Plot Comparison

JC 9/9/2017

Note

Plot Comparison based on plan.

```
Data & Libs
```

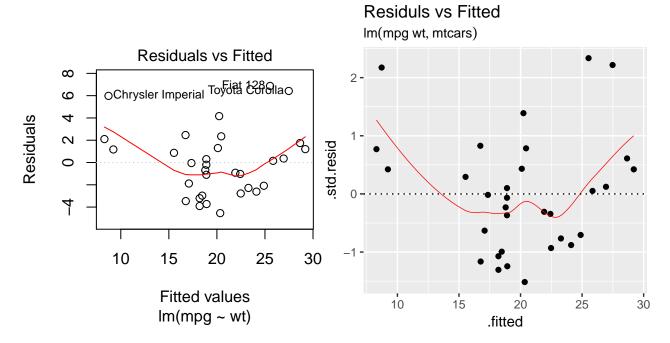
```
library(tidyverse)
library(broom)
library(gridExtra)
# Traditional Model
lm.1 <- lm(mpg ~ wt,data=mtcars)</pre>
# Generic name
My.Mod <- lm.1
# Tidy output
Tidy.Mod <- augment(My.Mod)
# Attributes
class(My.Mod)
## [1] "lm"
attributes (My. Mod)
## $names
## [1] "coefficients" "residuals"
                                         "effects"
                                                         "rank"
   [5] "fitted.values" "assign"
                                         "qr"
                                                         "df.residual"
## [9] "xlevels"
                        "call"
                                         "terms"
                                                         "model"
##
## $class
## [1] "lm"
head(Tidy.Mod,4)
##
          .rownames mpg
                            wt .fitted
                                           .se.fit
## 1
          Mazda RX4 21.0 2.620 23.28261 0.6335798 -2.2826106 0.04326896
## 2 Mazda RX4 Wag 21.0 2.875 21.91977 0.5714319 -0.9197704 0.03519677
         Datsun 710 22.8 2.320 24.88595 0.7359177 -2.0859521 0.05837573
## 4 Hornet 4 Drive 21.4 3.215 20.10265 0.5384424 1.2973499 0.03125017
       .sigma
                  .cooksd .std.resid
## 1 3.067494 0.013274072 -0.7661677
## 2 3.093068 0.001723963 -0.3074305
## 3 3.072127 0.015439367 -0.7057525
## 4 3.088268 0.003020558 0.4327511
```

Notes axis scale/ aspect slightly different Text annotations logic \sim highlighting "Wigglyness" of line different \sim slightly diff parameter?

Plot1: Residules v Fitted

```
plot(My.Mod,which=1)

D1 <- augment(My.Mod) %>%
    ggplot(aes(x=.fitted,y=.std.resid)) +
    geom_point() +
    geom_smooth(se=FALSE,colour="red",size=.25) +
    geom_hline(yintercept=0,linetype=3) +
    labs(title="Residuls vs Fitted",subtitle=My.Mod$call)
D1
```

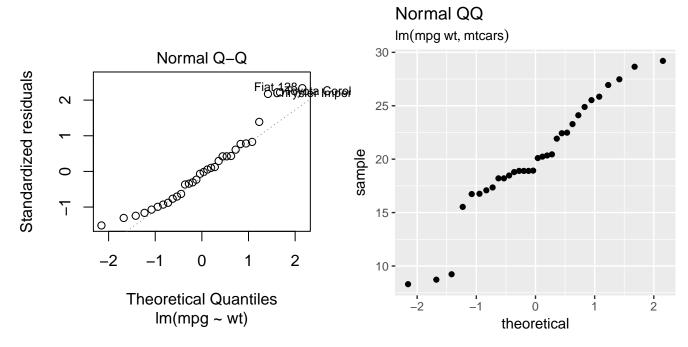


Plot2: Normal QQ

```
plot(My.Mod,which=2)

D2 <- augment(My.Mod) %>%
    ggplot(aes(sample=.fitted)) +
    stat_qq() +
    labs(title="Normal QQ",subtitle=My.Mod$call)

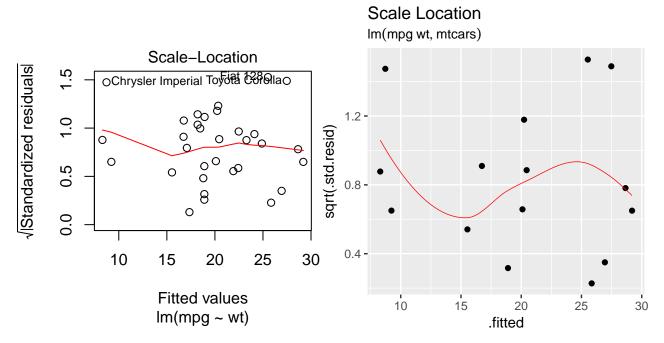
D2
```



Plot3: Scale Location

```
plot(My.Mod,which=3)

D3 <- augment(My.Mod) %>%
    ggplot(aes(x=.fitted, y=sqrt(.std.resid))) +
    geom_point()+
    labs(title="Scale Location",subtitle=My.Mod$call) +
    geom_smooth(se=FALSE,colour="red",size=.25)
D3
```



 ${\bf Questions.} \dots$

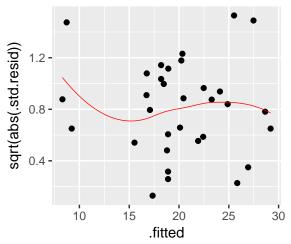
Missing Values? Why? Line wigliness.What parameters?

Missing Values

Raw Tidy.Mod\$.std.resid [1] -0.76616765 -0.30743051 -0.70575249 0.43275114 -0.06681879 [6] -0.23148309 -1.30552216 1.38889709 0.78392687 0.10010803 ## [11] -0.36728706 0.29288651 -0.01683789 -0.63159969 0.42296071 ## [16] 0.76979873 2.17353314 2.33490215 0.61035691 2.21708271 ## [21] -0.87964013 -0.99313634 -1.24418015 -1.16279098 **##** [26] 0.12244407 0.05177187 0.42254270 -1.51549710 -0.93086929 ## [31] -1.07151943 -0.34388215 # Why so many NAN's? sqrt(Tidy.Mod\$.std.resid) [1] NaNNaN NaN 0.6578382 NaN NaN NaN [8] 1.1785148 0.8853964 0.3163985 NaN ## NaN 0.5411899 NaN ## [15] 0.6503543 0.8773817 1.4742907 1.5280387 0.7812534 1.4889871 NaN ## [22] NaN 0.9097910 0.3499201 0.2275343 0.6500328 NaN NaN ## [29] NaN NaN NaN NaN #Try Abs value sqrt(abs(Tidy.Mod\$.std.resid)) [1] 0.8753100 0.5544642 0.8400908 0.6578382 0.2584933 0.4811269 1.1425945 [8] 1.1785148 0.8853964 0.3163985 0.6060421 0.5411899 0.1297609 0.7947325 ## [15] 0.6503543 0.8773817 1.4742907 1.5280387 0.7812534 1.4889871 0.9378913 **##** [22] 0.9965623 1.1154282 1.0783279 0.9097910 0.3499201 0.2275343 0.6500328 ## [29] 1.2310553 0.9648157 1.0351422 0.5864147 D3abs <- augment(My.Mod) %>% ggplot(aes(x=.fitted, y=sqrt(abs(.std.resid)))) + geom point()+ labs(title="Test SL with abs value", subtitle=My.Mod\$call) + geom_smooth(se=FALSE,colour="red",size=.25) D3abs

Test SL with abs value

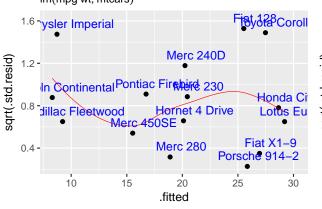
Im(mpg wt, mtcars)



with text labels added

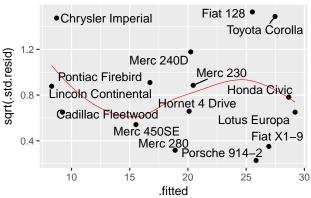
Scale Location

Im(mpg wt, mtcars)



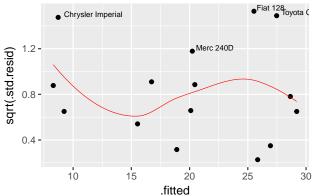
Scale Location

Im(mpg wt, mtcars)



Scale Location

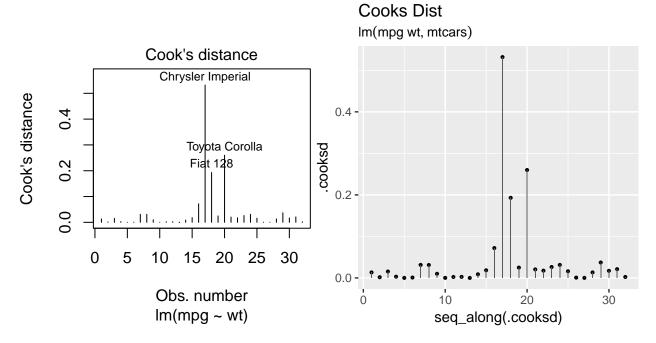
Im(mpg wt, mtcars)



Plot4: Cooks Distance

```
#Cooks Distane
plot(My.Mod,which=4)

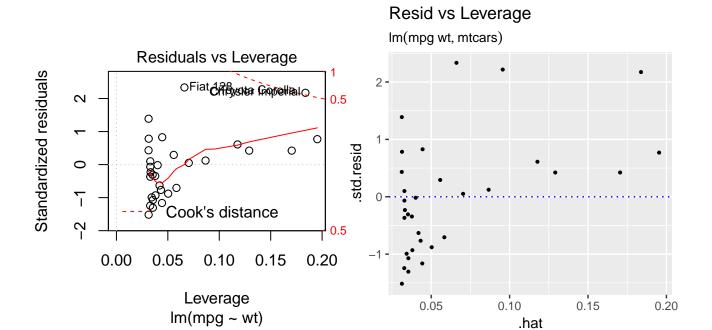
D4 <- augment(My.Mod) %>%
    ggplot(aes(x=seq_along(.cooksd), y=.cooksd)) +
    geom_point(size = .75) +
    geom_col(width = .1) +
    labs(title="Cooks Dist",subtitle=My.Mod$call)
D4
```



Plot5: Resid vs Leverage

```
# Resid vs Leverage
plot(My.Mod,which=5)

D5 <- augment(My.Mod) %>%
    ggplot(aes(x=.hat, y=.std.resid)) +
    geom_point(size=.75) +
    geom_hline(yintercept=0,linetype=3,colour="blue") +
    labs(title="Resid vs Leverage",subtitle=My.Mod$call)
D5
```



0.05

0.15

.hat

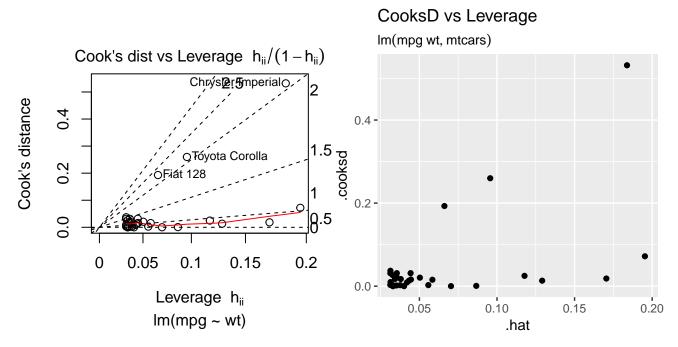
0.20

Plot6: CooksSD vs Leverage

```
plot(My.Mod,which=6)

D6 <- augment(My.Mod) %>%
    ggplot(aes(x=.hat, y=.cooksd)) +
    geom_point() +
    labs(title="CooksD vs Leverage",subtitle=My.Mod$call)

D6
```



Panel

```
#library(gridExtra)
D.Panel <- grid.arrange(D1, D2, D3, D4, D5, D6, ncol=2)
D.Panel
      Residuls vs Fitted
                                                         Normal QQ
      Im(mpg wt, mtcars)
                                                         Im(mpg wt, mtcars)
                                                     30 -
25 -
20 -
15 -
10 -
                                                  sample
                   15
                                              30
                                     25
          10
                            20
                        .fitted
                                                                        theoretical
                                                         Cooks Dist
       Scale Location
       Im(mpg wt, mtcars)
                                                         Im(mpg wt, mtcars)
sqrt(.std.resid)
                                                  .cooksd
                                                     0.2 -
   0.8 -
   0.4 -
                                                     0.0
                                                                     10
                    15
                                                                                 20
                                                                                             30
                             20
           10
                                              30
                                      25
                                                                   seq_along(.cooksd)
                         .fitted
                                                         CooksD vs Leverage
      Resid vs Leverage
                                                         Im(mpg wt, mtcars)
      Im(mpg wt, mtcars)
                                                     0.4 -
                                                     0.2 -
                                 0.15
                                             0.20
                      0.10
                                                                                    0.15
          0.05
                                                             0.05
                                                                         0.10
                                                                                                0.20
                         .hat
                                                                            .hat
## TableGrob (3 x 2) "arrange": 6 grobs
            cells
                      name
## 1 1 (1-1,1-1) arrange gtable[layout]
## 2 2 (1-1,2-2) arrange gtable[layout]
## 3 3 (2-2,1-1) arrange gtable[layout]
## 4 4 (2-2,2-2) arrange gtable[layout]
## 5 5 (3-3,1-1) arrange gtable[layout]
## 6 6 (3-3,2-2) arrange gtable[layout]
```

