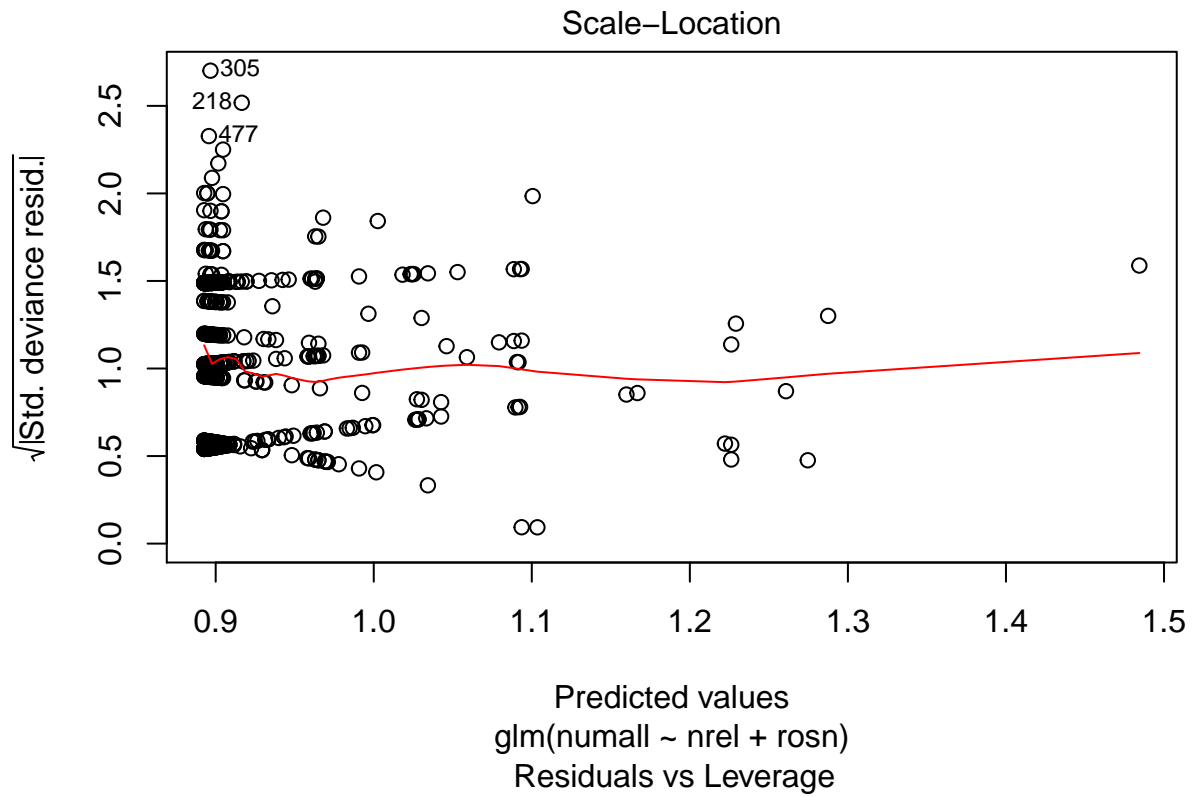
# hypothesis tests for statistical significance
Anova(N.mod.0, test = "LR")

## Analysis of Deviance Table (Type II tests)
##
## Response: numall
##      LR Chisq Df Pr(>Chisq)
## nrel   6.9884  1  0.008204 **
## rosn    0.0270  1  0.869573
## ---
## Signif. codes:  0 '***' 0.001 '**' 0.01 '*' 0.05 '.' 0.1 ' ' 1

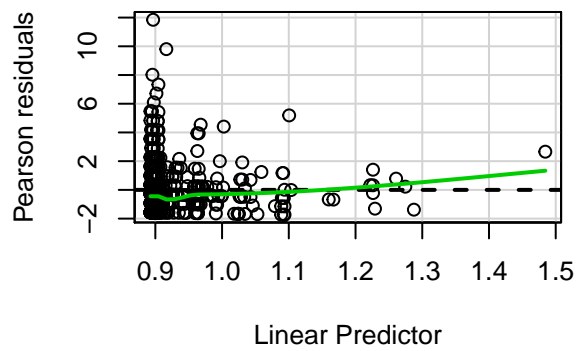
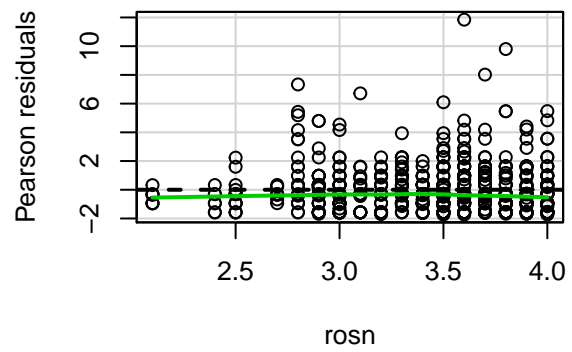
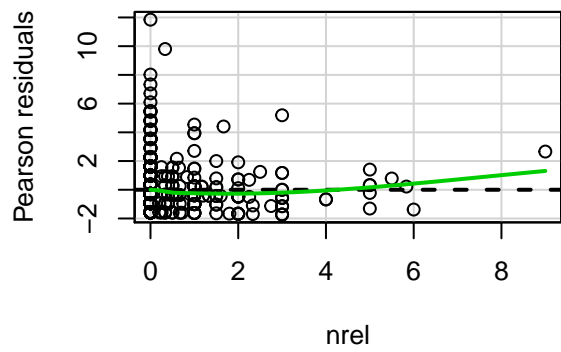
# diagnostic plots
plot(N.mod.0)

```



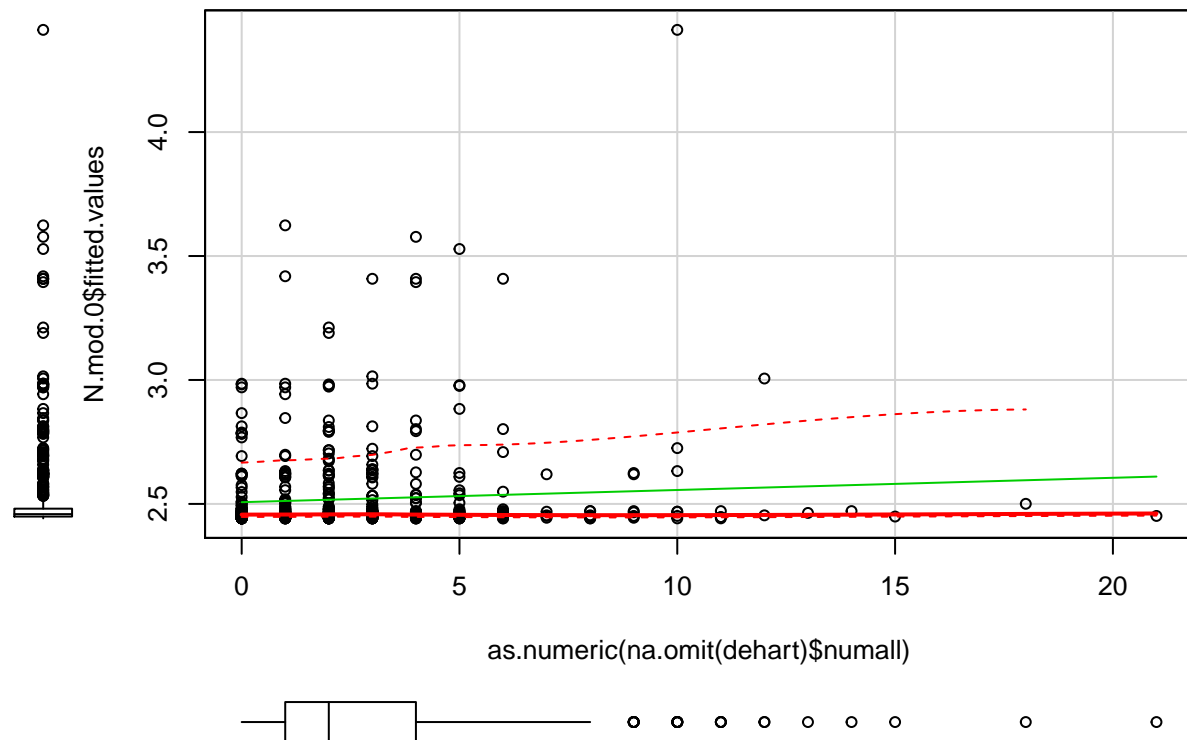


```
# Pearson residuals
residualPlots(N.mod.0)
```



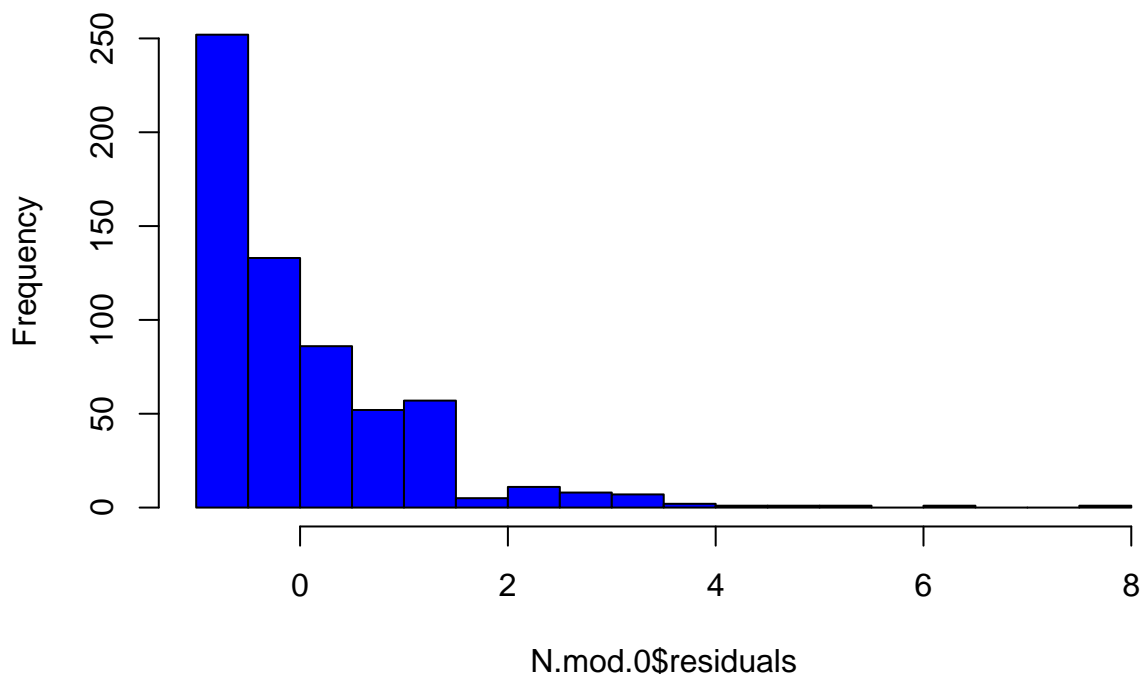
```
##      Test stat Pr(>|t|)
## nrel      7.028  0.008
## rosn      2.364  0.124
```

```
# scatterplot of actual and fitted values
scatterplot(as.numeric(na.omit(dehart)$numall), N.mod.0$fitted.values)
```



```
# histogram of residuals
hist(N.mod.0$residuals, breaks = 20, col = "blue")
```

Histogram of N.mod.0\$residuals



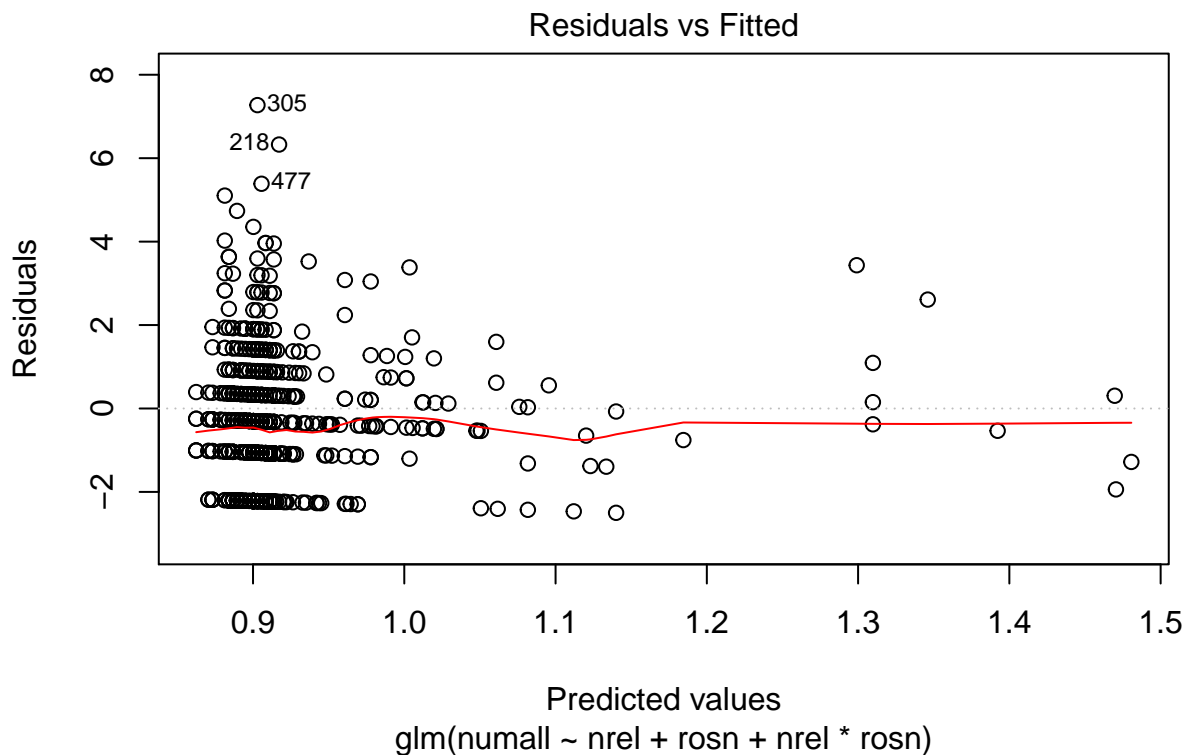
```
N.mod.1 <- glm(formula = numall ~ nrel + rosn + nrel*rosn, family = poisson(link = "log"), data = na.omit(dehart))
summary(N.mod.1)
```

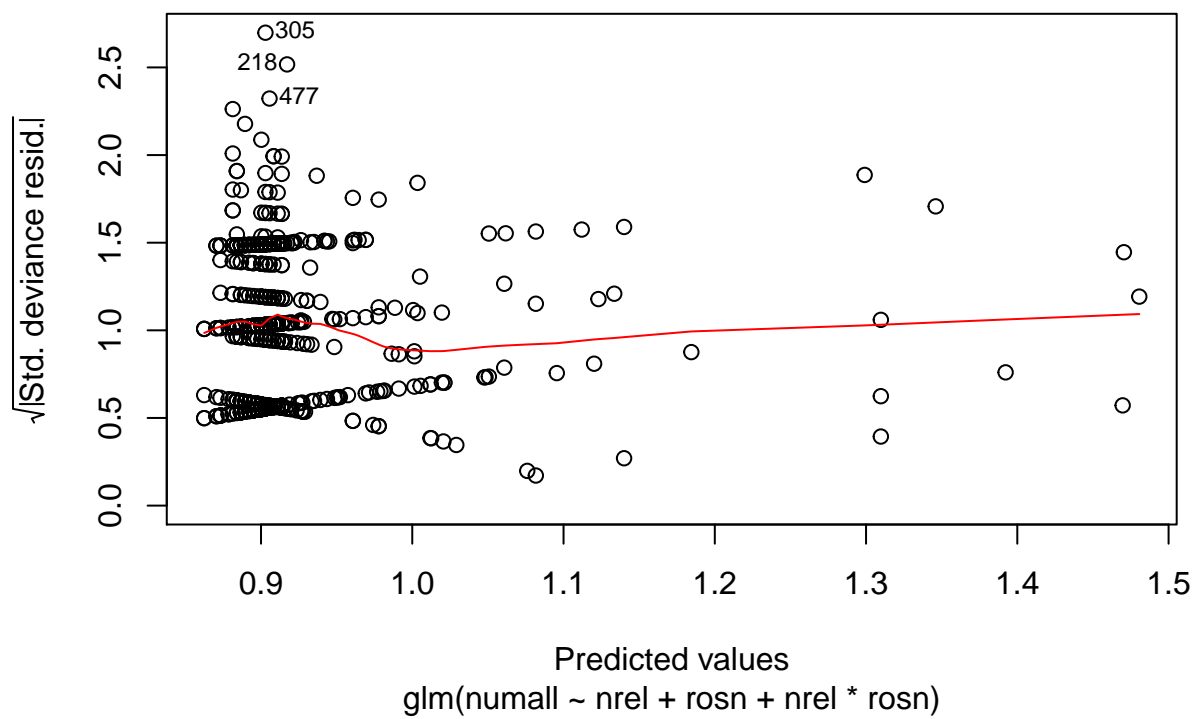
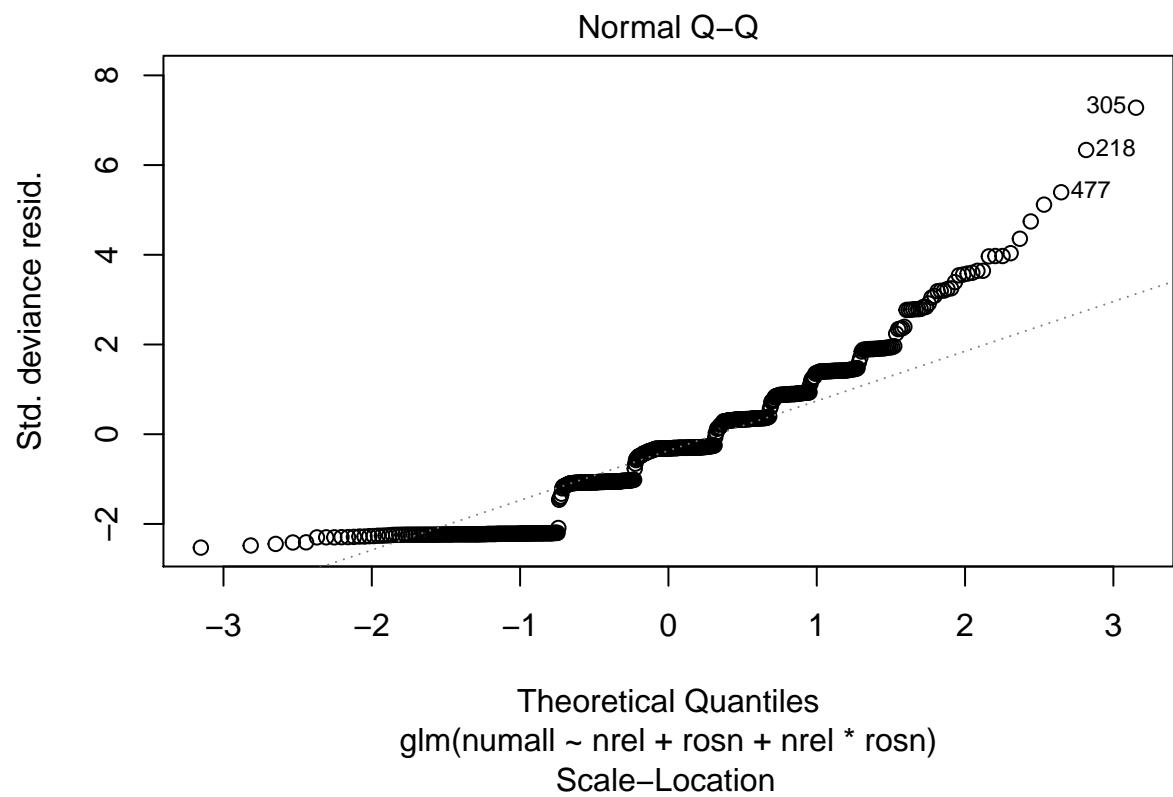
```
##
## Call:
## glm(formula = numall ~ nrel + rosn + nrel * rosn, family = poisson(link = "log"),
##      data = na.omit(dehart))
##
## Deviance Residuals:
##      Min       1Q   Median       3Q      Max
## -2.5007  -1.0966  -0.3156   0.3795   7.2719
##
## Coefficients:
##              Estimate Std. Error z value Pr(>|z|)
## (Intercept)  0.80573    0.22392   3.598  0.00032 ***
## nrel         0.45430    0.23218   1.957  0.05039 .
## rosn         0.02700    0.06477   0.417  0.67682
## nrel:rosn    -0.11252    0.06722  -1.674  0.09416 .
## ---
## Signif. codes:  0 '***' 0.001 '**' 0.01 '*' 0.05 '.' 0.1 ' ' 1
##
## (Dispersion parameter for poisson family taken to be 1)
##
##      Null deviance: 1583.5  on 617  degrees of freedom
## Residual deviance: 1573.8  on 614  degrees of freedom
## AIC: 2943.7
##
```

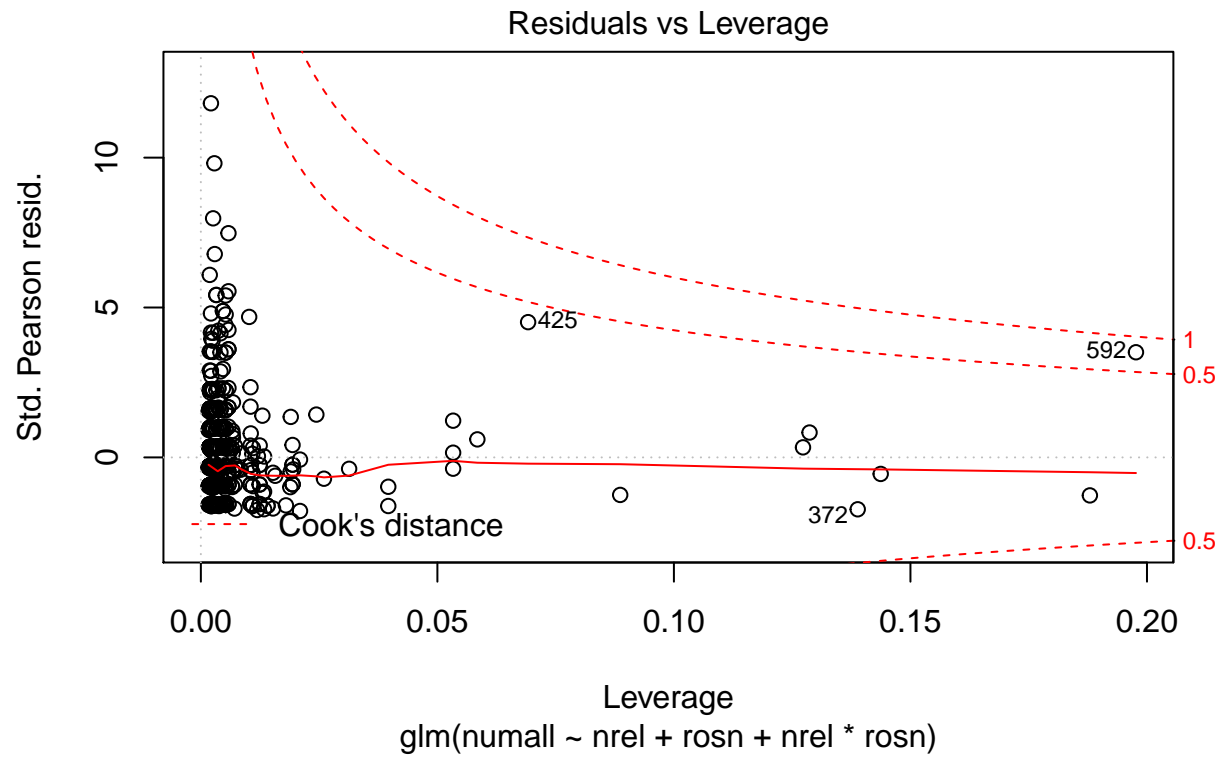
```
## Number of Fisher Scoring iterations: 5
# hypothesis tests for statistical significance
Anova(N.mod.1, test = "LR")

## Analysis of Deviance Table (Type II tests)
##
## Response: numall
##          LR Chisq Df Pr(>Chisq)
## nrel      6.9884  1  0.008204 **
## rosn      0.0270  1  0.869573
## nrel:rosn  2.7409  1  0.097809 .
## ---
## Signif. codes:  0 '***' 0.001 '**' 0.01 '*' 0.05 '.' 0.1 ' ' 1

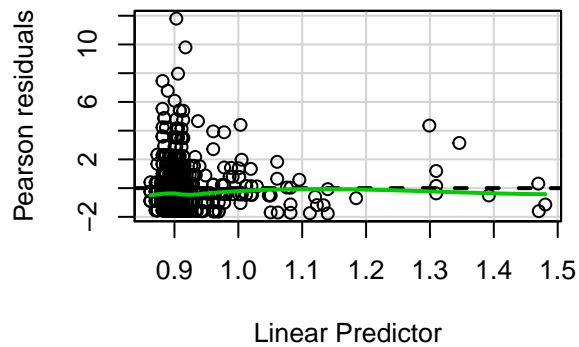
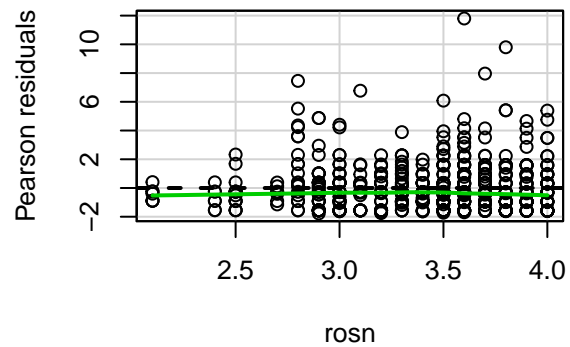
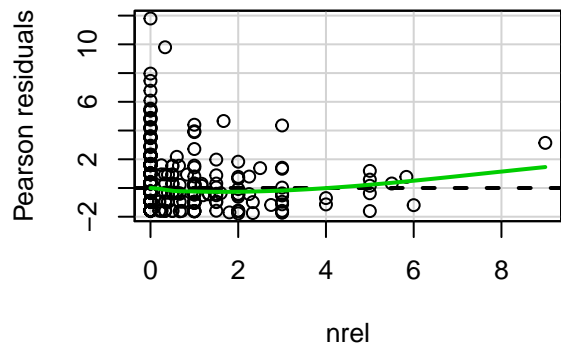
# diagnostic plots
plot(N.mod.1)
```







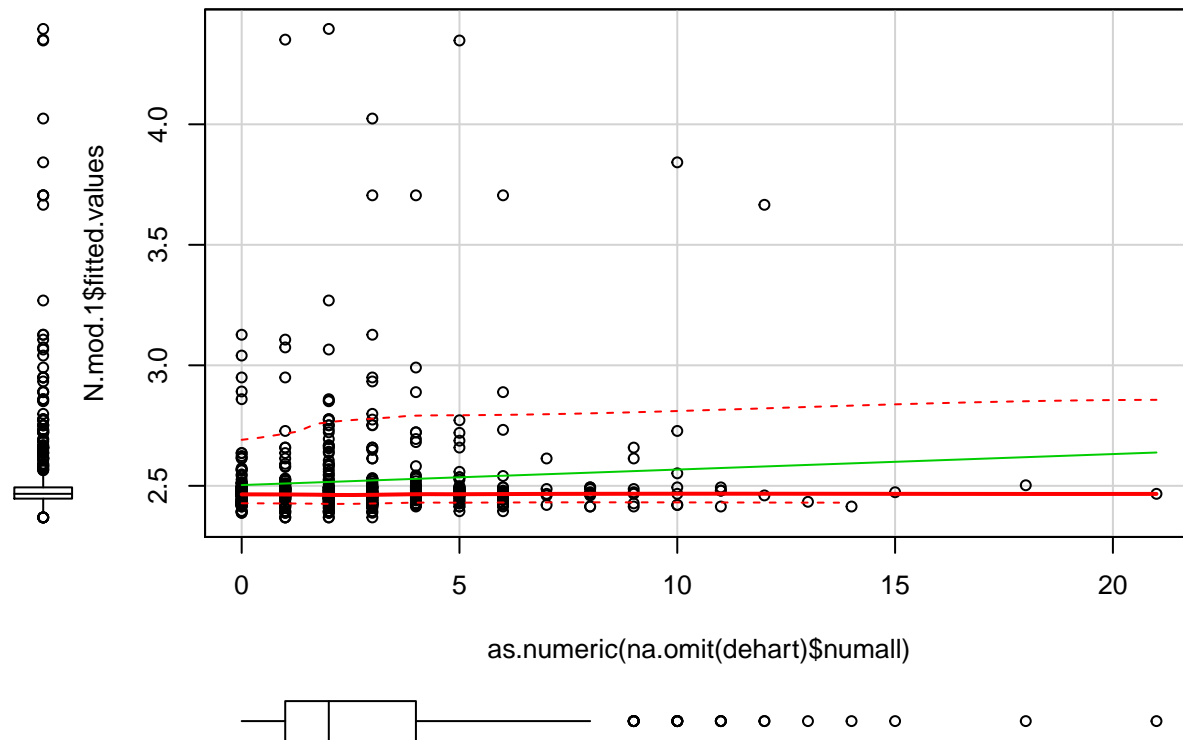
```
# Pearson residuals
residualPlots(N.mod.1)
```



```
##      Test stat Pr(>|t|)
## nrel      8.826   0.003
## rosn      1.934   0.164
```

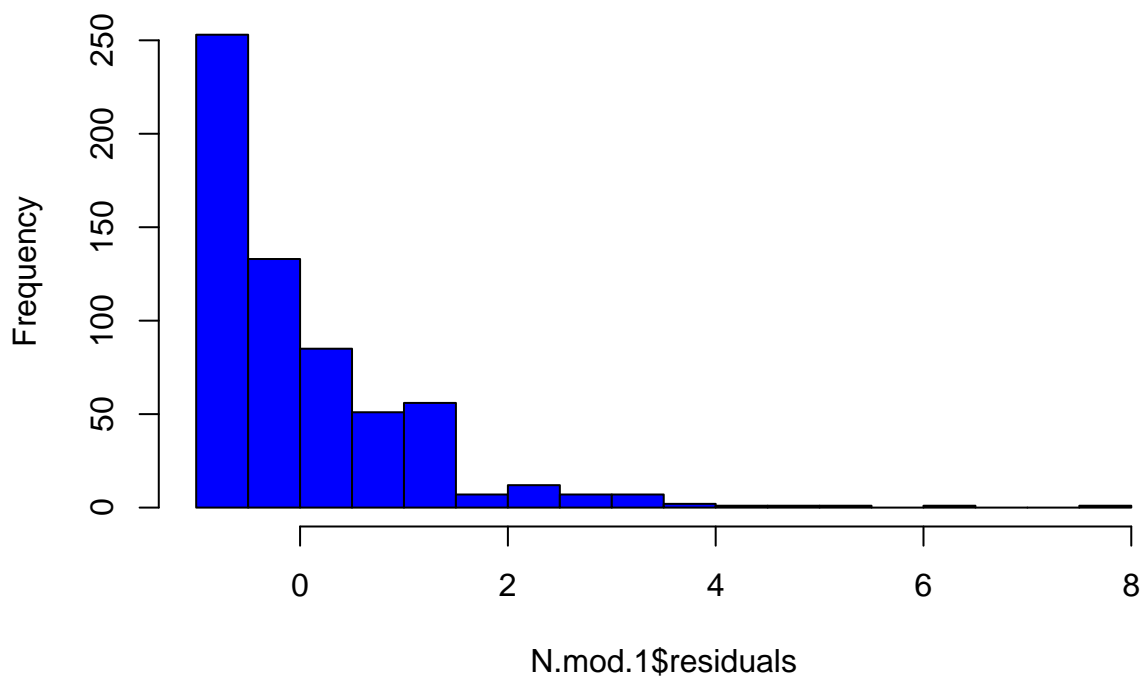


```
# scatterplot of actual and fitted values
scatterplot(as.numeric(na.omit(dehart)$numall), N.mod.1$fitted.values)
```



```
# histogram of residuals
hist(N.mod.1$residuals, breaks = 20, col = "blue")
```

Histogram of N.mod.1\$residuals



```

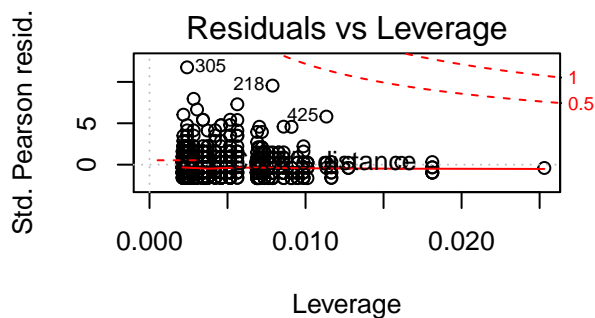
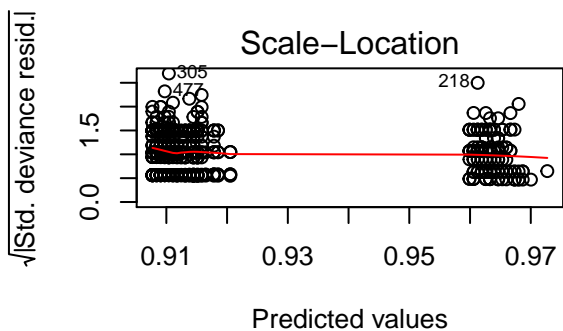
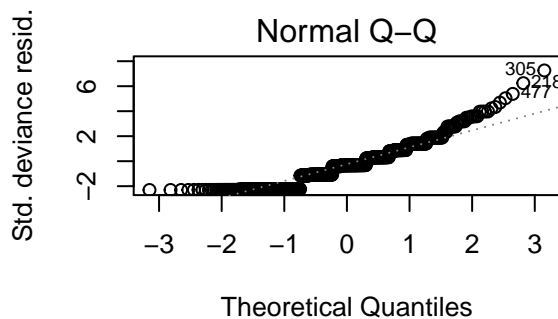
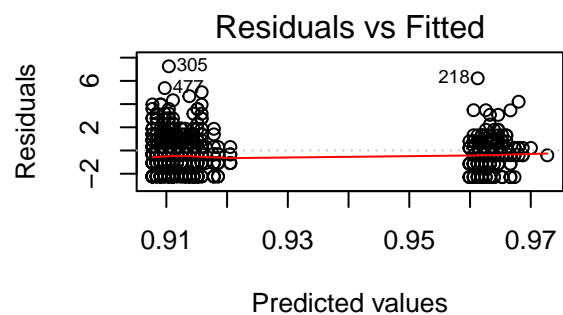
# AGAIN WITH FACTOR FOR NREL
dehart$nrel_f <- dehart$nrel
dehart$nrel_f[dehart$nrel_f != 0] = 1
dehart$nrel_f <- factor(dehart$nrel_f)
levels(dehart$nrel_f) <- c("low", "high")

N.mod.Oa <- glm(formula = numall ~ nrel_f + rosn, family = poisson(link = "log"), data = na.omit(dehart),
summary(N.mod.Oa)

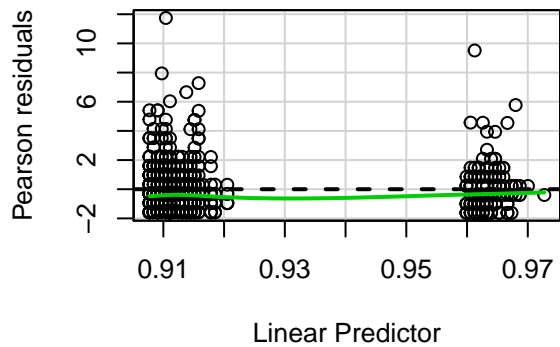
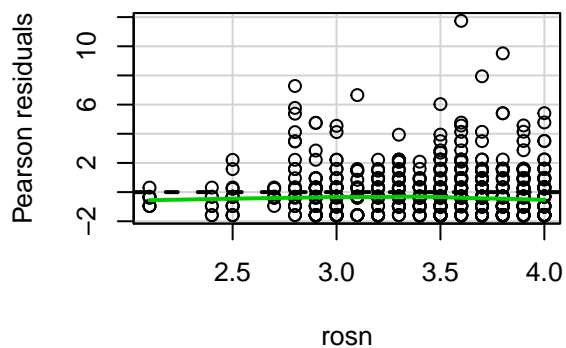
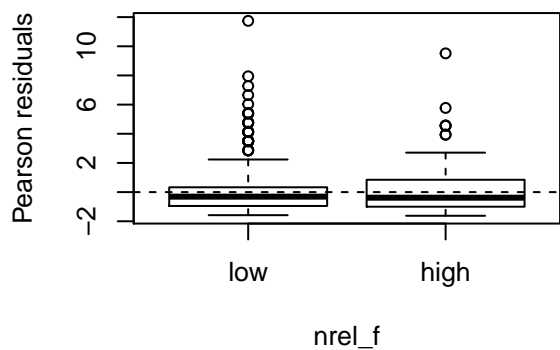
##
## Call:
## glm(formula = numall ~ nrel_f + rosn, family = poisson(link = "log"),
##      data = na.omit(dehart))
##
## Deviance Residuals:
##      Min       1Q   Median       3Q      Max
## -2.2939  -1.1442  -0.3229   0.6693   7.2528
##
## Coefficients:
##              Estimate Std. Error z value Pr(>|z|)
## (Intercept)  0.934688   0.208906   4.474 7.67e-06 ***
## nrel_fhigh    0.052194   0.059103   0.883   0.377
## rosn         -0.006743   0.060503  -0.111   0.911
## ---
## Signif. codes:  0 '***' 0.001 '**' 0.01 '*' 0.05 '.' 0.1 ' ' 1
##
## (Dispersion parameter for poisson family taken to be 1)
##
##      Null deviance: 1583.5  on 617  degrees of freedom
## Residual deviance: 1582.8  on 615  degrees of freedom
## AIC: 2950.7
##
## Number of Fisher Scoring iterations: 5
# hypothesis tests for statistical significance
Anova(N.mod.Oa, test = "LR")

## Analysis of Deviance Table (Type II tests)
##
## Response: numall
##           LR Chisq Df Pr(>Chisq)
## nrel_f    0.77295  1    0.3793
## rosn      0.01241  1    0.9113
# diagnostic plots
par(mfrow=c(2,2))
plot(N.mod.Oa)

```

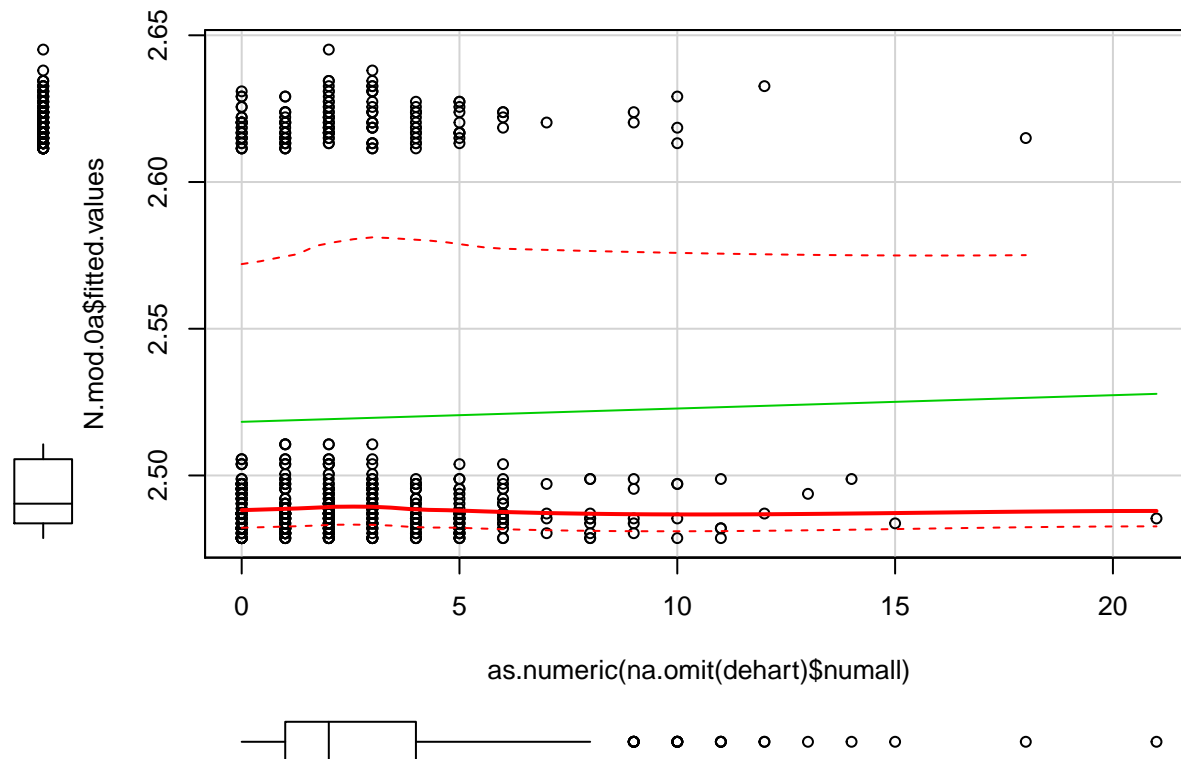


```
# Pearson residuals
residualPlots(N.mod.0a)
```



```
##      Test stat Pr(>|t|)
## nrel_f      NA      NA
## rosn      2.591  0.107
```

```
# scatterplot of actual and fitted values
scatterplot(as.numeric(na.omit(dehart)$numall), N.mod.0a$fitted.values)
```



```
# histogram of residuals
hist(N.mod.0a$residuals, breaks = 20, col = "blue")
```

```
N.mod.1a <- glm(formula = numall ~ nrel_f + rosn + nrel_f*rosn, family = poisson(link = "log"), data = na.omit(dehart))
summary(N.mod.1a)
```

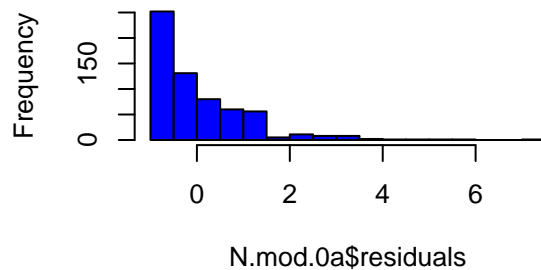
```
##
## Call:
## glm(formula = numall ~ nrel_f + rosn + nrel_f * rosn, family = poisson(link = "log"),
##      data = na.omit(dehart))
##
## Deviance Residuals:
##      Min       1Q   Median       3Q      Max
## -2.4504  -1.1137  -0.3339   0.5868   7.2275
##
## Coefficients:
##              Estimate Std. Error z value Pr(>|z|)
## (Intercept)    0.74179    0.23394   3.171  0.00152 **
## nrel_fhigh     1.06068    0.52477   2.021  0.04326 *
## rosn           0.04959    0.06768   0.733  0.46375
## nrel_fhigh:rosn -0.29204    0.15154  -1.927  0.05397 .
## ---
## Signif. codes:  0 '***' 0.001 '**' 0.01 '*' 0.05 '.' 0.1 ' ' 1
##
## (Dispersion parameter for poisson family taken to be 1)
##
```

```
##      Null deviance: 1583.5  on 617  degrees of freedom
## Residual deviance: 1579.1  on 614  degrees of freedom
## AIC: 2949
##
## Number of Fisher Scoring iterations: 5
# hypothesis tests for statistical significance
Anova(N.mod.1a, test = "LR")

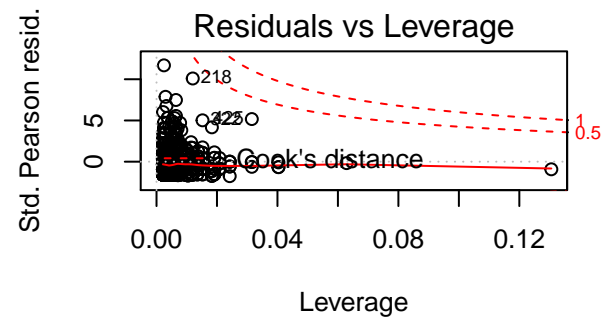
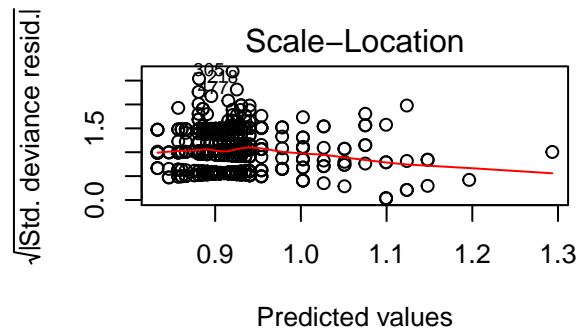
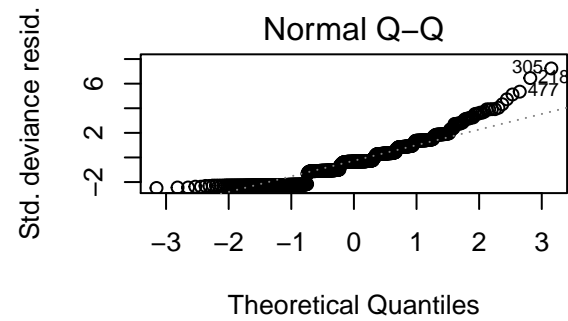
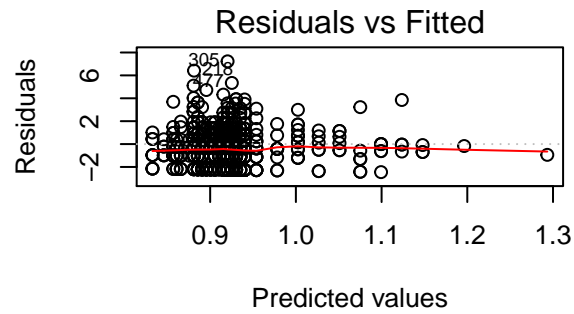
## Analysis of Deviance Table (Type II tests)
##
## Response: numall
##           LR Chisq Df Pr(>Chisq)
## nrel_f      0.7729  1   0.37931
## rosn        0.0124  1   0.91129
## nrel_f:rosn  3.6538  1   0.05594 .
## ---
## Signif. codes:  0 '***' 0.001 '**' 0.01 '*' 0.05 '.' 0.1 ' ' 1

# diagnostic plots
par(mfrow=c(2,2))
```

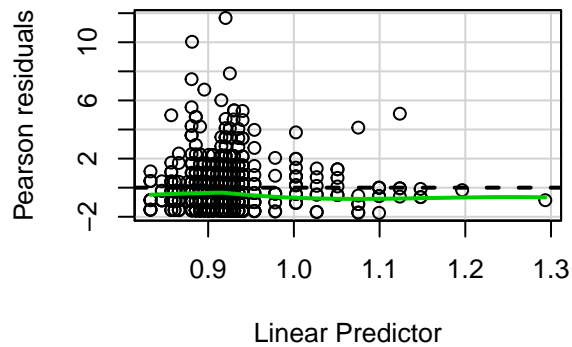
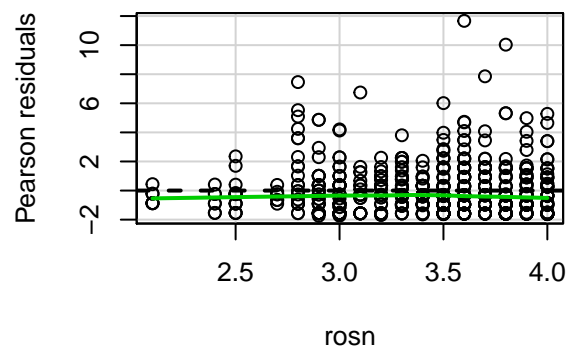
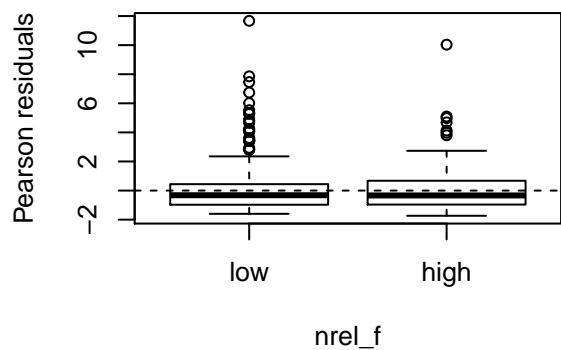
Histogram of N.mod.0a\$residuals



```
plot(N.mod.1a)
```

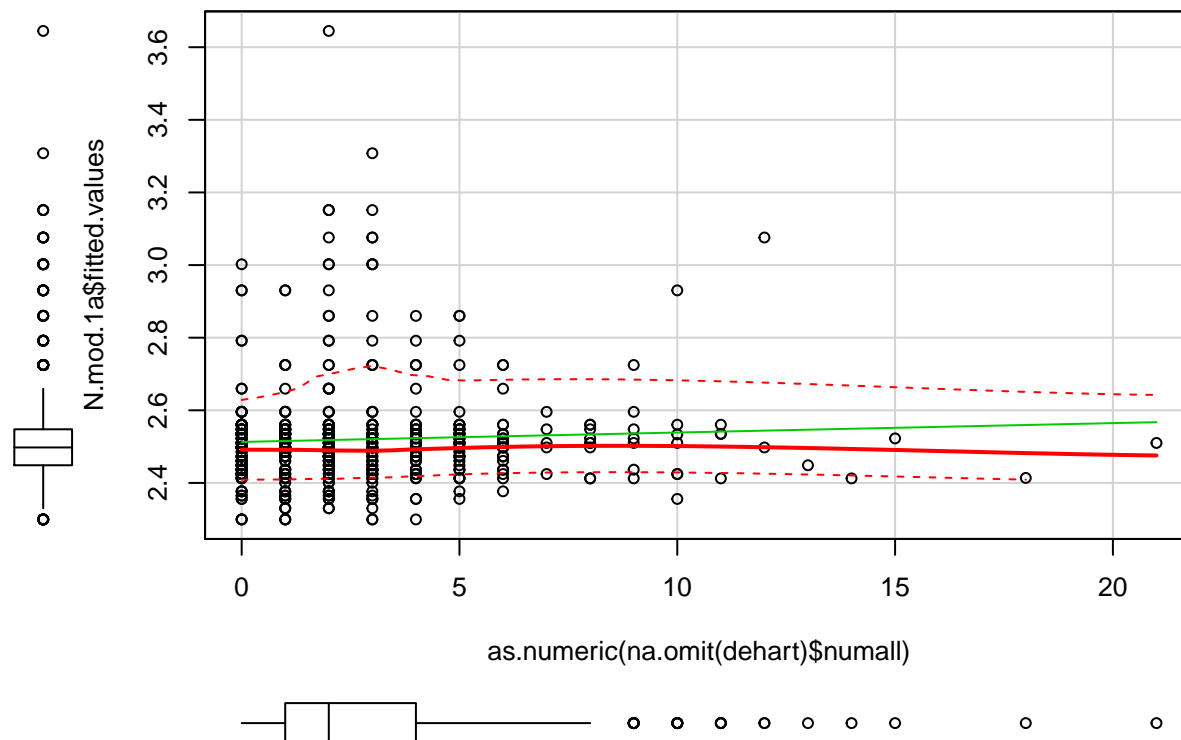


```
# Pearson residuals
residualPlots(N.mod.1a)
```



```
##      Test stat Pr(>|t|)
## nrel_f      NA      NA
## rosn      2.118  0.146
```

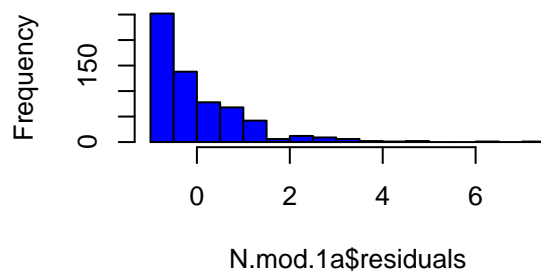
```
# scatterplot of actual and fitted values
scatterplot(as.numeric(na.omit(dehart)$numall), N.mod.1a$fitted.values)
```



```
# histogram of residuals
hist(N.mod.1a$residuals, breaks = 20, col = "blue")
```

little significance for anything

Histogram of N.mod.1a\$residuals



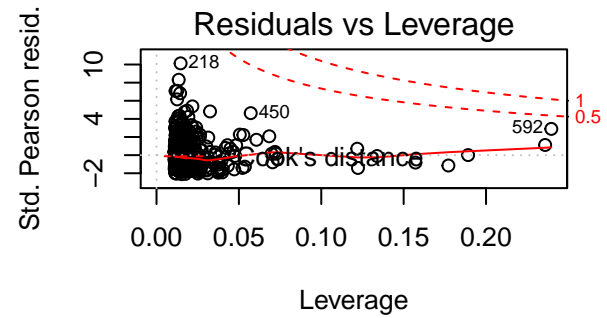
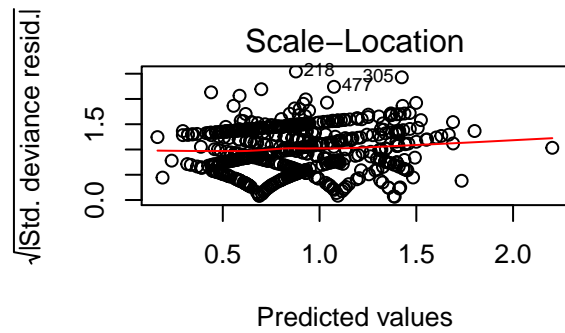
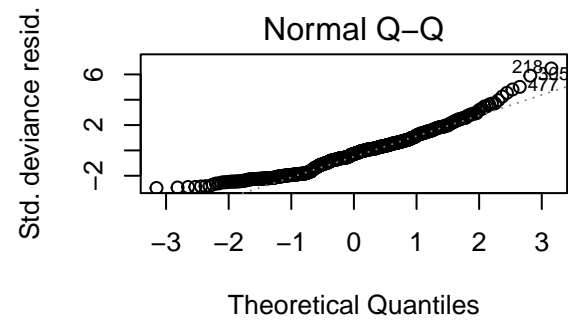
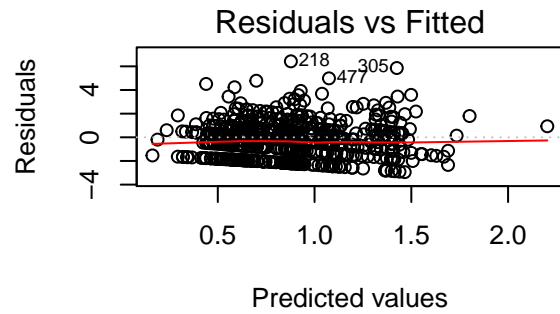
```
# INTERMEDIATE MODEL; DAYS OF THE WEEK
```

```
N.mod.2 <- glm(formula = numall ~ nrel + prel + rosn + nrel*rosn + prel*rosn + dayweek_f, family = poisson)
summary(N.mod.2)
```

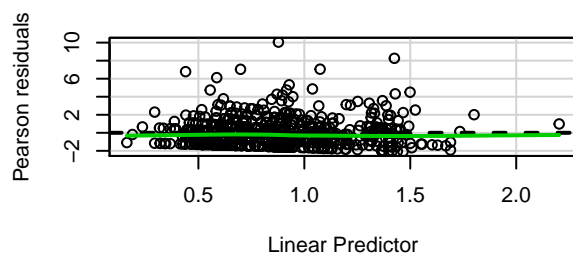
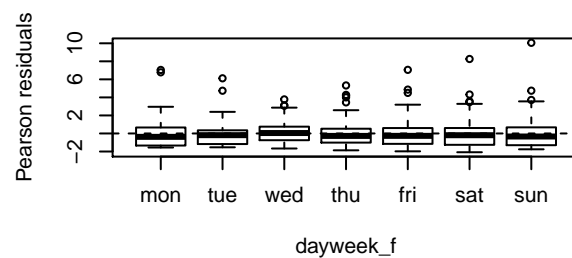
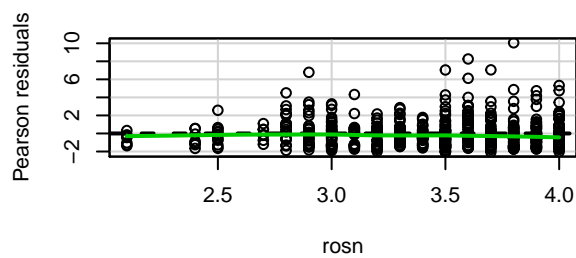
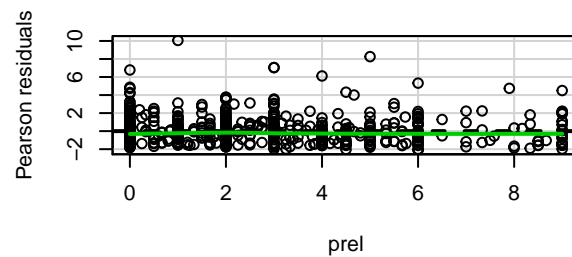
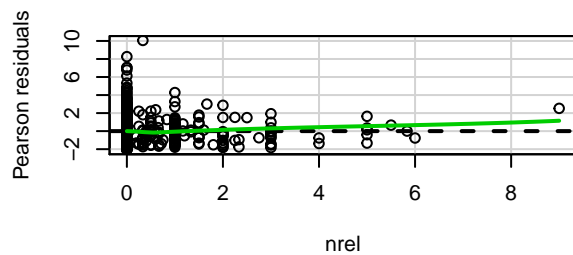
```
##
## Call:
## glm(formula = numall ~ nrel + prel + rosn + nrel * rosn + prel *
##      rosn + dayweek_f, family = poisson(link = "log"), data = na.omit(dehart))
```

```
##
## Deviance Residuals:
##      Min       1Q   Median       3Q      Max
## -2.9432  -1.5586  -0.2800   0.5938   6.4240
##
## Coefficients:
##              Estimate Std. Error z value Pr(>|z|)
## (Intercept)  -0.32074    0.35651  -0.900  0.368296
## nrel          0.75464    0.23851   3.164  0.001556 **
## prel          0.26556    0.07484   3.549  0.000387 ***
## rosn          0.26221    0.10174   2.577  0.009955 **
## dayweek_ftue -0.14614    0.11041  -1.324  0.185635
## dayweek_fwed -0.09892    0.10809  -0.915  0.360100
## dayweek_ftthu 0.19492    0.10104   1.929  0.053711 .
## dayweek_ffri  0.36276    0.09749   3.721  0.000199 ***
## dayweek_fsat  0.66487    0.09225   7.207  5.7e-13 ***
## dayweek_fsun  0.18081    0.10180   1.776  0.075714 .
## nrel:rosn     -0.19302    0.06875  -2.808  0.004991 **
## prel:rosn     -0.06616    0.02216  -2.986  0.002830 **
## ---
## Signif. codes:  0 '***' 0.001 '**' 0.01 '*' 0.05 '.' 0.1 ' ' 1
##
## (Dispersion parameter for poisson family taken to be 1)
##
##      Null deviance: 1583.5  on 617  degrees of freedom
## Residual deviance: 1416.1  on 606  degrees of freedom
## AIC: 2802
##
## Number of Fisher Scoring iterations: 5
# hypothesis tests for statistical significance
Anova(N.mod.2, test = "LR")

## Analysis of Deviance Table (Type II tests)
##
## Response: numall
##           LR Chisq Df Pr(>Chisq)
## nrel         10.510  1  0.001187 **
## prel         16.789  1  4.178e-05 ***
## rosn          0.089  1  0.765921
## dayweek_f    116.305  6 < 2.2e-16 ***
## nrel:rosn      7.642  1  0.005702 **
## prel:rosn      8.807  1  0.003000 **
## ---
## Signif. codes:  0 '***' 0.001 '**' 0.01 '*' 0.05 '.' 0.1 ' ' 1
# diagnostic plots
par(mfrow=c(2,2))
plot(N.mod.2)
```

```
# Pearson residuals
residualPlots(N.mod.2)
```

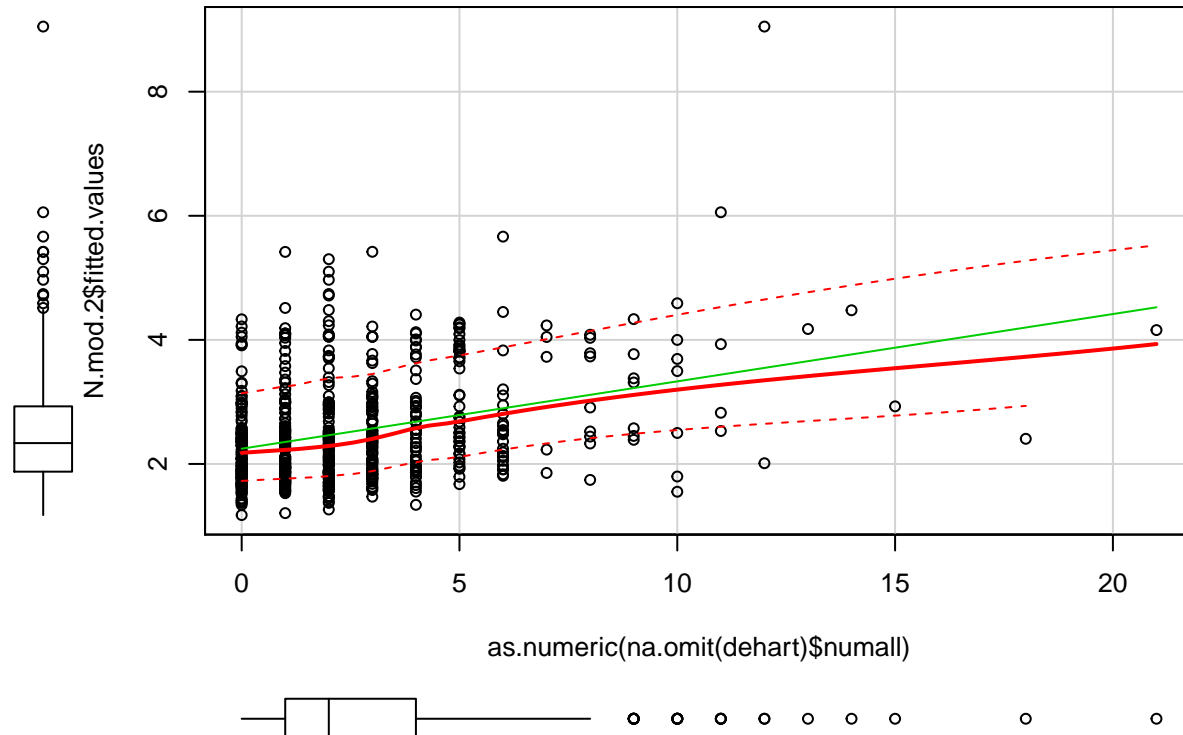


```
##          Test stat Pr(>|t|)
## nrel      6.696    0.010
```

```
## prel          0.000    0.996
## rosn          2.867    0.090
## dayweek_f      NA      NA
```

```
# scatterplot of actual and fitted values
```

```
scatterplot(as.numeric(na.omit(dehart)$numall), N.mod.2$fitted.values)
```



```
# histogram of residuals
```

```
hist(N.mod.2$residuals, breaks = 20, col = "blue")
```

```
# AGAIN WITH FACTOR FOR NREL
```

```
N.mod.2a <- glm(formula = numall ~ nrel + prel + rosn + nrel*rosn + prel*rosn + dayweek_f, family = poisson)
summary(N.mod.2a)
```

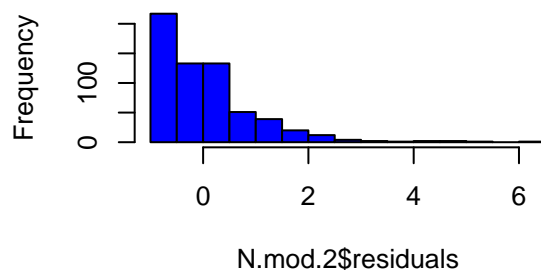
```
##
## Call:
## glm(formula = numall ~ nrel + prel + rosn + nrel * rosn + prel *
##      rosn + dayweek_f, family = poisson(link = "log"), data = na.omit(dehart))
##
## Deviance Residuals:
##      Min       1Q   Median       3Q      Max
## -2.9432  -1.5586  -0.2800   0.5938   6.4240
##
## Coefficients:
##              Estimate Std. Error z value Pr(>|z|)
## (Intercept)  -0.32074    0.35651  -0.900  0.368296
## nrel           0.75464    0.23851   3.164  0.001556 **
## prel          0.26556    0.07484   3.549  0.000387 ***
## rosn          0.26221    0.10174   2.577  0.009955 **
## dayweek_ftue -0.14614    0.11041  -1.324  0.185635
```

```
## dayweek_fwed -0.09892    0.10809   -0.915  0.360100
## dayweek_fthu  0.19492    0.10104    1.929  0.053711 .
## dayweek_ffri  0.36276    0.09749    3.721  0.000199 ***
## dayweek_fsat  0.66487    0.09225    7.207  5.7e-13 ***
## dayweek_fsun  0.18081    0.10180    1.776  0.075714 .
## nrel:rosl      -0.19302    0.06875   -2.808  0.004991 **
## prel:rosl      -0.06616    0.02216   -2.986  0.002830 **
## ---
## Signif. codes:  0 '***' 0.001 '**' 0.01 '*' 0.05 '.' 0.1 ' ' 1
##
## (Dispersion parameter for poisson family taken to be 1)
##
## Null deviance: 1583.5  on 617  degrees of freedom
## Residual deviance: 1416.1  on 606  degrees of freedom
## AIC: 2802
##
## Number of Fisher Scoring iterations: 5
# hypothesis tests for statistical significance
Anova(N.mod.2a, test = "LR")
```

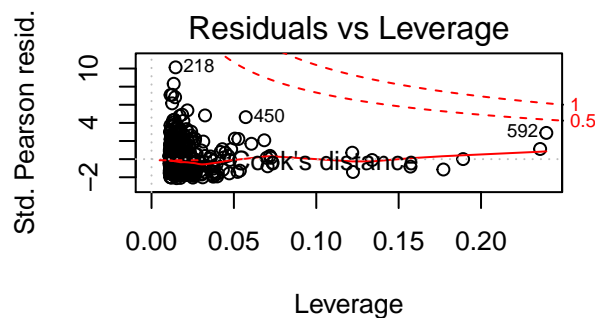
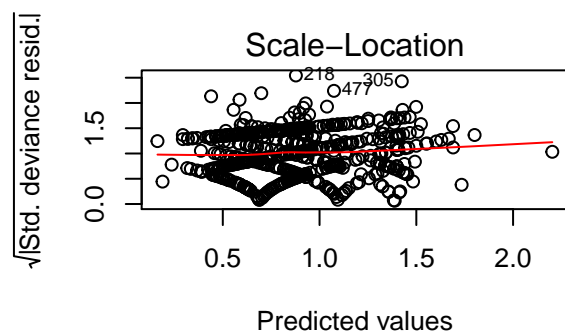
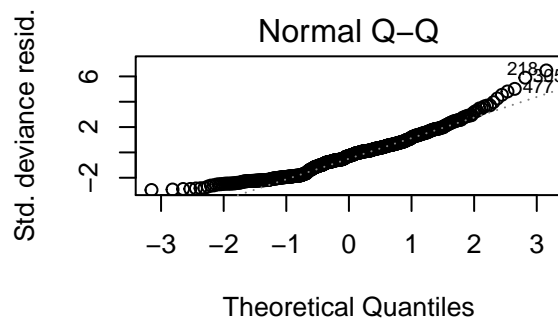
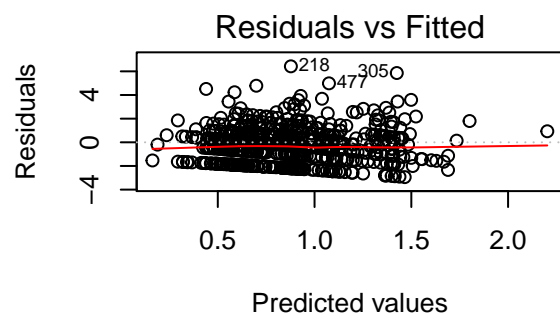
```
## Analysis of Deviance Table (Type II tests)
##
## Response: numall
##          LR Chisq Df Pr(>Chisq)
## nrel         10.510  1  0.001187 **
## prel         16.789  1  4.178e-05 ***
## rosl          0.089  1  0.765921
## dayweek_f    116.305  6 < 2.2e-16 ***
## nrel:rosl      7.642  1  0.005702 **
## prel:rosl      8.807  1  0.003000 **
## ---
## Signif. codes:  0 '***' 0.001 '**' 0.01 '*' 0.05 '.' 0.1 ' ' 1
```

```
# diagnostic plots
par(mfrow=c(2,2))
```

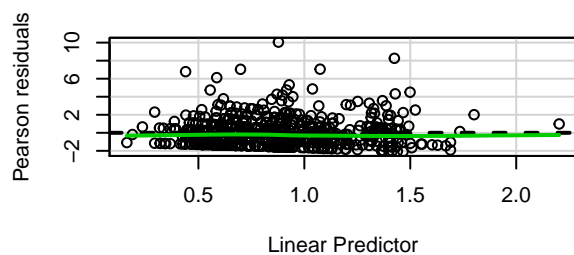
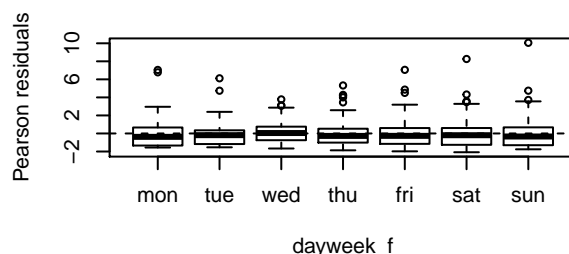
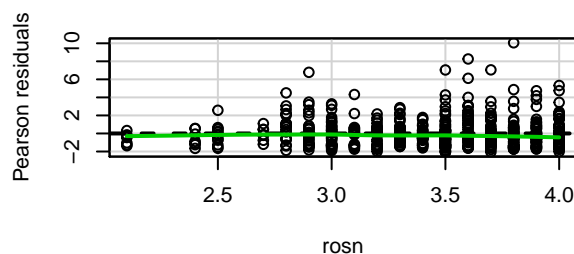
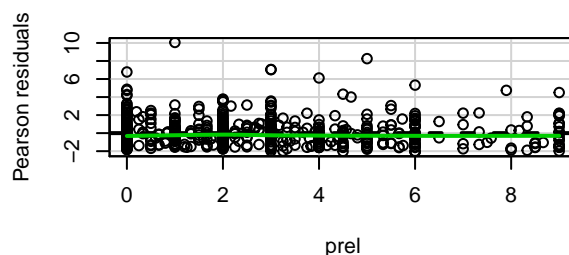
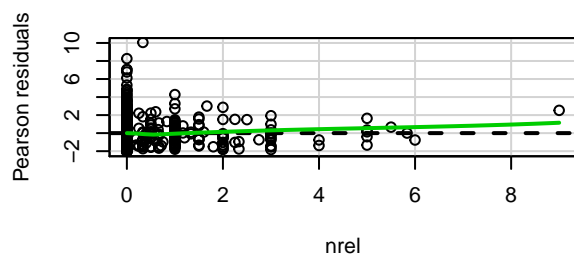
Histogram of N.mod.2\$residuals



```
plot(N.mod.2a)
```



```
# Pearson residuals
residualPlots(N.mod.2a)
```

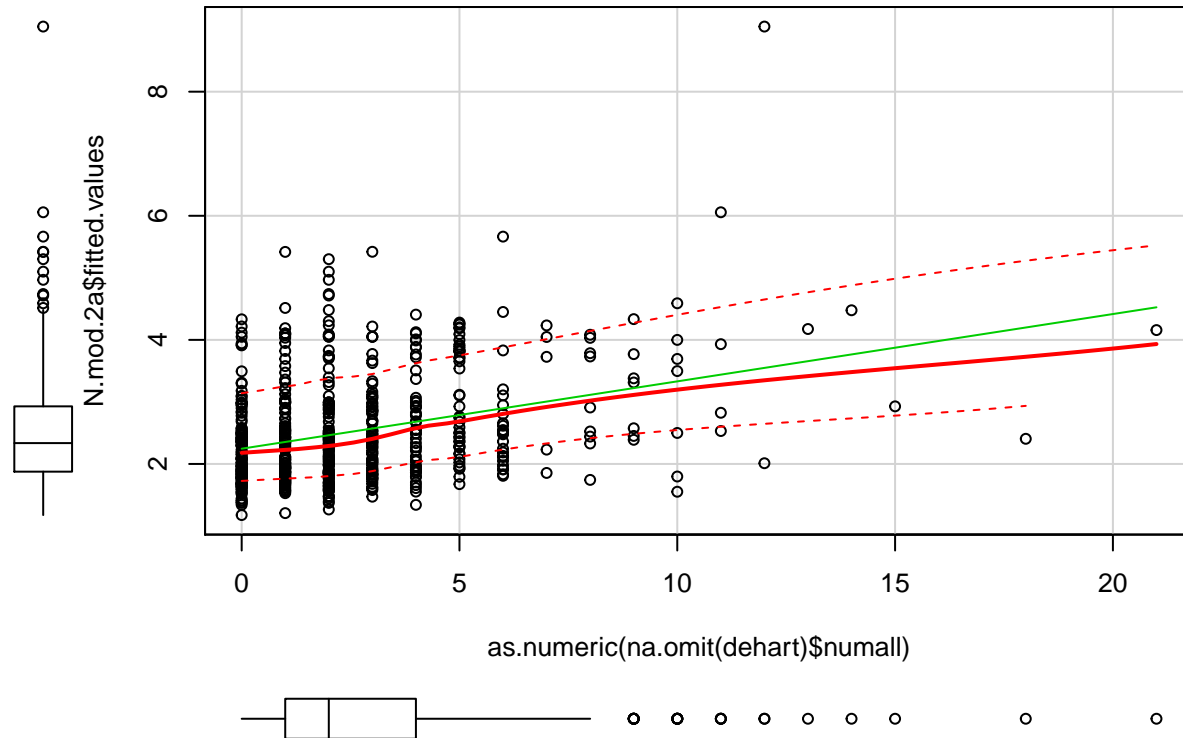


```
##          Test stat Pr(>|t|)
## nrel      6.696    0.010
```

```
## prel          0.000    0.996
## rosn          2.867    0.090
## dayweek_f      NA      NA
```

```
# scatterplot of actual and fitted values
```

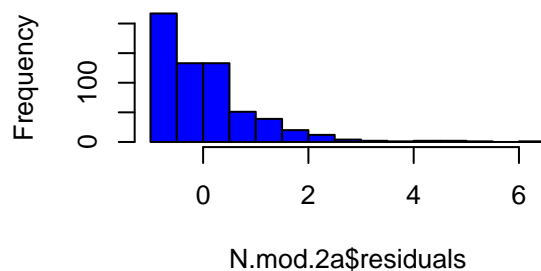
```
scatterplot(as.numeric(na.omit(dehart)$numall), N.mod.2a$fitted.values)
```



```
# histogram of residuals
```

```
hist(N.mod.2a$residuals, breaks = 20, col = "blue")
```

Histogram of N.mod.2a\$residuals



```
# EVERYTHING MODEL
```

```
N.mod.3 <- glm(formula = numall ~ nrel + prel + negevent + posevent + rosn + nrel*rosl + prel*rosl + da
summary(N.mod.3)
```

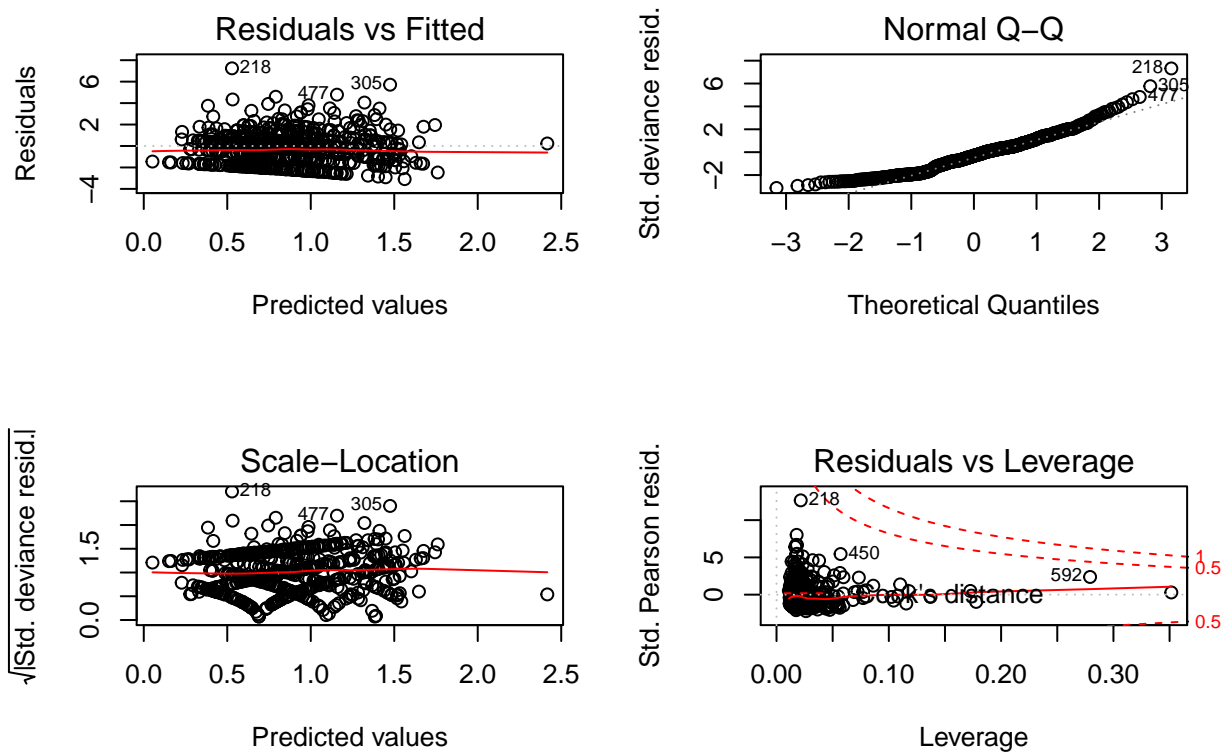
```
##
## Call:
## glm(formula = numall ~ nrel + prel + negevent + posevent + rosn +
##      nrel * rosn + prel * rosn + dayweek_f + gender_f, family = poisson(link = "log"),
##      data = na.omit(dehart))
```

```

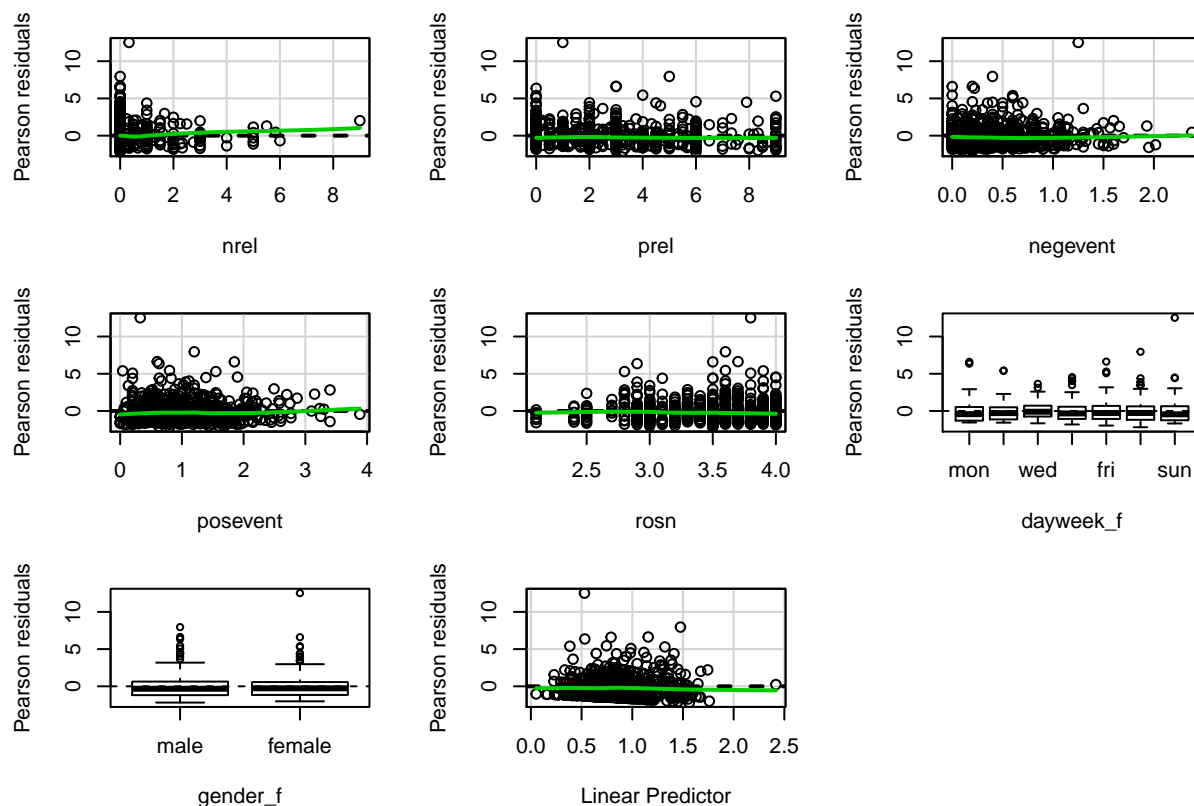
##
## Deviance Residuals:
##      Min       1Q   Median       3Q      Max
## -3.0888  -1.5326  -0.3009   0.5579   7.2406
##
## Coefficients:
##              Estimate Std. Error z value Pr(>|z|)
## (Intercept)  -0.18977    0.35862  -0.529 0.596697
## nrel          0.67296    0.24626   2.733 0.006282 **
## prel         0.23071    0.07679   3.004 0.002660 **
## negevent     -0.28695    0.07873  -3.645 0.000268 ***
## posevent      0.05717    0.04553   1.256 0.209197
## rosn          0.25820    0.10247   2.520 0.011742 *
## dayweek_ftue -0.14105    0.11041  -1.278 0.201401
## dayweek_fwed -0.06779    0.10818  -0.627 0.530885
## dayweek_fthu  0.20608    0.10109   2.039 0.041491 *
## dayweek_ffri  0.35610    0.09750   3.652 0.000260 ***
## dayweek_fsat  0.65559    0.09222   7.109 1.17e-12 ***
## dayweek_fsun  0.15534    0.10198   1.523 0.127684
## gender_ffemale -0.12184    0.05287  -2.304 0.021200 *
## nrel:rosl     -0.15359    0.07125  -2.156 0.031108 *
## prel:rosl     -0.05709    0.02252  -2.535 0.011243 *
## ---
## Signif. codes:  0 '***' 0.001 '**' 0.01 '*' 0.05 '.' 0.1 ' ' 1
##
## (Dispersion parameter for poisson family taken to be 1)
##
##      Null deviance: 1583.5  on 617  degrees of freedom
## Residual deviance: 1393.9  on 603  degrees of freedom
## AIC: 2785.9
##
## Number of Fisher Scoring iterations: 5
# hypothesis tests for statistical significance
Anova(N.mod.3, test = "LR")

## Analysis of Deviance Table (Type II tests)
##
## Response: numall
##              LR Chisq Df Pr(>Chisq)
## nrel          23.990  1  9.684e-07 ***
## prel           8.845  1  0.002938 **
## negevent      13.739  1  0.000210 ***
## posevent       1.558  1  0.211933
## rosn           0.152  1  0.696398
## dayweek_f    109.411  6 < 2.2e-16 ***
## gender_f       5.302  1  0.021296 *
## nrel:rosl      4.527  1  0.033371 *
## prel:rosl      6.355  1  0.011707 *
## ---
## Signif. codes:  0 '***' 0.001 '**' 0.01 '*' 0.05 '.' 0.1 ' ' 1
# diagnostic plots
par(mfrow=c(2,2))
plot(N.mod.3)

```



```
# Pearson residuals
residualPlots(N.mod.3)
```

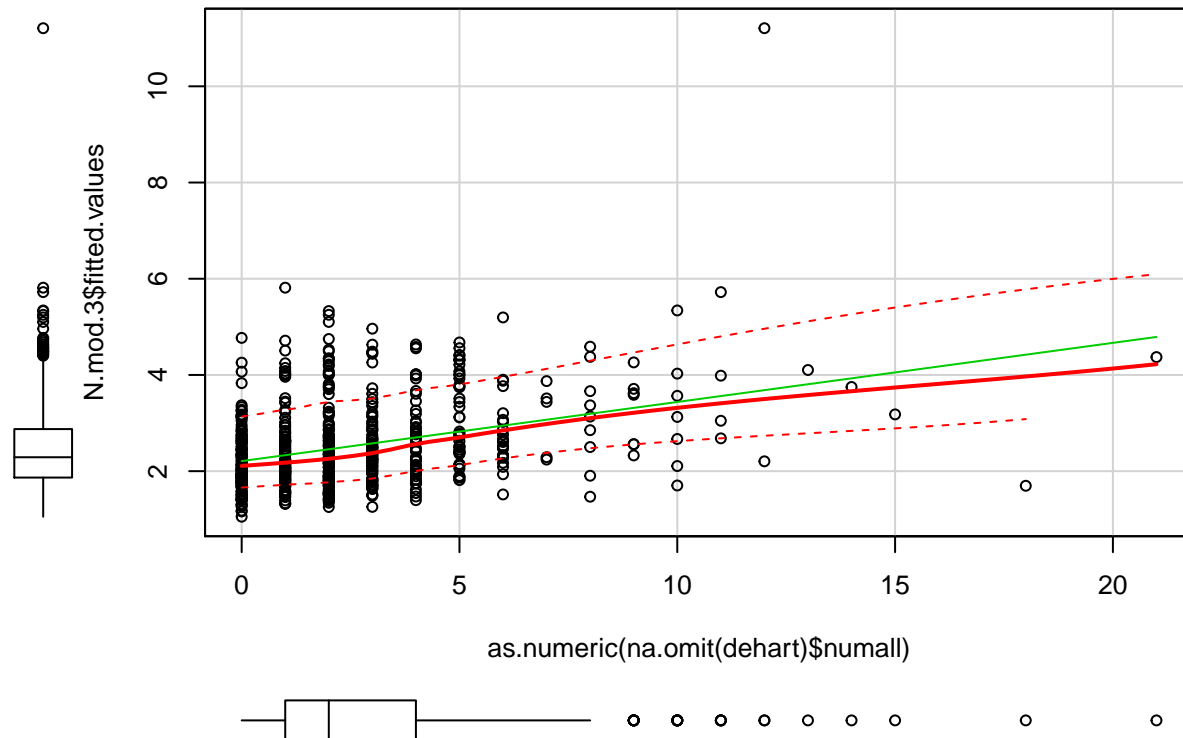


```
##          Test stat Pr(>|t|)
## nrel      4.860    0.027
```

```
## prel          0.086    0.769
## negevent      0.434    0.510
## posevent     1.369    0.242
## rosn         1.595    0.207
## dayweek_f      NA      NA
## gender_f       NA      NA
```

```
# scatterplot of actual and fitted values
```

```
scatterplot(as.numeric(na.omit(dehart)$numall), N.mod.3$fitted.values)
```



```
# histogram of residuals
```

```
hist(N.mod.3$residuals, breaks = 20, col = "blue")
```

```
# AGAIN WITH FACTOR FOR NREL
```

```
N.mod.3a <- glm(formula = numall ~ nrel_f + prel + negevent + posevent + rosn + nrel_f*rosn + prel*rosn
summary(N.mod.3a)
```

```
##
## Call:
## glm(formula = numall ~ nrel_f + prel + negevent + posevent +
##      rosn + nrel_f * rosn + prel * rosn + dayweek_f + gender_f,
##      family = poisson(link = "log"), data = na.omit(dehart))
##
## Deviance Residuals:
##      Min       1Q   Median       3Q      Max
## -3.0351  -1.4924  -0.3336   0.5733   6.8959
##
## Coefficients:
##              Estimate Std. Error z value Pr(>|z|)
## (Intercept)   -0.29535    0.36850  -0.801  0.42285
```



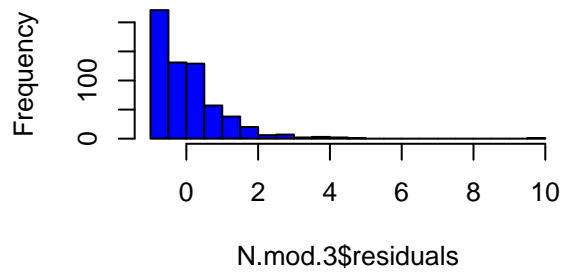
```

## nrel_fhigh      1.27887      0.54581      2.343      0.01913 *
## prel           0.23940      0.07676      3.119      0.00181 **
## negevent       -0.18588      0.07493     -2.481      0.01311 *
## posevent       0.06225      0.04583      1.358      0.17441
## rosn           0.28017      0.10473      2.675      0.00747 **
## dayweek_ftue   -0.13216      0.11070     -1.194      0.23254
## dayweek_fwed   -0.05320      0.10803     -0.492      0.62240
## dayweek_fthu    0.21604      0.10143      2.130      0.03317 *
## dayweek_ffri    0.38033      0.09755      3.899      9.66e-05 ***
## dayweek_fsat    0.67713      0.09217      7.347      2.03e-13 ***
## dayweek_fsun    0.17687      0.10213      1.732      0.08329 .
## gender_ffemale  -0.12023      0.05294     -2.271      0.02313 *
## nrel_fhigh:rosn -0.32065      0.15819     -2.027      0.04266 *
## prel:rosn      -0.06174      0.02250     -2.744      0.00606 **
## ---
## Signif. codes:  0 '***' 0.001 '**' 0.01 '*' 0.05 '.' 0.1 ' ' 1
##
## (Dispersion parameter for poisson family taken to be 1)
##
## Null deviance: 1583.5 on 617 degrees of freedom
## Residual deviance: 1411.2 on 603 degrees of freedom
## AIC: 2803.1
##
## Number of Fisher Scoring iterations: 5
# hypothesis tests for statistical significance
Anova(N.mod.3a, test = "LR")

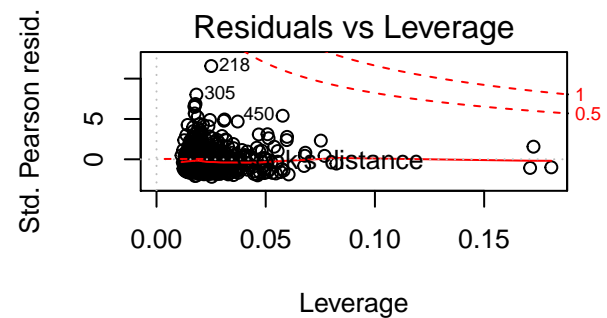
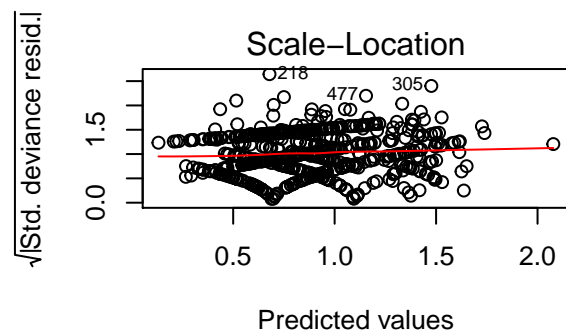
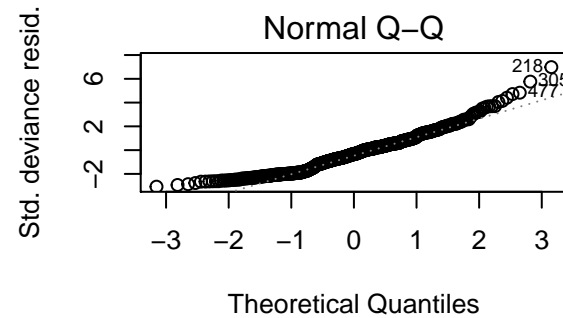
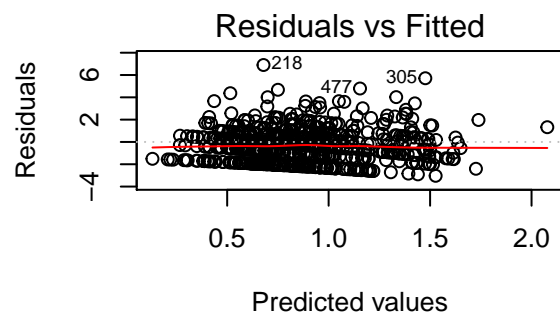
## Analysis of Deviance Table (Type II tests)
##
## Response: numall
##          LR Chisq Df Pr(>Chisq)
## nrel_f      7.187  1  0.007344 **
## prel       5.871  1  0.015388 *
## negevent    6.308  1  0.012018 *
## posevent    1.820  1  0.177352
## rosn        0.101  1  0.751226
## dayweek_f  114.269  6 < 2.2e-16 ***
## gender_f    5.151  1  0.023234 *
## nrel_f:rosn  4.044  1  0.044319 *
## prel:rosn   7.444  1  0.006364 **
## ---
## Signif. codes:  0 '***' 0.001 '**' 0.01 '*' 0.05 '.' 0.1 ' ' 1
# diagnostic plots
par(mfrow=c(2,2))

```

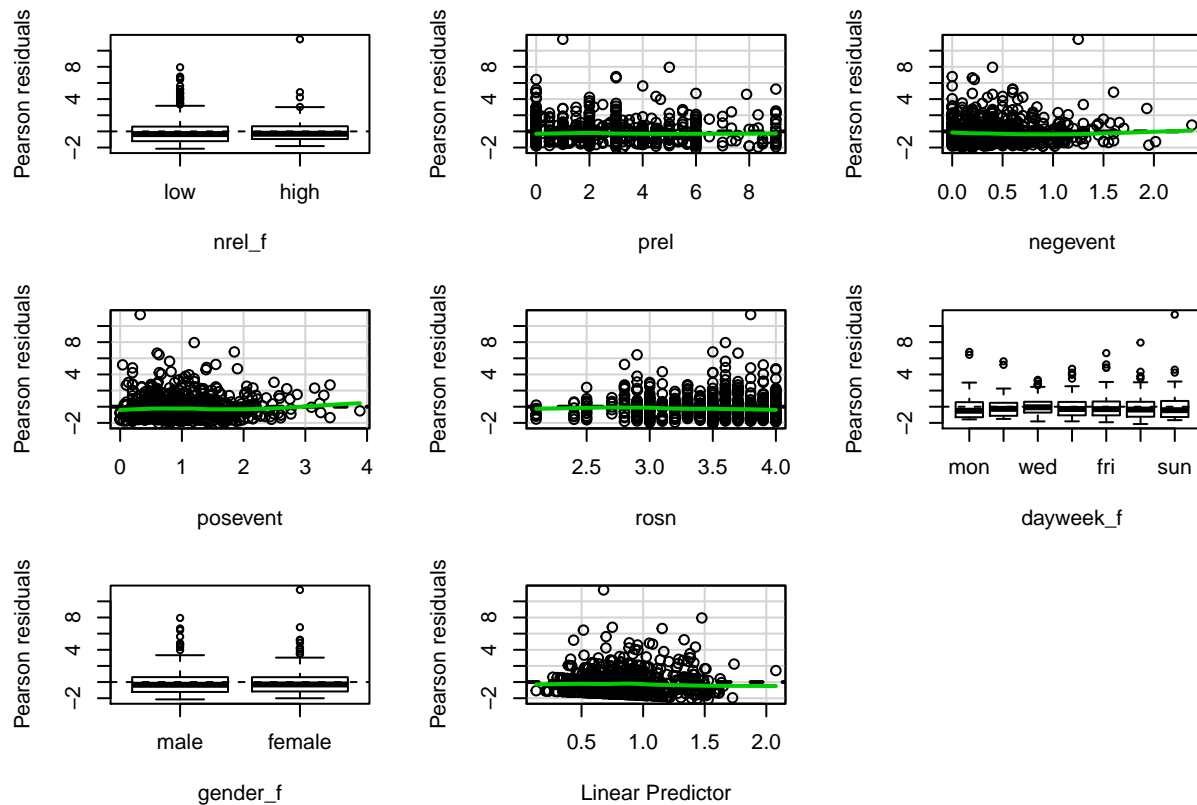
Histogram of N.mod.3\$residuals



```
plot(N.mod.3a)
```

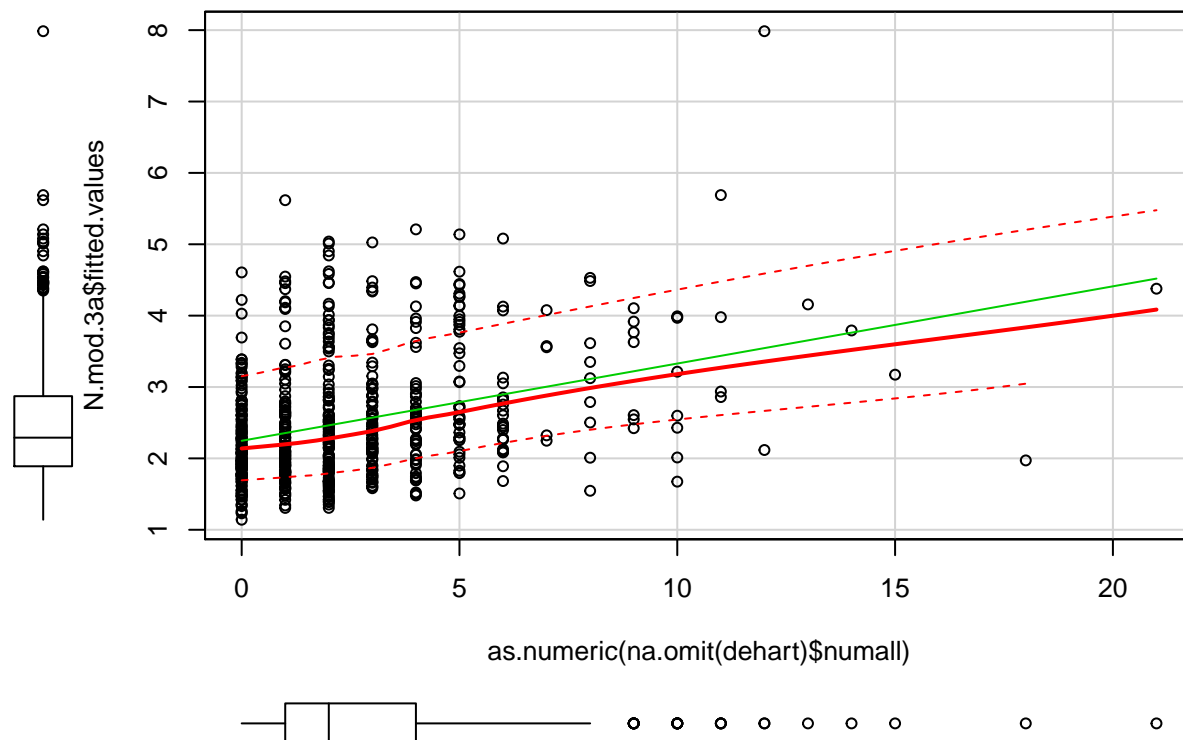


```
# Pearson residuals  
residualPlots(N.mod.3a)
```



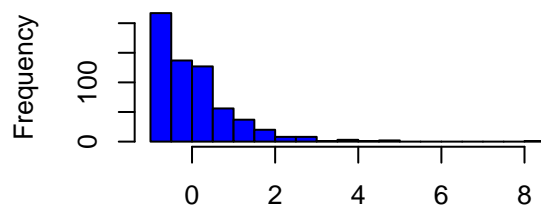
```
##          Test stat Pr(>|t|)
## nrel_f          NA      NA
## prel           0.668    0.414
## negevent        2.283    0.131
## posevent        3.083    0.079
## rosn            2.133    0.144
## dayweek_f       NA      NA
## gender_f       NA      NA
```

```
# scatterplot of actual and fitted values
scatterplot(as.numeric(na.omit(dehart)$numall), N.mod.3a$fitted.values)
```



```
# histogram of residuals
hist(N.mod.3a$residuals, breaks = 20, col = "blue")
```

Histogram of N.mod.3a\$residuals



N.mod.3a\$residuals

DESIRE TO DRINK

```
# DESIRE TO DRINK; ORDINAL LOGISTIC MODEL
# No significant interaction effect
```

```
# number of 'bins' :
```

```
N <- 3
```

```
dehart$desired_o <- cut(dehart$desired,N)
levels(dehart$desired_o) <- c("low","med","high")
```

```
library(package = MASS)
```

```
## Warning: package 'MASS' was built under R version 3.4.3
```

```
##
```

```
## Attaching package: 'MASS'
```

```
## The following object is masked from 'package:dplyr':
```

```
##
##      select
# BASIC MODEL

D.mod.0 <- polr(formula = desired_o ~ nrel + rosn, method = "logistic", data = na.omit(dehart))
summary(D.mod.0)

##
## Re-fitting to get Hessian

## Call:
## polr(formula = desired_o ~ nrel + rosn, data = na.omit(dehart),
##      method = "logistic")
##
## Coefficients:
##          Value Std. Error t value
## nrel  0.2327     0.0842   2.763
## rosn -0.6614     0.1820  -3.633
##
## Intercepts:
##          Value Std. Error t value
## low|med -3.2555   0.6391  -5.0940
## med|high -1.2734   0.6258  -2.0347
##
## Residual Deviance: 1301.456
## AIC: 1309.456

# hypothesis tests for statistical significance
Anova(D.mod.0, test = "LR")

## Analysis of Deviance Table (Type II tests)
##
## Response: desired_o
##      LR Chisq Df Pr(>Chisq)
## nrel   7.9511  1  0.0048058 **
## rosn  13.4152  1  0.0002496 ***
## ---
## Signif. codes:  0 '***' 0.001 '**' 0.01 '*' 0.05 '.' 0.1 ' ' 1

D.mod.1 <- polr(formula = desired_o ~ nrel + rosn + nrel*rosn, method = "logistic", data = na.omit(dehart))
summary(D.mod.1)

##
## Re-fitting to get Hessian

## Call:
## polr(formula = desired_o ~ nrel + rosn + nrel * rosn, data = na.omit(dehart),
##      method = "logistic")
##
## Coefficients:
##          Value Std. Error t value
## nrel      0.7295     0.8564   0.8518
## rosn     -0.6242     0.1925  -3.2418
## nrel:rosn -0.1429     0.2448  -0.5838
##
## Intercepts:
```

```

##           Value Std. Error t value
## low|med -3.1279  0.6741    -4.6401
## med|high -1.1457  0.6619    -1.7311
##
## Residual Deviance: 1301.11
## AIC: 1311.11

# hypothesis tests for statistical significance
Anova(D.mod.1, test = "LR")

## Analysis of Deviance Table (Type II tests)
##
## Response: desired_o
##           LR Chisq Df Pr(>Chisq)
## nrel           7.9511  1  0.0048058 **
## rosn          13.4152  1  0.0002496 ***
## nrel:rosn      0.3467  1  0.5559720
## ---
## Signif. codes:  0 '***' 0.001 '**' 0.01 '*' 0.05 '.' 0.1 ' ' 1

# factors for nrel
D.mod.0a <- polr(formula = desired_o ~ nrel_f + rosn, method = "logistic", data = na.omit(dehart))
summary(D.mod.0a)

##
## Re-fitting to get Hessian
## Call:
## polr(formula = desired_o ~ nrel_f + rosn, data = na.omit(dehart),
##       method = "logistic")
##
## Coefficients:
##           Value Std. Error t value
## nrel_fhigh  0.2042     0.1776   1.149
## rosn       -0.6536     0.1822  -3.588
##
## Intercepts:
##           Value Std. Error t value
## low|med -3.2532  0.6387    -5.0933
## med|high -1.2863  0.6256    -2.0562
##
## Residual Deviance: 1308.084
## AIC: 1316.084

# hypothesis tests for statistical significance
Anova(D.mod.0a, test = "LR")

## Analysis of Deviance Table (Type II tests)
##
## Response: desired_o
##           LR Chisq Df Pr(>Chisq)
## nrel_f      1.3231  1  0.2500292
## rosn       13.0747  1  0.0002993 ***
## ---
## Signif. codes:  0 '***' 0.001 '**' 0.01 '*' 0.05 '.' 0.1 ' ' 1

```

```
D.mod.1a <- polr(formula = desired_o ~ nrel_f + rosn + nrel_f*rosn, method = "logistic", data = na.omit(summary(D.mod.1a))
```

```
##
## Re-fitting to get Hessian

## Call:
## polr(formula = desired_o ~ nrel_f + rosn + nrel_f * rosn, data = na.omit(dehart),
##      method = "logistic")
##
## Coefficients:
##              Value Std. Error t value
## nrel_fhigh      1.9635      1.6524   1.188
## rosn            -0.5624      0.2006  -2.804
## nrel_fhigh:rosn -0.5057      0.4720  -1.071
##
## Intercepts:
##              Value Std. Error t value
## low|med    -2.9419    0.6995   -4.2058
## med|high   -0.9726    0.6889   -1.4118
##
## Residual Deviance: 1306.926
## AIC: 1316.926
```

```
# hypothesis tests for statistical significance
Anova(D.mod.1a, test = "LR")
```

```
## Analysis of Deviance Table (Type II tests)
##
## Response: desired_o
##              LR Chisq Df Pr(>Chisq)
## nrel_f          1.3231  1  0.2500292
## rosn           13.0747  1  0.0002993 ***
## nrel_f:rosn     1.1583  1  0.2818095
## ---
## Signif. codes:  0 '***' 0.001 '**' 0.01 '*' 0.05 '.' 0.1 ' ' 1
```

```
# INTERMEDIATE MODEL; DAYS OF THE WEEK
```

```
D.mod.2 <- polr(formula = desired_o ~ nrel + prel + rosn + nrel*rosn + prel*rosn + dayweek_f, method = summary(D.mod.2)
```

```
##
## Re-fitting to get Hessian

## Call:
## polr(formula = desired_o ~ nrel + prel + rosn + nrel * rosn +
##      prel * rosn + dayweek_f, data = na.omit(dehart), method = "logistic")
##
## Coefficients:
##              Value Std. Error t value
## nrel          1.04024      0.88741   1.1722
## prel          0.40201      0.25436   1.5805
## rosn         -0.41787      0.28431  -1.4698
## dayweek_ftue  0.17013      0.28610   0.5947
## dayweek_fwed  0.24405      0.28909   0.8442
```

```
## dayweek_fthu 0.43213 0.29240 1.4779
## dayweek_ffri 0.58517 0.28812 2.0310
## dayweek_fsat 0.72353 0.29169 2.4804
## dayweek_fsun -0.36303 0.29268 -1.2404
## nrel:rosl -0.21341 0.25329 -0.8425
## prel:rosl -0.08328 0.07412 -1.1237
##
```

```
## Intercepts:
```

```
## Value Std. Error t value
## low|med -1.8847 0.9971 -1.8901
## med|high 0.1882 0.9927 0.1896
##
```

```
## Residual Deviance: 1265.545
```

```
## AIC: 1291.545
```

```
# hypothesis tests for statistical significance
```

```
Anova(D.mod.2, test = "LR")
```

```
## Analysis of Deviance Table (Type II tests)
```

```
##
```

```
## Response: desired_o
```

```
## LR Chisq Df Pr(>Chisq)
## nrel 11.8245 1 0.0005846 ***
## prel 12.9135 1 0.0003262 ***
## rosl 14.5145 1 0.0001391 ***
## dayweek_f 19.7301 6 0.0030929 **
## nrel:rosl 0.7236 1 0.3949748
## prel:rosl 1.2785 1 0.2581742
```

```
## ---
```

```
## Signif. codes: 0 '***' 0.001 '**' 0.01 '*' 0.05 '.' 0.1 ' ' 1
```

```
# AGAIN WITH FACTOR FOR NREL
```

```
D.mod.2a <- polr(formula = desired_o ~ nrel + prel + rosl + nrel*rosl + prel*rosl + dayweek_f, method =
summary(D.mod.2a)
```

```
##
```

```
## Re-fitting to get Hessian
```

```
## Call:
```

```
## polr(formula = desired_o ~ nrel + prel + rosl + nrel * rosl +
## prel * rosl + dayweek_f, data = na.omit(dehart), method = "logistic")
```

```
##
```

```
## Coefficients:
```

```
## Value Std. Error t value
## nrel 1.04024 0.88741 1.1722
## prel 0.40201 0.25436 1.5805
## rosl -0.41787 0.28431 -1.4698
## dayweek_ftue 0.17013 0.28610 0.5947
## dayweek_fwed 0.24405 0.28909 0.8442
## dayweek_fthu 0.43213 0.29240 1.4779
## dayweek_ffri 0.58517 0.28812 2.0310
## dayweek_fsat 0.72353 0.29169 2.4804
## dayweek_fsun -0.36303 0.29268 -1.2404
## nrel:rosl -0.21341 0.25329 -0.8425
## prel:rosl -0.08328 0.07412 -1.1237
```

```
##
```



```

## Intercepts:
##      Value Std. Error t value
## low|med -1.8847  0.9971   -1.8901
## med|high  0.1882  0.9927    0.1896
##
## Residual Deviance: 1265.545
## AIC: 1291.545

# hypothesis tests for statistical significance
Anova(D.mod.2a, test = "LR")

## Analysis of Deviance Table (Type II tests)
##
## Response: desired_o
##      LR Chisq Df Pr(>Chisq)
## nrel      11.8245  1 0.0005846 ***
## prel      12.9135  1 0.0003262 ***
## rosn      14.5145  1 0.0001391 ***
## dayweek_f  19.7301  6 0.0030929 **
## nrel:rosn   0.7236  1 0.3949748
## prel:rosn   1.2785  1 0.2581742
## ---
## Signif. codes:  0 '***' 0.001 '**' 0.01 '*' 0.05 '.' 0.1 ' ' 1

# EVERYTHING MODEL

D.mod.3 <- polr(formula = desired_o ~ nrel + prel + negevent + posevent + rosn + nrel*rosn + prel*rosn +
summary(D.mod.3)

##
## Re-fitting to get Hessian

## Call:
## polr(formula = desired_o ~ nrel + prel + negevent + posevent +
##      rosn + nrel * rosn + prel * rosn + dayweek_f + gender_f,
##      data = na.omit(dehart), method = "logistic")
##
## Coefficients:
##      Value Std. Error t value
## nrel      1.06457   0.89038  1.1956
## prel      0.36682   0.25715  1.4265
## negevent   0.07835   0.23000  0.3406
## posevent   0.29010   0.14827  1.9565
## rosn     -0.38060   0.28738 -1.3244
## dayweek_ftue 0.17462   0.28700  0.6085
## dayweek_fwed 0.23381   0.29036  0.8052
## dayweek_fthu 0.44950   0.29310  1.5336
## dayweek_ffri 0.57864   0.28920  2.0008
## dayweek_fsat 0.73233   0.29259  2.5029
## dayweek_fsun -0.36900   0.29467 -1.2522
## gender_ffemale -0.37536   0.16101 -2.3313
## nrel:rosn   -0.21804   0.25446 -0.8569
## prel:rosn   -0.08630   0.07465 -1.1561
##
## Intercepts:
##      Value Std. Error t value

```

```
## low|med -1.7482 1.0062 -1.7375
## med|high 0.3456 1.0024 0.3448
##
## Residual Deviance: 1257.573
## AIC: 1289.573
```

```
# hypothesis tests for statistical significance
Anova(D.mod.3, test = "LR")
```

```
## Analysis of Deviance Table (Type II tests)
##
## Response: desired_o
##          LR Chisq Df Pr(>Chisq)
## nrel          9.8935 1 0.0016586 **
## prel          3.3432 1 0.0674818 .
## negevent       0.1161 1 0.7332954
## posevent       3.8755 1 0.0489964 *
## rosn          12.8532 1 0.0003369 ***
## dayweek_f     19.9499 6 0.0028268 **
## gender_f       5.4631 1 0.0194223 *
## nrel:rosn       0.7440 1 0.3883749
## prel:rosn       1.3545 1 0.2444937
## ---
## Signif. codes:  0 '***' 0.001 '**' 0.01 '*' 0.05 '.' 0.1 ' ' 1
```

```
# AGAIN WITH FACTOR FOR NREL
```

```
D.mod.3a <- polr(formula = desired_o ~ nrel_f + prel + negevent + posevent + rosn + nrel_f*rosn + prel*
summary(D.mod.3a)
```

```
##
## Re-fitting to get Hessian
## Call:
## polr(formula = desired_o ~ nrel_f + prel + negevent + posevent +
##       rosn + nrel_f * rosn + prel * rosn + dayweek_f + gender_f,
##       data = na.omit(dehart), method = "logistic")
##
## Coefficients:
##              Value Std. Error t value
## nrel_fhigh      2.37635      1.7146  1.3860
## prel           0.37847      0.2555  1.4811
## negevent        0.31361      0.2197  1.4276
## posevent        0.27271      0.1475  1.8492
## rosn           -0.31480      0.2917 -1.0791
## dayweek_ftue    0.19285      0.2878  0.6702
## dayweek_fwed    0.26519      0.2904  0.9132
## dayweek_fthu    0.45382      0.2936  1.5457
## dayweek_ffri    0.61483      0.2894  2.1248
## dayweek_fsat    0.75681      0.2923  2.5889
## dayweek_fsun   -0.33017      0.2954 -1.1178
## gender_ffemale  -0.36240      0.1610 -2.2515
## nrel_fhigh:rosn -0.61124      0.4895 -1.2486
## prel:rosn       -0.09397      0.0742 -1.2665
##
## Intercepts:
##              Value Std. Error t value
```

```

## low|med -1.4866 1.0214 -1.4555
## med|high 0.5897 1.0189 0.5788
##
## Residual Deviance: 1265.037
## AIC: 1297.037
# hypothesis tests for statistical significance
Anova(D.mod.3a, test = "LR")

## Analysis of Deviance Table (Type II tests)
##
## Response: desired_o
##          LR Chisq Df Pr(>Chisq)
## nrel_f      1.6008  1 0.2057878
## prel        2.1860  1 0.1392746
## negevent     2.0408  1 0.1531332
## posevent     3.4588  1 0.0629154 .
## rosn        13.1261  1 0.0002912 ***
## dayweek_f    19.9268  6 0.0028537 **
## gender_f      5.0941  1 0.0240073 *
## nrel_f:rosn   1.5731  1 0.2097587
## prel:rosn     1.6256  1 0.2023108
## ---
## Signif. codes:  0 '***' 0.001 '**' 0.01 '*' 0.05 '.' 0.1 ' ' 1

```