Regression

Code ▼

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```
library(car)
library(QuantPsyc)
library(boot)
library(dplyr) # data mainpulation
library(cowplot)
library(ggplot2)
```

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```
albumsales1<-lm(formula=sales~adverts, data=df1)

# Interpreting a simple regression
summary(albumsales1)</pre>
```

```
Call:
lm(formula = sales ~ adverts, data = df1)
Residuals:
    Min
                   Median
                               30
              10
                                       Max
-152.949 -43.796
                   -0.393
                          37.040 211.866
Coefficients:
            Estimate Std. Error t value Pr(>|t|)
(Intercept) 1.341e+02 7.537e+00 17.799 <2e-16 ***
adverts
           9.612e-02 9.632e-03
                                 9.979
                                         <2e-16 ***
Signif. codes: 0 '***' 0.001 '**' 0.05 '.' 0.1 ' ' 1
Residual standard error: 65.99 on 198 degrees of freedom
                             Adjusted R-squared: 0.3313
Multiple R-squared: 0.3346,
F-statistic: 99.59 on 1 and 198 DF, p-value: < 2.2e-16
```

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df2<- read.delim('/home/atrides/Desktop/R/statistics_with_R/07_Regression/Da
ta Files/Album Sales 2.dat', header=TRUE)</pre>

head(df2)

	adverts <dbl></dbl>	sales <int></int>	airplay <int></int>	attract <int></int>
1	10.256	330	43	10
2	985.685	120	28	7
3	1445.563	360	35	7
4	1188.193	270	33	7
5	574.513	220	44	5
6	568.954	170	19	5
6 rows				

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```
albumsales2<-lm(sales~adverts, data=df2)
albumsales3<-lm(sales~adverts+airplay+attract, data=df2) # or use update(alb
umsales2, .~.+attract+airplay)</pre>
```

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summary(albumsales2)

```
Call:
lm(formula = sales ~ adverts, data = df2)
Residuals:
    Min
              1Q Median
                               30
                                       Max
-152.949 -43.796
                   -0.393 37.040 211.866
Coefficients:
            Estimate Std. Error t value Pr(>|t|)
(Intercept) 1.341e+02 7.537e+00 17.799 <2e-16 ***
                                9.979 <2e-16 ***
adverts
           9.612e-02 9.632e-03
Signif. codes: 0 '***' 0.001 '**' 0.01 '*' 0.05 '.' 0.1 ' ' 1
Residual standard error: 65.99 on 198 degrees of freedom
Multiple R-squared: 0.3346, Adjusted R-squared: 0.3313
F-statistic: 99.59 on 1 and 198 DF, p-value: < 2.2e-16
```

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```
summary(albumsales3)
Call:
lm(formula = sales ~ adverts + airplay + attract, data = df2)
Residuals:
    Min
              10
                  Median
                               30
                                      Max
-121.324 -28.336
                  -0.451
                           28.967 144.132
Coefficients:
             Estimate Std. Error t value Pr(>|t|)
(Intercept) -26.612958 17.350001 -1.534
            0.084885
                       0.006923 12.261 < 2e-16 ***
adverts
airplay
             3.367425
                       0.277771 12.123 < 2e-16 ***
attract
            - - -
Signif. codes: 0 '***' 0.001 '**' 0.05 '.' 0.1 ' ' 1
Residual standard error: 47.09 on 196 degrees of freedom
Multiple R-squared: 0.6647,
                             Adjusted R-squared: 0.6595
F-statistic: 129.5 on 3 and 196 DF, p-value: < 2.2e-16
                                                                     Hide
# model parameters
print(lm.beta(albumsales3)) # standarized b balues
  adverts
           airplay
                    attract
0.5108462 0.5119881 0.1916834
                                                                     Hide
print(confint(albumsales3))
                  2.5 %
                            97.5 %
(Intercept) -60.82960967 7.60369295
            0.07123166 0.09853799
adverts
             2.81962186 3.91522848
airplay
attract
             6.27855218 15.89411823
                                                                     Hide
# comparing Models
anova(albumsales2, albumsales3)
```

```
Analysis of Variance Table
Model 1: sales ~ adverts
Model 2: sales ~ adverts + airplay + attract
  Res.Df RSS Df Sum of Sq F Pr(>F)
1
     198 862264
2
     196 434575 2 427690 96.447 < 2.2e-16 ***
Signif. codes: 0 '***' 0.001 '**' 0.05 '.' 0.1 ' ' 1
                                                                            Hide
cat("AIC model2: ", AIC(albumsales2),"\nBIC model2: ", BIC(albumsales2))
AIC model2: 2247.375
BIC model2: 2257.27
                                                                            Hide
cat("AIC_model3: ", AIC(albumsales3),"\nBIC_model3: ", BIC(albumsales3))
AIC model3: 2114.337
BIC model3: 2130.828
                                                                            Hide
# outliers and influential cases
df2$residuals <-resid(albumsales3)</pre>
df2$standarized.residuals <- rstandard(albumsales3)</pre>
df2$studentized.residuals <- rstudent(albumsales3)</pre>
df2$cooks <- cooks.distance(albumsales3)</pre>
df2$dfbeta <- dfbeta(albumsales3)</pre>
df2$dffit <- dffits(albumsales3)</pre>
df2$leverage <- hatvalues(albumsales3)</pre>
df2$covratio <- covratio(albumsales3)</pre>
# saving this data
write.table(df2,'albumSalesWithDiagnosticsData.dat', sep='\t', row.names = F
ALSE)
large_resid <- dplyr::filter(df2, standarized.residuals>2 | standarized.resi
duals< -2)
# these cases are to be analyzed coz they have somewhat large residuals
large resid
```

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```
# now lets see cooks distance , leverage , covariance ratio for 'these' case
s
k = 3 #number of predictors
n = 200 #number of objervations
average_leverage = (k+1)/n
average_leverage
```

```
[1] 0.02
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```
cvr_low<- 1-3*average_leverage
cvr_high<- 1+3*average_leverage

large_resid$cov_ideal_low <- cvr_low
large_resid$cov_ideal_high <- cvr_high

large_resid</pre>
```

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```
# from this large residual model we conclude that
```

Most of our 12 potential outliers have CVR values within or just outside t he boundaries.

none of them has a Cook's distance greater than 1, so none of the cases i s having an undue influence on the model.

So , Note:

- # i) Look at standardized residuals and check that no more than 5% of cases have absolute values above 2,
- # and that no more than about 1% have absolute values above 2.5. Any case with a value above about 3 could be an outlier.
- # ii)Look at the values of Cook's distance: any value above 1 indicates a ca se that might be influencing the model.
- # iii)Calculate the average leverage (the number of predictors plus 1, divid ed by the sample size)
- # and then look for values greater than twice or three times this avera ge value
- # iv)Calculate the upper and lower limit of acceptable values for the covariance ratio, CVR.
- # The upper limit is 1 plus three times the average leverage, whereas
- # the lower limit is 1 minus three times the average leverage.
- # Cases that have a CVR falling outside these limits may be problemat ic

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```
-----
# Testing various assumptions
# i) Assumptions of Independent Errors
car::durbinWatsonTest(albumsales3) # hence assumption is valid here
lag Autocorrelation D-W Statistic p-value
          0.0026951
                       1.949819
                                   0.738
Alternative hypothesis: rho != 0
                                                                      Hide
# ii) Assumption of no multicollinearity
vif_<- car::vif(albumsales3)</pre>
print(vif_)
adverts airplay attract
1.014593 1.042504 1.038455
                                                                      Hide
print(mean(vif ))
[1] 1.03185
                                                                      Hide
# the assumption of multicollinearity is followed too
                                                                      Hide
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```
# iii) Assumption about the Residuals
df2$fitted.values<- fitted.values(albumsales3)
df2$std_fitted.values<- (fitted.values(albumsales3)-mean(fitted.values(album
sales3)))/sd(fitted.values(albumsales3))
resid_plot<- ggplot(df2, aes(standarized.residuals,std_fitted.values))
resid_plot<- resid_plot+geom_point()+geom_smooth(formula='y~x',method = "l
m",alpha=0.1)

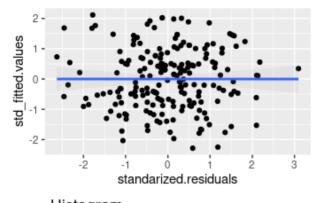
resid_qq<- ggplot(df2, aes(sample=studentized.residuals))
resid_qq<- resid_qq+stat_qq()+ggtitle('QQ-Plot')
resid_qq</pre>
```

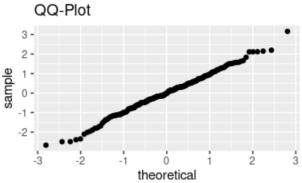


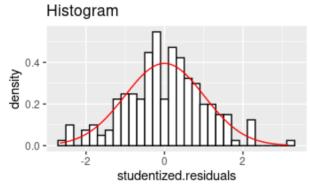
```
histresid<- ggplot(df2, aes(studentized.residuals))
histresid<- histresid+geom_histogram(aes(y=..density..),colour='black', fill
='white')+
    ggtitle('Histogram')+
    stat_function(fun = dnorm, args = list(mean=0, sd=sd(df2$studentized.residuals, na.rm = TRUE)), colour='red')

plot_grid(resid_plot, resid_qq, histresid,ncol=2, nrow=2)</pre>
```

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







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this assumption was also met