

# Linear regression of the Cats Dataset from MASS

*Jon Kinsey*

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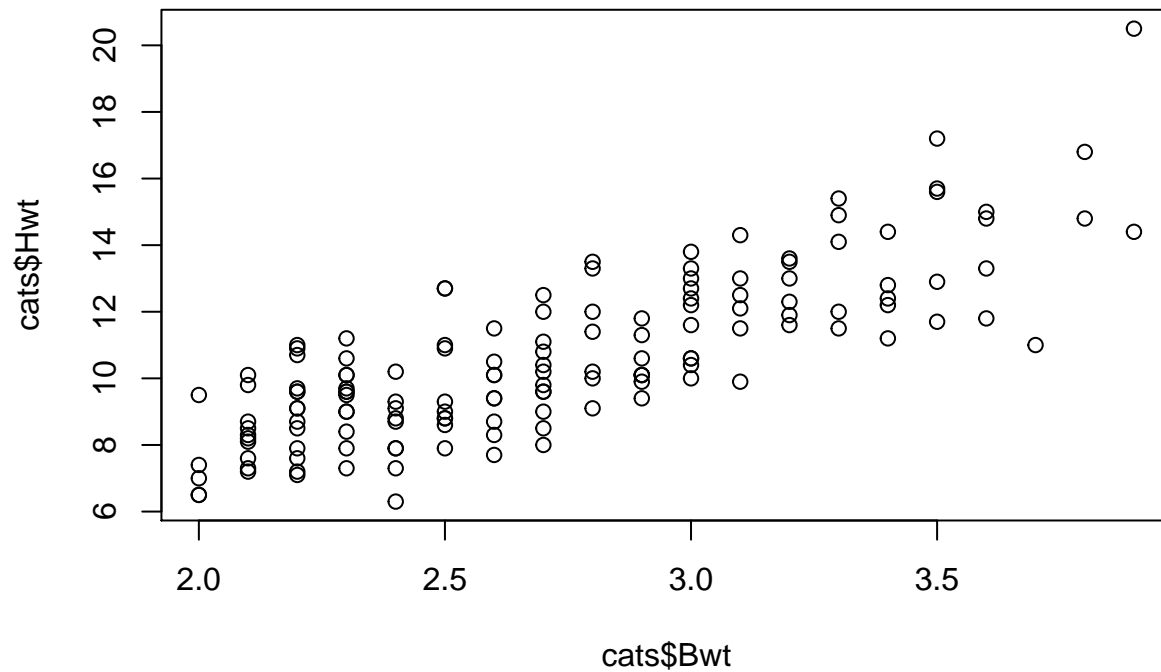
```
# The heart and body weights of samples of male and female cats used for
# digitalis experiments. The cats were all adult, over 2 kg body weight.
# This data frame contains the following columns:
# Sex : Factor with levels "F" and "M".
# Bwt : body weight in kg.
# Hwt : heart weight in g.
# Reference:
# R. A. Fisher (1947) The analysis of covariance method for the relation between
# a part and the whole, Biometrics 3, 65-68.
#
library(MASS)
data(cats)
str(cats)
```

```
## 'data.frame': 144 obs. of 3 variables:
## $ Sex: Factor w/ 2 levels "F","M": 1 1 1 1 1 1 1 1 1 1 ...
## $ Bwt: num 2 2 2 2.1 2.1 2.1 2.1 2.1 2.1 2.1 ...
## $ Hwt: num 7 7.4 9.5 7.2 7.3 7.6 8.1 8.2 8.3 8.5 ...
```

```
summary(cats)
```

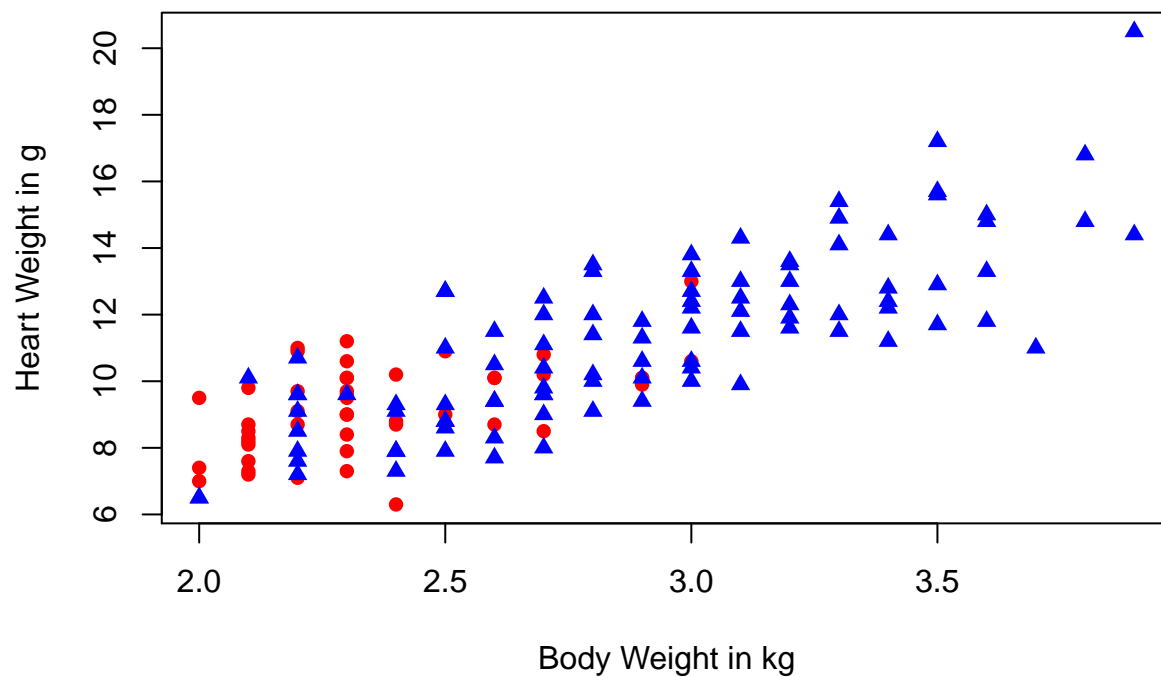
```
## Sex      Bwt      Hwt
## F:47  Min.   :2.000  Min.   : 6.30
## M:97  1st Qu.:2.300  1st Qu.: 8.95
##      Median :2.700  Median :10.10
##      Mean   :2.724  Mean   :10.63
##      3rd Qu.:3.025  3rd Qu.:12.12
##      Max.   :3.900  Max.   :20.50
```

```
#
plot(cats$Bwt, cats$Hwt)
```



```
# a more detailed plot separating males, females
with(cats, plot(Bwt, Hwt, type="n", xlab="Body Weight in kg",
               ylab="Heart Weight in g",
               main="Heart Weight vs. Body Weight of Cats"))
with(cats, points(Bwt[Sex=="F"], Hwt[Sex=="F"], pch=16, col="red"))
with(cats, points(Bwt[Sex=="M"], Hwt[Sex=="M"], pch=17, col="blue"))
```

### Heart Weight vs. Body Weight of Cats



```
# A Pearson product-moment correlation coefficient can be calculated using
# the cor( ) function
with(cats, cor(Bwt, Hwt))
```

```
## [1] 0.8041274
```

```
# Pearson's r = .804 indicates a strong positive relationship.
#
# regression fit:
attach(cats)
lm(Hwt ~ Bwt)
```

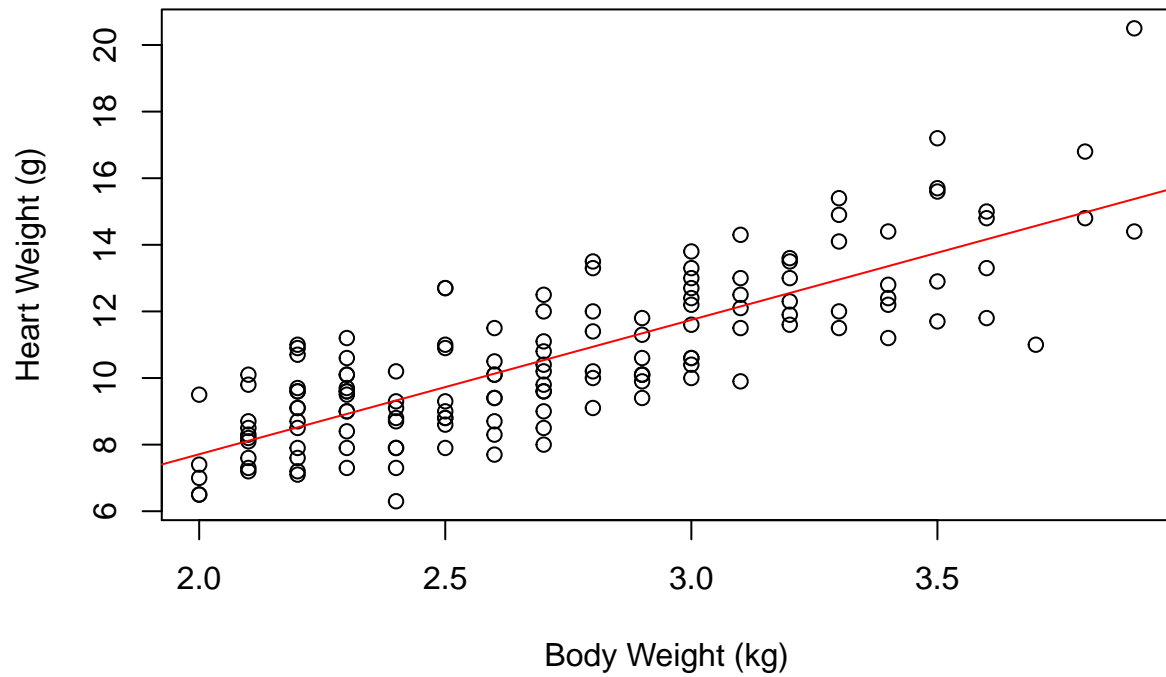
```
##
## Call:
## lm(formula = Hwt ~ Bwt)
##
## Coefficients:
## (Intercept)      Bwt
##      -0.3567      4.0341
```

```
# So the fitted regression equation is: Hwt = 4.0341 (Bwt) - 0.3567.
lmfit <- lm(Hwt ~ Bwt)
summary(lmfit)
```

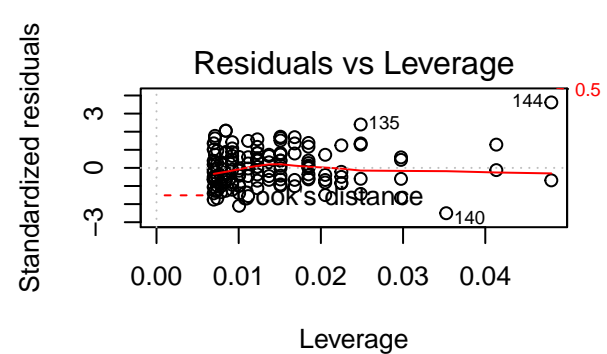
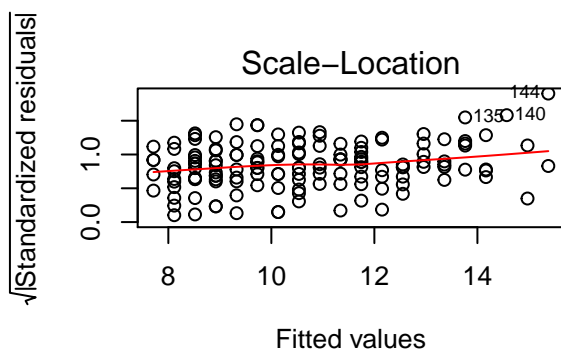
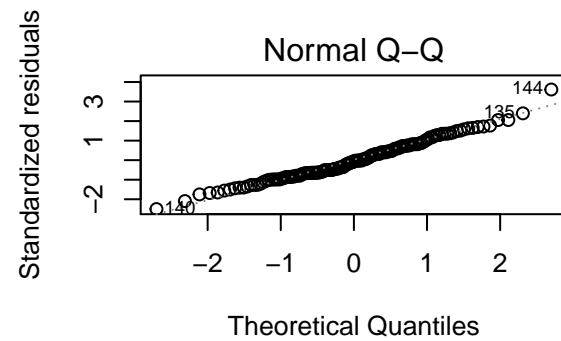
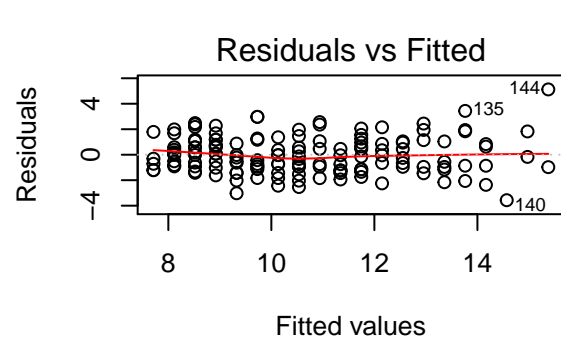
```
##
## Call:
## lm(formula = Hwt ~ Bwt)
##
## Residuals:
##      Min       1Q   Median       3Q      Max
## -3.5694 -0.9634 -0.0921  1.0426  5.1238
##
## Coefficients:
##              Estimate Std. Error t value Pr(>|t|)
## (Intercept)  -0.3567     0.6923  -0.515   0.607
## Bwt           4.0341     0.2503  16.119 <2e-16 ***
## ---
## Signif. codes:  0 '***' 0.001 '**' 0.01 '*' 0.05 '.' 0.1 ' ' 1
##
## Residual standard error: 1.452 on 142 degrees of freedom
## Multiple R-squared:  0.6466, Adjusted R-squared:  0.6441
## F-statistic: 259.8 on 1 and 142 DF, p-value: < 2.2e-16
```

```
# The three stars for Bwt indicate that the significance of the Bwt
# coefficient is between 0 and 0.001.
#
# plotting the regression line on a scatterplot
plot(Hwt ~ Bwt,
     xlab = "Body Weight (kg)", ylab = "Heart Weight (g)",
     main="Scatterplot of Body Weight vs. Heart Weight")
abline(lmfit, col="red")
```

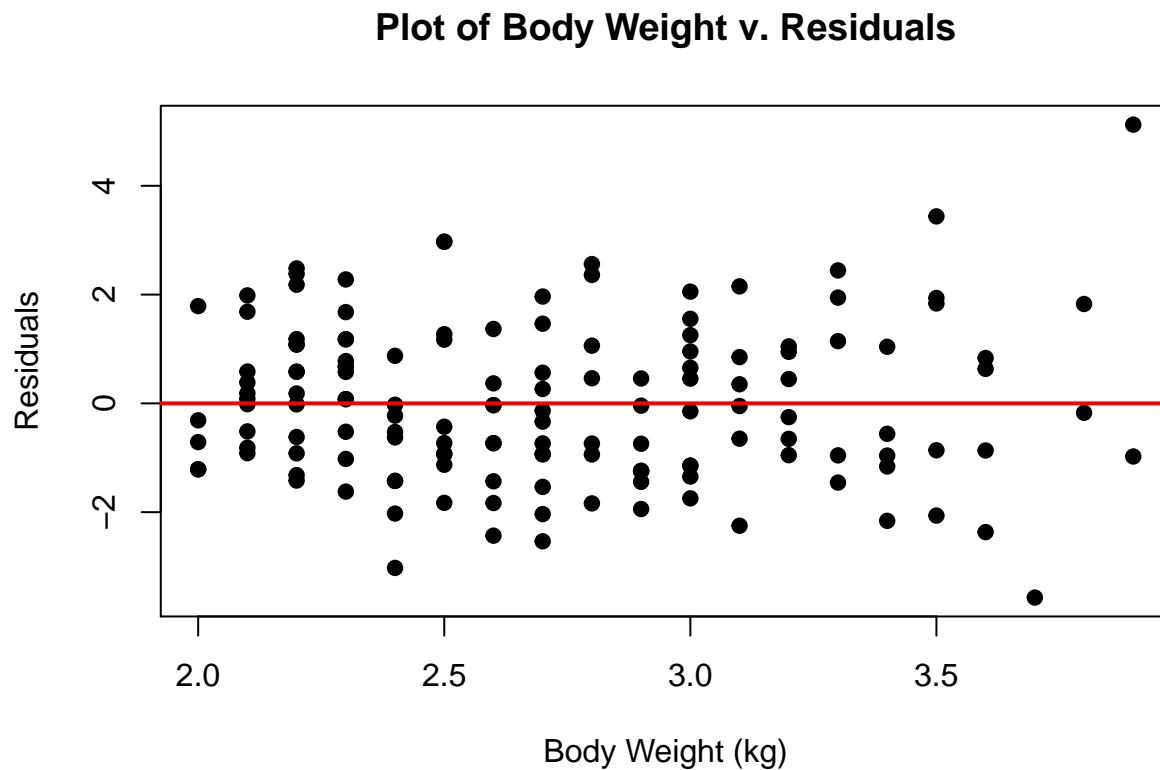
## Scatterplot of Body Weight vs. Heart Weight



```
#
par(mfrow=c(2,2)) # configure output window for 4 plots, 2x2
plot(lmfit)
```

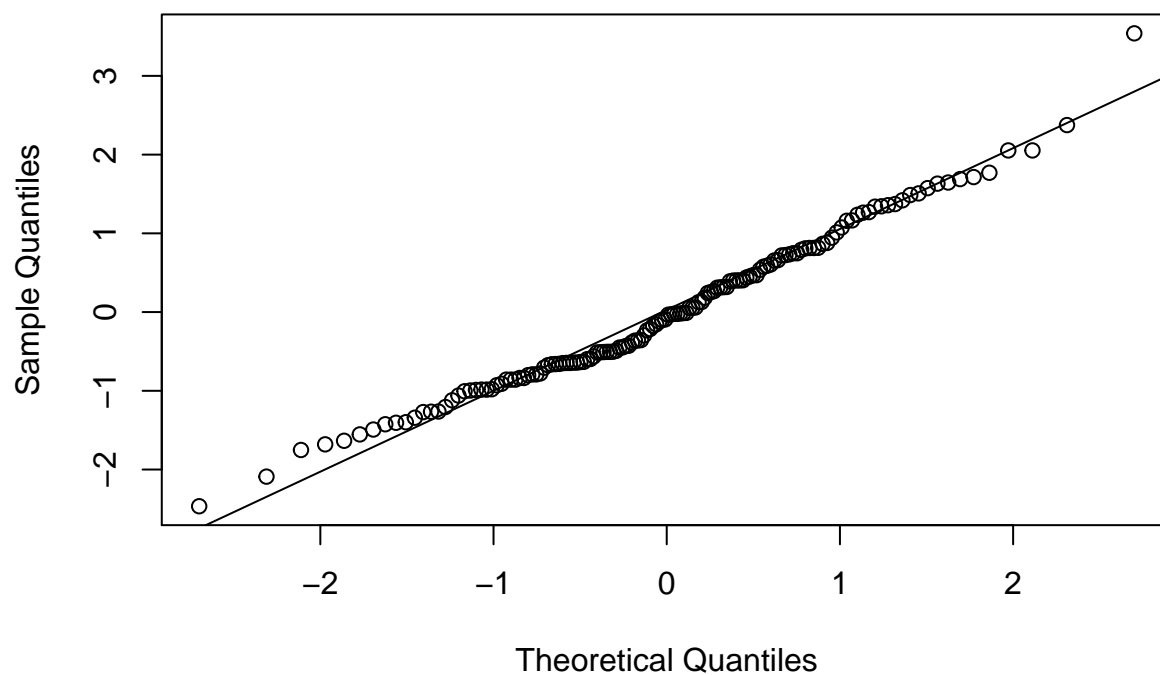


```
#
# plot of the residuals
par(mfrow=c(1,1))
res = lmfit$residuals
plot(Bwt, res, pch = 19, xlab = "Body Weight (kg)", ylab = "Residuals",
     main = "Plot of Body Weight v. Residuals")
abline(h = 0, col = "red", lwd = 2)
```



```
#
res = scale(res)
# replot the Q-Q plot
# Note: An ideal Q-Q plot has points falling more or less on the diagonal line,
# indicating that our residuals are approximately normally distributed.
qqnorm(res)
qqline(res)
```

## Normal Q-Q Plot



```
#  
# find the expected heart weight for a cat that weighs 3 kg  
newObs <- data.frame(Bwt=3)  
predict(lmfit, newObs, interval="predict")
```

```
##      fit      lwr      upr  
## 1 11.74553 8.861263 14.62979
```

```
# going back to the heart-body plot we see that 11.75 gms looks right
```