

Lab 9 Solutions

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We'll begin by loading some packages.

```
library(MASS)
library(plyr)
library(ggplot2)
library(reshape)
```

```
##
## Attaching package: 'reshape'
## The following objects are masked from 'package:plyr':
##
##      rename, round_any
```

```
library(e1071)
```

Let's form our favourite birthwt data set.

```
# Rename the columns to have more descriptive names
colnames(birthwt) <- c("birthwt.below.2500", "mother.age", "mother.weight",
  "race", "mother.smokes", "previous.prem.labor", "hypertension", "uterine.irr",
  "physician.visits", "birthwt.grams")

# Transform variables to factors with descriptive levels
birthwt <- transform(birthwt,
  race = as.factor(mapvalues(race, c(1, 2, 3),
    c("white", "black", "other"))),
  mother.smokes = as.factor(mapvalues(mother.smokes,
    c(0,1), c("no", "yes"))),
  hypertension = as.factor(mapvalues(hypertension,
    c(0,1), c("no", "yes"))),
  uterine.irr = as.factor(mapvalues(uterine.irr,
    c(0,1), c("no", "yes")))
)
```

ANOVA with birthwt data

(a) Create a new factor that categorizes the number of physician visits into three levels: 0, 1, 2, 3 or more.

```
pvb <- unique(birthwt$physician.visits)
repvb <- as.character(pvb)
for (i in 4:length(repvb)) {
  repvb[i] = "3 or more"
}
birthwt <- transform(birthwt, phys.visit.binned1 = as.factor(mapvalues(physician.visits, pvb, repvb )))
birthwt$phys.visit.binned1
```

```
##      [1] 0          3          1          3 or more 0          0          1
##      [8] 1          1          0          0          1          0          3 or more
##     [15] 0          0          0          3          0          1          3 or more
##     [22] 3          1          0          3 or more 0          0          3 or more
```

```
## [29] 0      1      1      1      1      1      0
## [36] 3 or more 3 or more 0      3 or more 1      3 or more 3 or more
## [43] 1      0      0      0      3 or more 0      3 or more
## [50] 0      1      0      0      3 or more 0      0
## [57] 0      0      0      0      0      3 or more 0
## [64] 0      0      1      3 or more 3 or more 1      3 or more
## [71] 0      3 or more 1      0      0      0      1
## [78] 3 or more 0      0      1      0      0      0
## [85] 0      0      0      0      0      1      0
## [92] 3 or more 0      0      0      1      1      0
## [99] 0      1      1      0      0      1      0
## [106] 0      1      0      3 or more 3 or more 3 or more 1
## [113] 3 or more 1      0      1      0      0      3 or more
## [120] 1      1      0      1      0      3 or more 3 or more
## [127] 1      0      1      1      0      3 or more 0
## [134] 0      0      0      1      1      0      1
## [141] 0      0      0      1      0      3 or more 3 or more
## [148] 0      0      0      1      3 or more 0      0
## [155] 0      0      3      1      0      0      0
## [162] 1      0      0      0      0      3 or more 0
## [169] 1      0      1      0      0      0      0
## [176] 0      1      3      0      3 or more 1      3
## [183] 0      0      3 or more 3 or more 0      0      3
## Levels: 0 1 3 3 or more
```

```
phys.visit.binned <- birthwt$physician.visits
phys.visit.binned[phys.visit.binned >= 3] <- "3.or.more"
birthwt <- transform(birthwt, phys.visit.binned = as.factor(phys.visit.binned))
birthwt$phys.visit.binned
```

```
## [1] 0      3.or.more 1      2      0      0      1
## [8] 1      1      0      0      1      0      2
## [15] 0      0      0      3.or.more 0      1      2
## [22] 3.or.more 1      0      2      0      0      2
## [29] 0      1      1      1      1      1      0
## [36] 2      2      0      2      1      2      2
## [43] 1      0      0      0      3.or.more 0      2
## [50] 0      1      0      0      2      0      0
## [57] 0      0      0      0      0      2      0
## [64] 0      0      1      2      3.or.more 1      2
## [71] 0      2      1      0      0      0      1
## [78] 3.or.more 0      0      1      0      0      0
## [85] 0      0      0      0      0      1      0
## [92] 2      0      0      0      1      1      0
## [99] 0      1      1      0      0      1      0
## [106] 0      1      0      2      3.or.more 2      1
## [113] 2      1      0      1      0      0      2
## [120] 1      1      0      1      0      2      2
## [127] 1      0      1      1      0      2      0
## [134] 0      0      0      1      1      0      1
## [141] 0      0      0      1      0      2      2
## [148] 0      0      0      1      2      0      0
## [155] 0      0      3.or.more 1      0      0      0
## [162] 1      0      0      0      0      3.or.more 0
## [169] 1      0      1      0      0      0      0
```

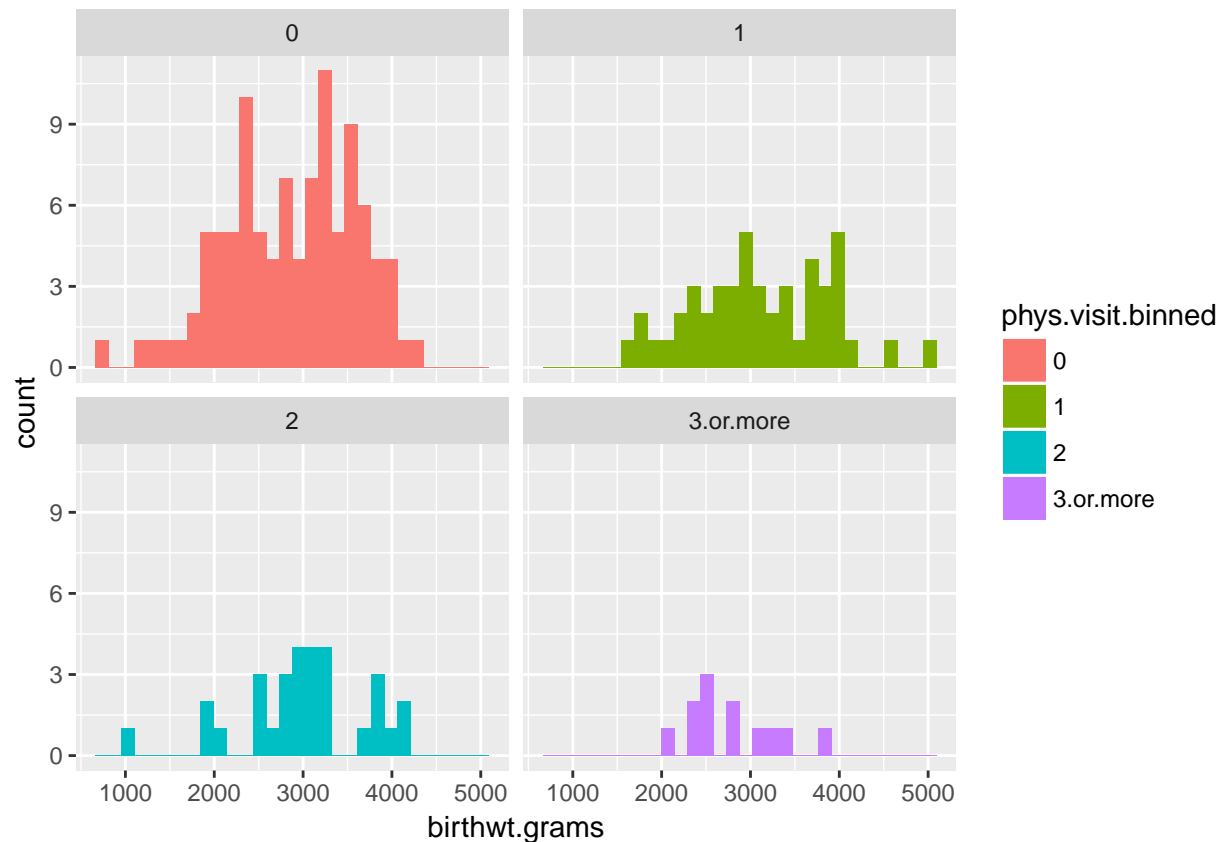
```
## [176] 0          1          3.or.more 0          2          1          3.or.more
## [183] 0          0          2          2          0          0          3.or.more
## Levels: 0 1 2 3.or.more
```

Hint: One way of doing this is with mapvalues, by mapping all instances of 3, 4,... etc, to “3 or more”.

Histogram of Birthweight

```
qplot(data=birthwt, x = birthwt.grams, facets = ~phys.visit.binned, geom = "histogram", fill = phys.visit.binned)
```

```
## `stat_bin()` using `bins = 30`. Pick better value with `binwidth`.
```



Check for Skewness

```
with(data=birthwt, skewness(birthwt.grams))
```

```
## [1] -0.205337
```

```
with(data=birthwt, skewness(birthwt.grams[phys.visit.binned=="0"]))
```

```
## [1] -0.3924435
```

```
with(data=birthwt, skewness(birthwt.grams[phys.visit.binned=="1"]))
```

```
## [1] 0.07103918
```

```
with(data=birthwt, skewness(birthwt.grams[phys.visit.binned=="2"]))
```

```
## [1] -0.5105783
```

```
with(data=birthwt, skewness(birthwt.grams[phys.visit.binned=="3.or.more"]))
```

```
## [1] 0.5874174
```

(b) Run an ANOVA to determine whether the average birth weight varies across number of physician visits.

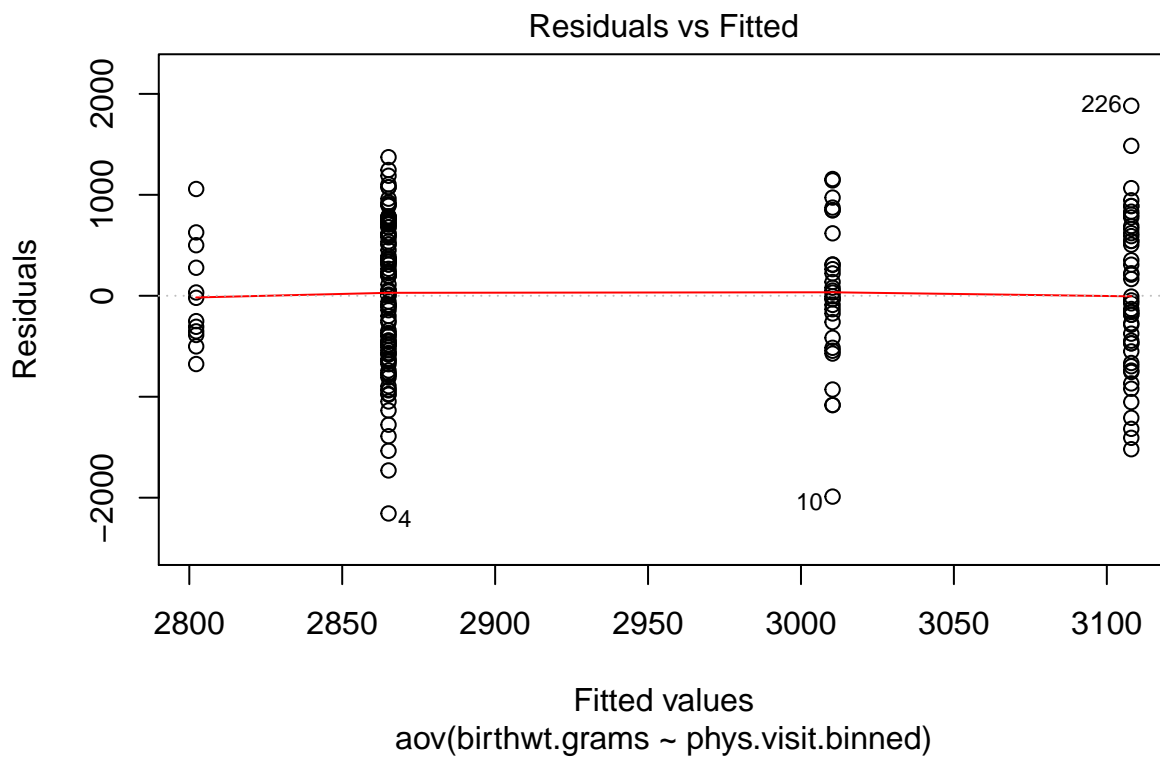
```
aov.birthwt <- aov(birthwt.grams ~ phys.visit.binned, data = birthwt)
```

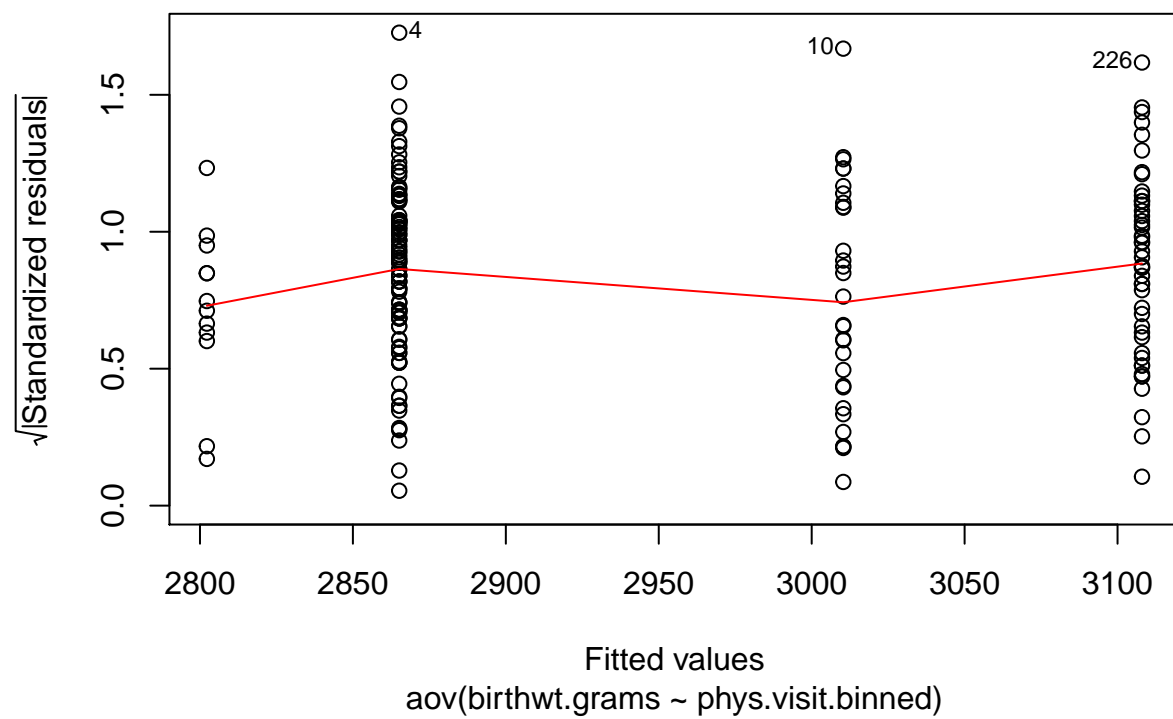
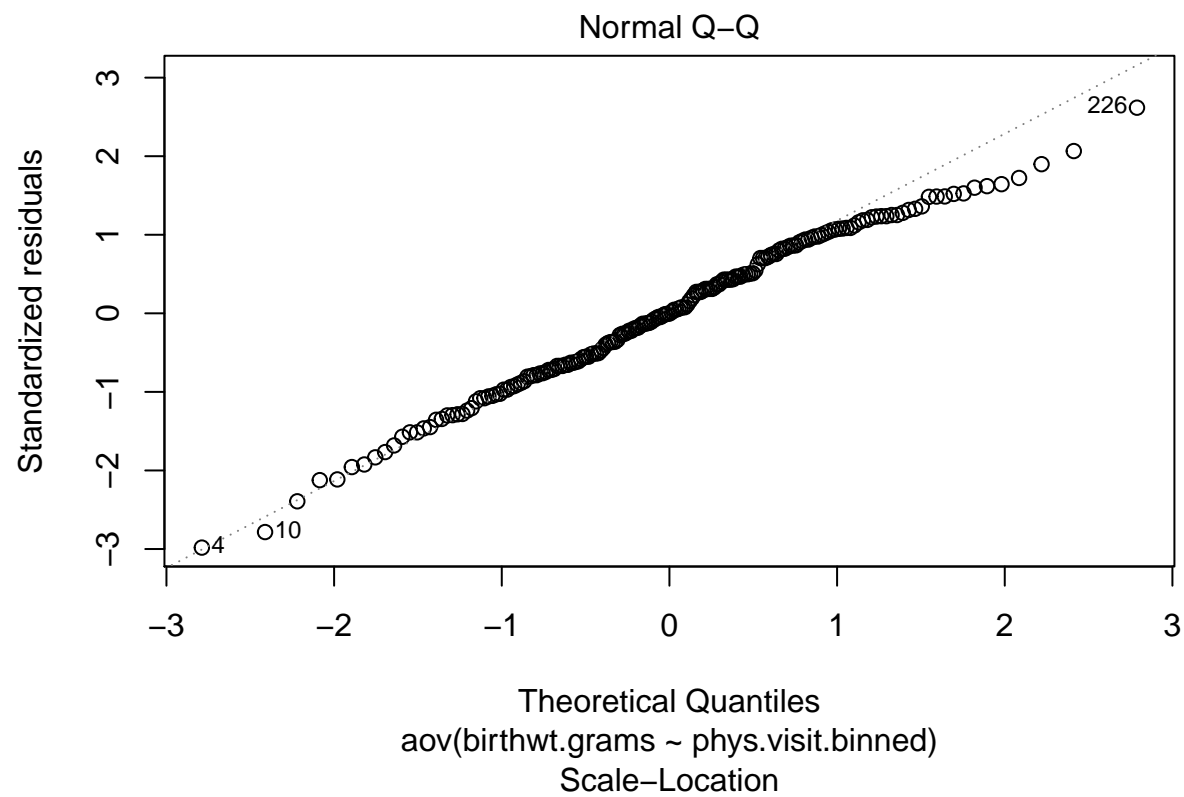
```
summary(aov.birthwt)
```

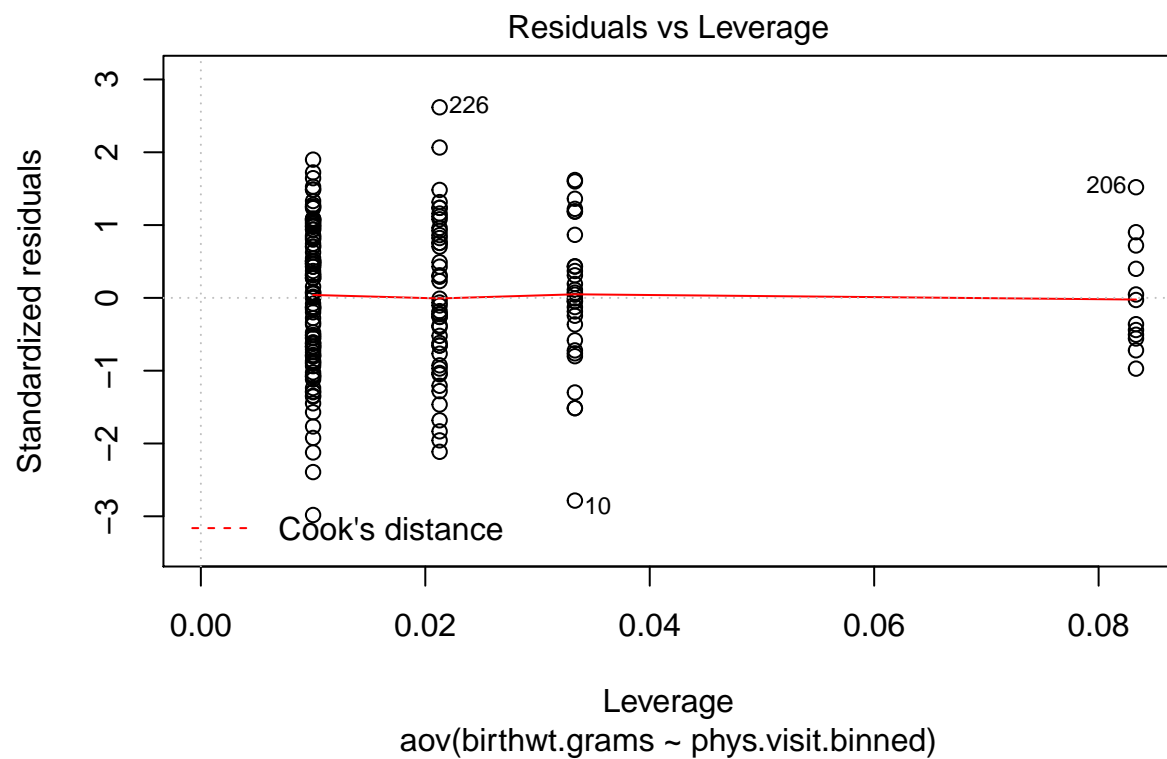
```
##              Df    Sum Sq Mean Sq F value Pr(>F)
## phys.visit.binned  3  2259057   753019    1.426   0.237
## Residuals        185  97710599   528165
```

The p-value is greater than 0.05, so the variation in birthweight across number of physician visits is not statistically significant.

```
plot(aov.birthwt)
```







Three-way ANOVA

```
twaov.birthwt <- aov(birthwt.grams ~ race+mother.smokes+phys.visit.binned, data = birthwt)
summary(twaov.birthwt)
```

```
##              Df    Sum Sq Mean Sq F value    Pr(>F)
## race           2  5015725  2507863    5.227 0.006205 **
## mother.smokes  1  7322575  7322575   15.262 0.000132 ***
## phys.visit.binned 3   311098   103699    0.216 0.885120
## Residuals     182 87320257   479782
## ---
## Signif. codes:  0 '***' 0.001 '**' 0.01 '*' 0.05 '.' 0.1 ' ' 1
```