

R08- Multiple Regression Examples

HCI/PSYCH 522
Iowa State University

April 7, 2022

Overview

- Children's heights depending on
 - mother's heights
 - + father's heights
 - + gender
- Wool breaks depending on
 - tension (L, M, H)
 - + type (A, B)

Multiple regression

Recall the simple linear regression model is

$$Y_i \stackrel{ind}{\sim} N(\mu_i, \sigma^2), \quad \mu_i = \beta_0 + \beta_1 X_{i,1} + \cdots + \beta_p X_{i,p}$$

where for observation i

- Y_i is the dependent variable and
- $X_{i,p}$ is the p^{th} independent variable.

Interpretation

- β_0 is mean of the dependent variable when all independent variables (X's) are 0
- β_p for $p \neq 0$, is the mean increase in the dependent variable for each unit increase in the associated independent variable
- R^2 is the proportion of variability in the dependent variable explained by the model

heights

ex0726

##	Gender	Family	Height	Father	Mother
## 1	male	1	73.2	78.5	67.0
## 2	female	1	69.2	78.5	67.0
## 3	female	1	69.0	78.5	67.0
## 4	female	1	69.0	78.5	67.0
## 5	male	2	73.5	75.5	66.5
## 6	male	2	72.5	75.5	66.5
## 7	female	2	65.5	75.5	66.5
## 8	female	2	65.5	75.5	66.5
## 9	male	3	71.0	75.0	64.0
## 10	female	3	68.0	75.0	64.0
## 11	male	4	70.5	75.0	64.0
## 12	male	4	68.5	75.0	64.0
## 13	female	4	67.0	75.0	64.0
## 14	female	4	64.5	75.0	64.0
## 15	female	4	63.0	75.0	64.0
## 16	male	5	72.0	75.0	58.5
## 17	male	5	69.0	75.0	58.5

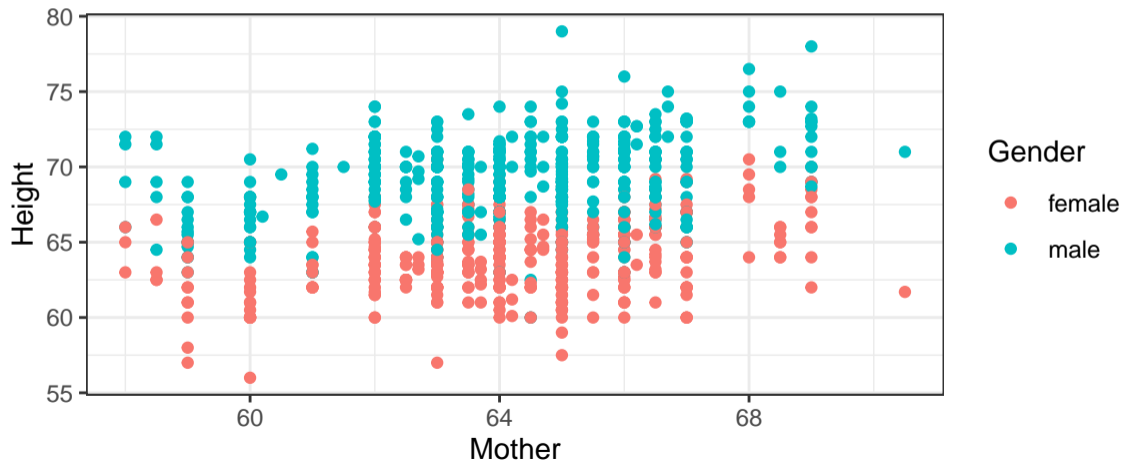
heights

```
summary(ex0726)
```

##	Gender	Family	Height	Father	Mother
##	female:453	Min. : 1.0	Min. :56.00	Min. :62.0	Min. :58.00
##	male :480	1st Qu.: 59.0	1st Qu.:64.00	1st Qu.:68.0	1st Qu.:63.00
##		Median :106.0	Median :66.50	Median :69.0	Median :64.00
##		Mean :106.2	Mean :66.74	Mean :69.2	Mean :64.09
##		3rd Qu.:158.0	3rd Qu.:69.70	3rd Qu.:71.0	3rd Qu.:65.50
##		Max. :205.0	Max. :79.00	Max. :78.5	Max. :70.50

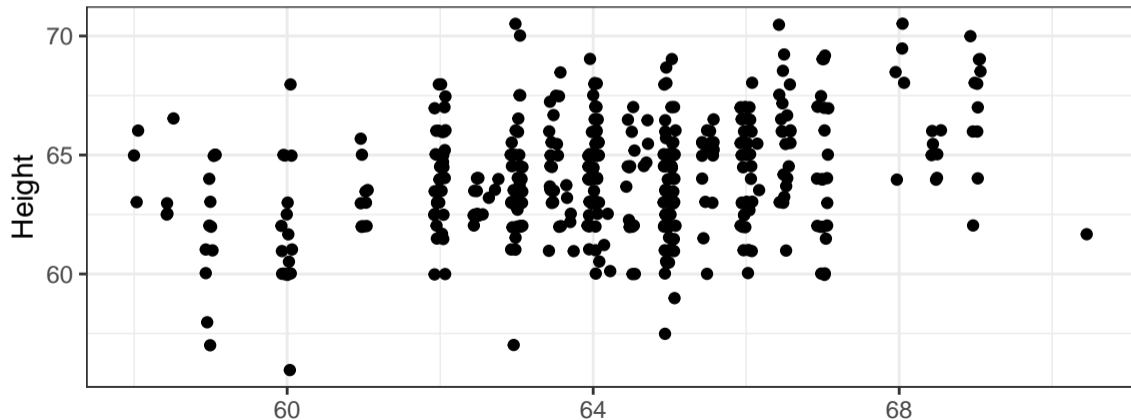
heights

```
ggplot(ex0726, aes(x=Mother, y=Height, color = Gender)) + geom_point()
```



heights

```
ggplot(ex0726 %>% filter(Gender == "female"),  
  aes(x=Mother, y=Height)) + geom_jitter()
```



```
m <- lm(Height ~ Mother, data = ex0726 %>% filter(Gender == "female"),)
summary(m)

##
## Call:
## lm(formula = Height ~ Mother, data = ex0726 %>% filter(Gender ==
##   "female"))
##
## Residuals:
##      Min       1Q   Median       3Q      Max
## -6.8739 -1.5331  0.0813  1.4445  6.7629
##
## Coefficients:
##              Estimate Std. Error t value Pr(>|t|)
## (Intercept) 43.67884    3.00214  14.549 < 2e-16 ***
## Mother       0.31839    0.04677   6.807 3.18e-11 ***
## ---
## Signif. codes:  0 '***' 0.001 '**' 0.01 '*' 0.05 '.' 0.1 ' ' 1
##
## Residual standard error: 2.246 on 451 degrees of freedom
## Multiple R-squared:  0.09318, Adjusted R-squared:  0.09117
## F-statistic: 46.34 on 1 and 451 DF,  p-value: 3.181e-11
```

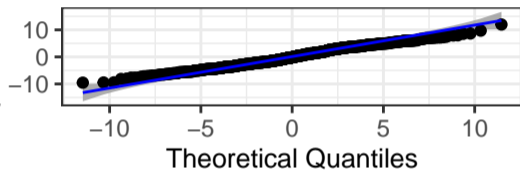

heights

```
m <- lm(Height ~ I(Mother-64), data = ex0726)
```

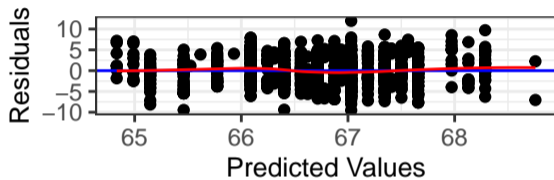
```
resid_panel(m, plots = c("qq", "resid", "index", "cookd"), smooth = TRUE, qqbands = TRUE)
```

Sample Quantiles

Q-Q Plot

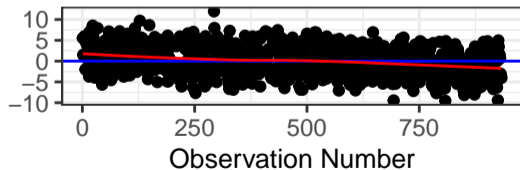


Residual Plot

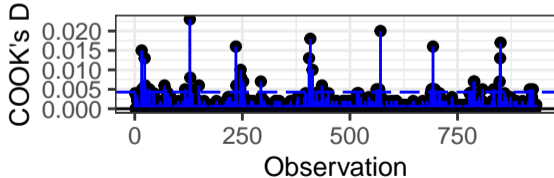


Residuals

Index Plot



COOK's D Plot



```
summary(m)
```

```
##
## Call:
## lm(formula = Height ~ I(Mother - 64), data = ex0726)
##
## Residuals:
##      Min       1Q   Median       3Q      Max
## -9.530 -2.629 -0.146  2.871 11.970
##
## Coefficients:
##              Estimate Std. Error t value Pr(>|t|)
## (Intercept)   66.71614    0.11499  580.185 < 2e-16 ***
## I(Mother - 64)  0.31403    0.05019   6.256 6.01e-10 ***
## ---
## Signif. codes:  0 '***' 0.001 '**' 0.01 '*' 0.05 '.' 0.1 ' ' 1
##
## Residual standard error: 3.51 on 931 degrees of freedom
## Multiple R-squared:  0.04034, Adjusted R-squared:  0.03931
## F-statistic: 39.14 on 1 and 931 DF,  p-value: 6.006e-10
```

heights

```
co <- coef(m)
co

##      (Intercept) I(Mother - 64)
##      66.7161395      0.3140259

ci <- confint(m)
ci

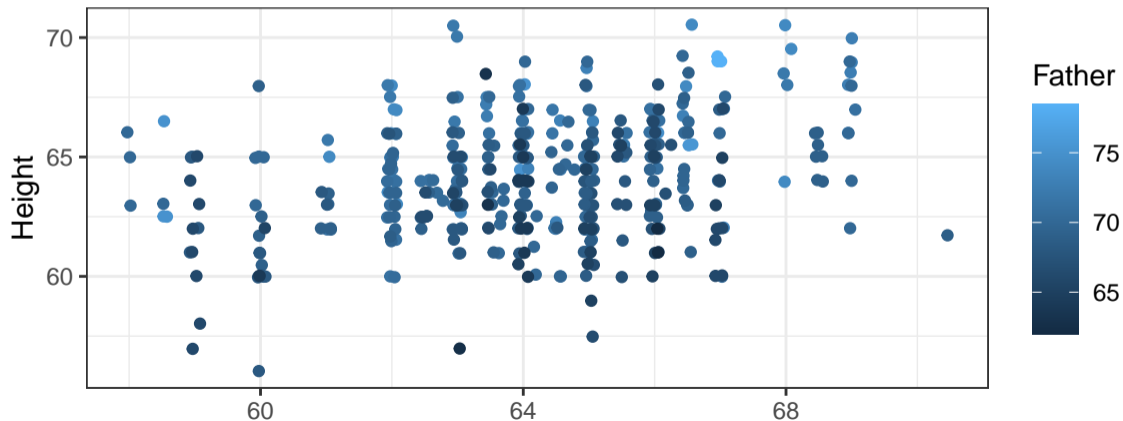
##              2.5 %      97.5 %
## (Intercept) 66.490468 66.9418112
## I(Mother - 64) 0.215519 0.4125328
```

heights

- When the mother's height is 64 in, the mean daughter's height is 66.7 in (66.5,66.9).
- For each inch increase in mother's height, the mean daughter's height increases by 0.31 (0.22,0.41) inches.
- The model with mother's height explains 4% of the variability in daughter's height.

heights

```
ggplot(ex0726 %>% filter(Gender == "female"),  
  aes(x=Mother, y=Height, color = Father)) + geom_jitter()
```



```
m <- lm(Height ~ Mother + Father, data = ex0726 %>% filter(Gender == "female"))
summary(m)
```

```
##
## Call:
## lm(formula = Height ~ Mother + Father, data = ex0726 %>% filter(Gender ==
##   "female"))
##
## Residuals:
##      Min       1Q   Median       3Q      Max
## -6.3726 -1.4040 -0.0423  1.4130  6.9325
##
## Coefficients:
##              Estimate Std. Error t value Pr(>|t|)
## (Intercept)  18.75770    3.60782   5.199 3.04e-07 ***
## Mother        0.30358    0.04206   7.218 2.27e-12 ***
## Father        0.37353    0.03590  10.406 < 2e-16 ***
## ---
## Signif. codes:  0 '***' 0.001 '**' 0.01 '*' 0.05 '.' 0.1 ' ' 1
##
## Residual standard error: 2.019 on 450 degrees of freedom
## Multiple R-squared:  0.2691, Adjusted R-squared:  0.2658
## F-statistic: 82.82 on 2 and 450 DF,  p-value: < 2.2e-16
```

heights

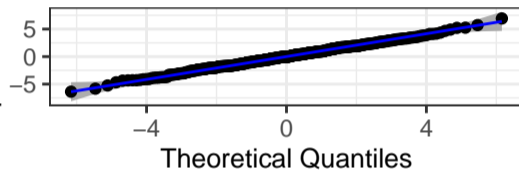
```
m <- lm(Height ~ I(Mother-64) + I(Father-69), data = ex0726 %>% filter(Gender == "female"))
```



```
resid_panel(m, plots = c("qq", "resid", "index", "cookd"), smooth = TRUE, qqbands = TRUE)
```

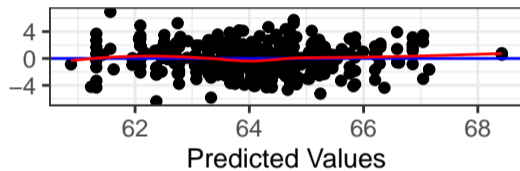
Sample Quantiles

Q-Q Plot



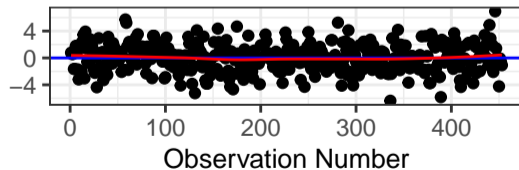
Residuals

Residual Plot



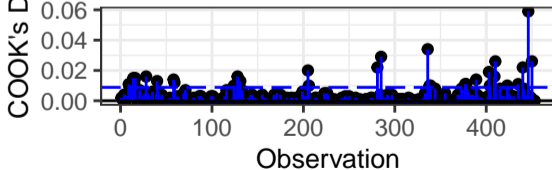
Residuals

Index Plot



COOK's D

COOK's D Plot



```
summary(m)
```

```
##  
## Call:  
## lm(formula = Height ~ I(Mother - 64) + I(Father - 69), data = ex0726 %>%  
##   filter(Gender == "female"))  
##  
## Residuals:  
##      Min       1Q   Median       3Q      Max   
## -6.3726 -1.4040 -0.0423  1.4130  6.9325   
##  
## Coefficients:  
##              Estimate Std. Error t value Pr(>|t|)      
## (Intercept)   63.96048    0.09549  669.786 < 2e-16 ***  
## I(Mother - 64)  0.30358    0.04206   7.218 2.27e-12 ***  
## I(Father - 69)  0.37353    0.03590  10.406 < 2e-16 ***  
## ---  
## Signif. codes:  0 '***' 0.001 '**' 0.01 '*' 0.05 '.' 0.1 ' ' 1  
##  
## Residual standard error: 2.019 on 450 degrees of freedom  
## Multiple R-squared:  0.2691, Adjusted R-squared:  0.2658   
## F-statistic: 82.82 on 2 and 450 DF,  p-value: < 2.2e-16
```

heights

```
co <- coef(m)
co

##      (Intercept) I(Mother - 64) I(Father - 69)
##      63.9604846      0.3035837      0.3735279

ci <- confint(m)
ci

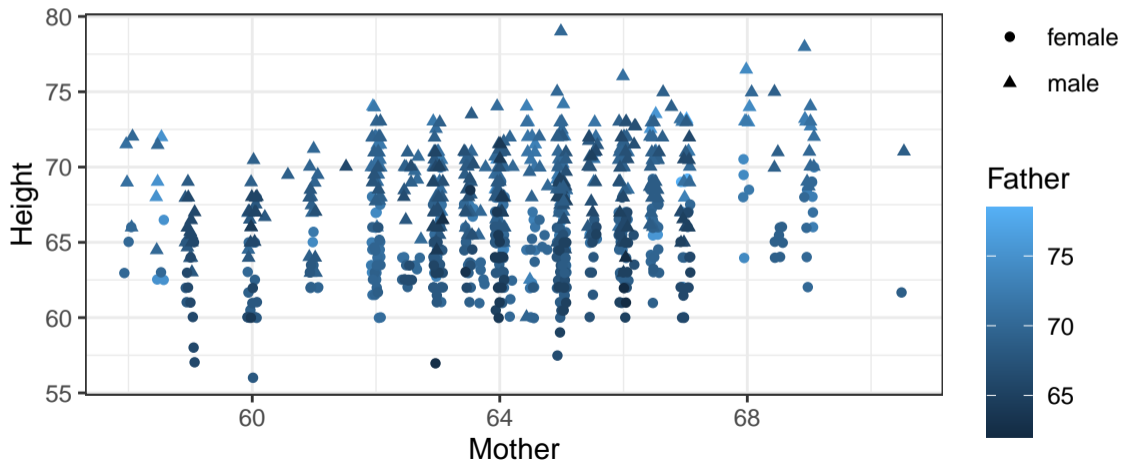
##              2.5 %      97.5 %
## (Intercept)  63.7728152  64.1481540
## I(Mother - 64)  0.2209226  0.3862448
## I(Father - 69)  0.3029841  0.4440717
```

heights

- When mother's height is 64 in and father's height is 69 in, the mean daughter's height is 64 in (63.8,64.1).
- For each inch increase in mother's height, the mean children's height increases by 0.3 (0.22,0.39) inches while holding father's height constant.
- For each inch increase in father's height, the mean daughter's height increases by 0.37 (0.3,0.44) inches while holding mother's height constant.
- The model with mother's and father's height explains 27% of the variability in daughter's height.

heights

```
ggplot(ex0726, aes(x=Mother, y=Height, color = Father, shape=Gender)) + geom_jitter()
```



```
m <- lm(Height ~ Mother + Father + Gender, data = ex0726)
summary(m)

##
## Call:
## lm(formula = Height ~ Mother + Father + Gender, data = ex0726)
##
## Residuals:
##      Min       1Q   Median       3Q      Max
## -9.5280 -1.4604  0.0996  1.4783  9.1161
##
## Coefficients:
##              Estimate Std. Error t value Pr(>|t|)
## (Intercept) 16.43221    2.72802   6.023 2.46e-09 ***
## Mother       0.31840    0.03102  10.263 < 2e-16 ***
## Father       0.39339    0.02868  13.718 < 2e-16 ***
## Gendermale   5.21902    0.14188  36.784 < 2e-16 ***
## ---
## Signif. codes:  0 '***' 0.001 '**' 0.01 '*' 0.05 '.' 0.1 ' ' 1
##
## Residual standard error: 2.165 on 929 degrees of freedom
## Multiple R-squared:  0.6358, Adjusted R-squared:  0.6346
## F-statistic: 540.5 on 3 and 929 DF,  p-value: < 2.2e-16
```

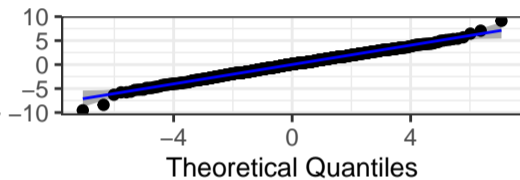
heights

```
m <- lm(Height ~ I(Mother-64) + I(Father-69) + Gender, data = ex0726)
```

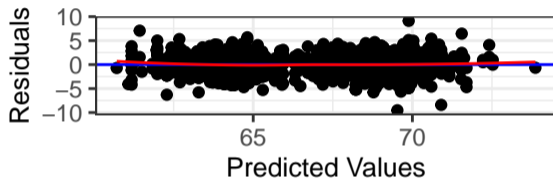
```
resid_panel(m, plots = c("qq", "resid", "index", "cookd"), smooth = TRUE, qqbands = TRUE)
```

Sample Quantiles

Q-Q Plot

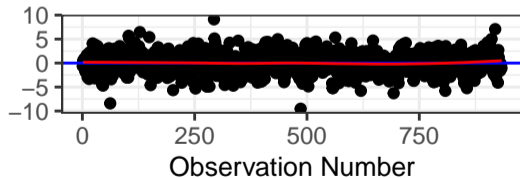


Residual Plot

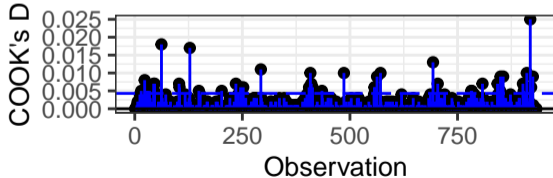


Residuals

Index Plot



COOK's D Plot




```
summary(m)
```

```
##
## Call:
## lm(formula = Height ~ I(Mother - 64) + I(Father - 69) + Gender,
##     data = ex0726)
##
## Residuals:
##      Min       1Q   Median       3Q      Max
## -9.5280 -1.4604  0.0996  1.4783  9.1161
##
## Coefficients:
##              Estimate Std. Error t value Pr(>|t|)
## (Intercept)   63.95311    0.10206   626.61 <2e-16 ***
## I(Mother - 64)  0.31840    0.03102   10.26 <2e-16 ***
## I(Father - 69)  0.39339    0.02868   13.72 <2e-16 ***
## Gendermale     5.21902    0.14188   36.78 <2e-16 ***
## ---
## Signif. codes:  0 '***' 0.001 '**' 0.01 '*' 0.05 '.' 0.1 ' ' 1
##
## Residual standard error: 2.165 on 929 degrees of freedom
## Multiple R-squared:  0.6358, Adjusted R-squared:  0.6346
## F-statistic: 540.5 on 3 and 929 DF,  p-value: < 2.2e-16
```

heights

```
co <- coef(m)
```

```
co
```

```
##      (Intercept) I(Mother - 64) I(Father - 69)      Gendermale
##      63.9531126      0.3183957      0.3933851      5.2190191
```

```
ci <- confint(m)
```

```
ci
```

```
##              2.5 %      97.5 %
## (Intercept)  63.7528127  64.1534126
## I(Mother - 64)  0.2575128  0.3792787
## I(Father - 69)  0.3371065  0.4496638
## Gendermale     4.9405709  5.4974674
```

heights

- When the mother's height is 64 in and father's height is 69 in, the mean daughter's height is 64 in (63.8,64.2).
- For each inch increase in mother's height, the mean child's height increases by 0.32 (0.26,0.38) inches while holding father's height and gender constant.
- For each inch increase in father's height, the mean child's height increases by 0.39 (0.34,0.45) inches while holding mother's height and gender constant.
- Male children are, on average, 5.22 (4.94,5.5) inches taller than female children while holding mother's and father's height constant.
- The model with mother's height, father's height, and gender explains 64% of the variability in child's height.

warpbreaks

```
em <- emmeans(m, pairwise ~ Gender, adjust = "none")
confint(em)

## $emmeans
##   Gender emmean      SE df lower.CL upper.CL
## female  64.06 0.10174 929   63.86   64.26
## male    69.28 0.09883 929   69.08   69.47
##
## Confidence level used: 0.95
##
## $contrasts
##   contrast      estimate      SE df lower.CL upper.CL
## female - male    -5.22 0.142 929   -5.5   -4.94
##
## Confidence level used: 0.95
```

warpbreaks

```
warpbreaks
```

```
##      breaks wool tension
## 1      26     A         L
## 2      30     A         L
## 3      54     A         L
## 4      25     A         L
## 5      70     A         L
## 6      52     A         L
## 7      51     A         L
## 8      26     A         L
## 9      67     A         L
## 10     18     A         M
## 11     21     A         M
## 12     29     A         M
## 13     17     A         M
## 14     12     A         M
## 15     18     A         M
## 16     35     A         M
## 17     30     A         M
```

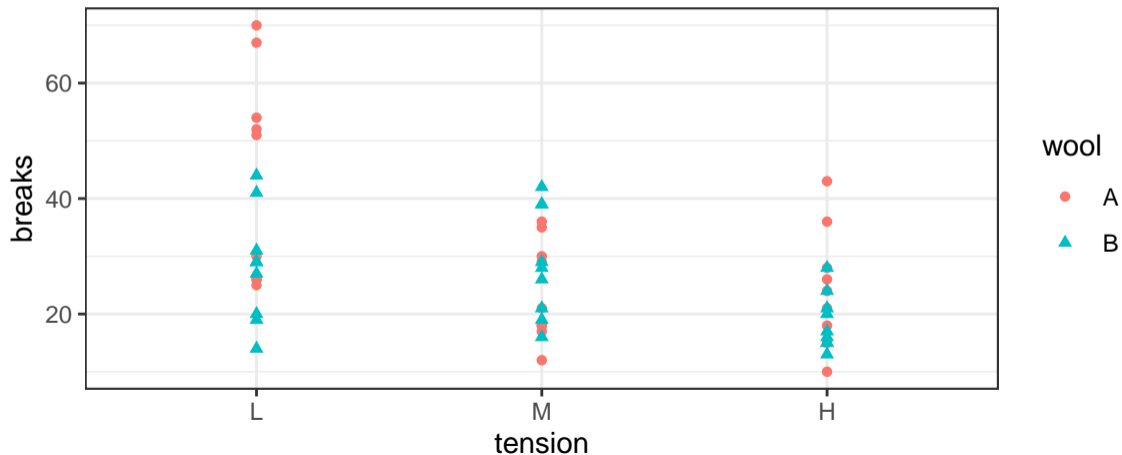
warpbreaks

```
summary(warpbreaks)
```

```
##      breaks      wool  tension
##  Min.    :10.00   A:27    L:18
##  1st Qu.:18.25   B:27    M:18
##  Median :26.00           H:18
##  Mean   :28.15
##  3rd Qu.:34.00
##  Max.   :70.00
```

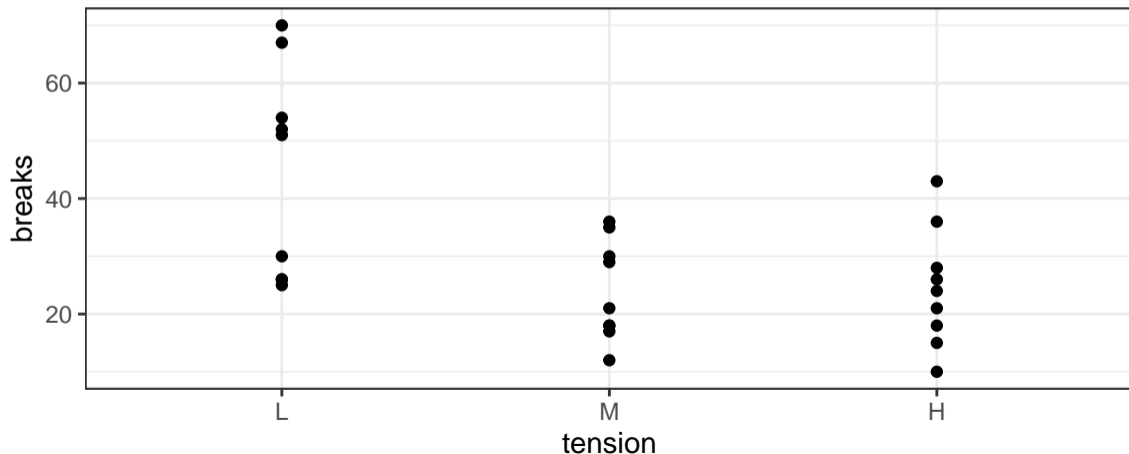
warpbreaks

```
ggplot(warpbreaks, aes(x=tension, y=breaks, color=wool, shape=wool)) + geom_point()
```



warpbreaks

```
ggplot(warpbreaks %>% filter(wool == "A"), aes(x=tension, y=breaks)) + geom_point()
```



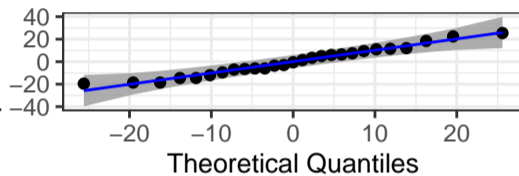
warpbreaks

```
m <- lm(breaks ~ tension, data = warpbreaks %>% filter(wool == "A"))
```

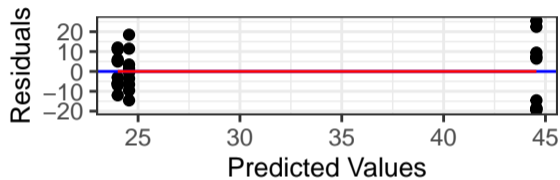
```
resid_panel(m, plots = c("qq", "resid", "index", "cookd"), smooth = TRUE, qqbands = TRUE)
```

Sample Quantiles

Q-Q Plot

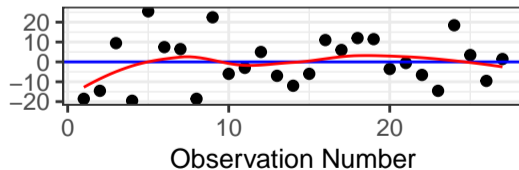


Residual Plot



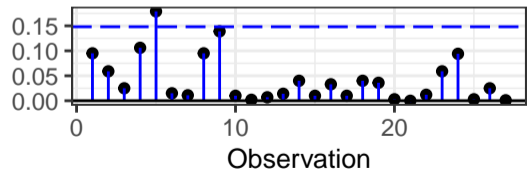
Residuals

Index Plot



COOK's D

COOK's D Plot



```
summary(m)
```

```
##
## Call:
## lm(formula = breaks ~ tension, data = warpbreaks %>% filter(wool ==
##   "A"))
##
## Residuals:
##      Min       1Q   Median       3Q      Max
## -19.5556  -8.2778  -0.5556   8.4444  25.4444
##
## Coefficients:
##              Estimate Std. Error t value Pr(>|t|)
## (Intercept)   44.556     4.338  10.271 2.91e-10 ***
## tensionM     -20.556     6.135  -3.351 0.00266 **
## tensionH     -20.000     6.135  -3.260 0.00332 **
## ---
## Signif. codes:  0 '***' 0.001 '**' 0.01 '*' 0.05 '.' 0.1 ' ' 1
##
## Residual standard error: 13.01 on 24 degrees of freedom
## Multiple R-squared:  0.3779, Adjusted R-squared:  0.326
## F-statistic: 7.288 on 2 and 24 DF,  p-value: 0.003363
```

warpbreaks

```
co <- coef(m)
co

## (Intercept)    tensionM    tensionH
##    44.55556    -20.55556    -20.00000

ci <- confint(m)
ci

##              2.5 %    97.5 %
## (Intercept)  35.60269  53.508421
## tensionM    -33.21682  -7.894291
## tensionH    -32.66126  -7.338736
```

warpbreaks

- For wool type A, when tension is low the mean number of breaks is 45 (36,54).
- For wool type A, moving from low to medium causes the mean number of breaks to decrease by 21 (8,33).
- For wool type A, moving from low to high causes the mean number of breaks to decrease by 20 (7,33).
- For wool type A, The model with tension 38% of the variability in number of breaks.

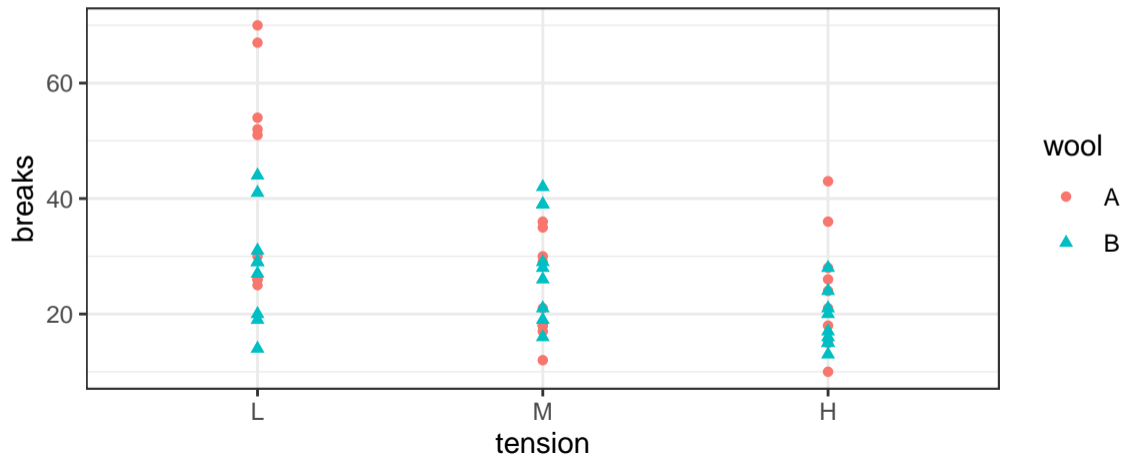
warpbreaks

```
em <- emmeans(m, pairwise ~ tension, adjust = "none")
confint(em)

## $emmeans
##   tension emmean   SE df lower.CL upper.CL
##   L         44.6 4.34 24    35.6    53.5
##   M         24.0 4.34 24    15.0    33.0
##   H         24.6 4.34 24    15.6    33.5
##
## Confidence level used: 0.95
##
## $contrasts
##   contrast estimate   SE df lower.CL upper.CL
##   L - M      20.556 6.13 24     7.89    33.2
##   L - H      20.000 6.13 24     7.34    32.7
##   M - H      -0.556 6.13 24    -13.22    12.1
##
## Confidence level used: 0.95
```

warpbreaks

```
ggplot(warpbreaks, aes(x=tension, y=breaks, color=wool, shape=wool)) + geom_point()
```



warpbreaks

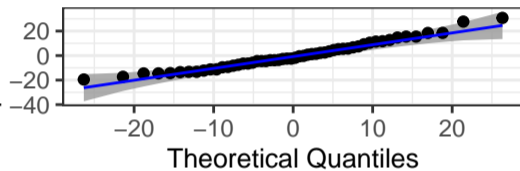
```
m <- lm(breaks ~ tension + wool, data = warpbreaks)
```



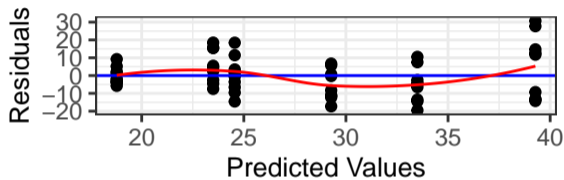
```
resid_panel(m, plots = c("qq", "resid", "index", "cookd"), smooth = TRUE, qqbands = TRUE)
```

Sample Quantiles

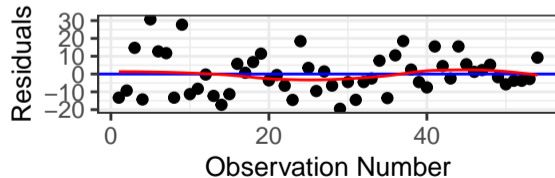
Q-Q Plot



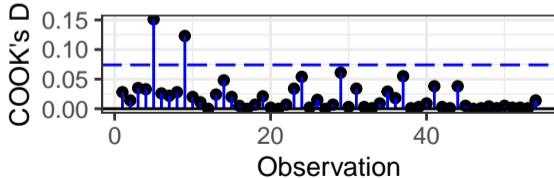
Residual Plot



Index Plot



COOK's D Plot



```
summary(m)
```

```
##
## Call:
## lm(formula = breaks ~ tension + wool, data = warpbreaks)
##
## Residuals:
##      Min       1Q   Median       3Q      Max
## -19.500  -8.083  -2.139   6.472  30.722
##
## Coefficients:
##              Estimate Std. Error t value Pr(>|t|)
## (Intercept)   39.278     3.162  12.423 < 2e-16 ***
## tensionM     -10.000     3.872  -2.582  0.012787 *
## tensionH     -14.722     3.872  -3.802  0.000391 ***
## woolB         -5.778     3.162  -1.827  0.073614 .
## ---
## Signif. codes:  0 '***' 0.001 '**' 0.01 '*' 0.05 '.' 0.1 ' ' 1
##
## Residual standard error: 11.62 on 50 degrees of freedom
## Multiple R-squared:  0.2691, Adjusted R-squared:  0.2253
## F-statistic: 6.138 on 3 and 50 DF,  p-value: 0.00123
```

warpbreaks

```
co <- coef(m)
co

## (Intercept)    tensionM    tensionH        woolB
##  39.277778   -10.000000   -14.722222   -5.777778

ci <- confint(m)
ci

##           2.5 %    97.5 %
## (Intercept)  32.92715  45.6284061
## tensionM    -17.77790  -2.2221006
## tensionH    -22.50012  -6.9443228
## woolB       -12.12841   0.5728505
```

warpbreaks

- For wool type A and low tension, the mean number of breaks is 39 (33,46).
- Moving from low to medium causes the mean number of breaks to decrease by 10 (2,18) for both wool types.
- Moving from low to high causes the mean number of breaks to decrease by 15 (7,23) for both wool types.
- On average, wool type B has 6 (-1,12) fewer breaks than wool type A.
- The model with tension and wool type explains 27% of the variability in number of breaks.

warpbreaks

```
em <- emmeans(m, pairwise ~ tension, adjust = "none")
confint(em)
```

```
## $emmeans
```

```
##   tension emmean   SE df lower.CL upper.CL
##   L           36.4 2.74 50    30.9    41.9
##   M           26.4 2.74 50    20.9    31.9
##   H           21.7 2.74 50    16.2    27.2
```

```
##
## Results are averaged over the levels of: wool
## Confidence level used: 0.95
```

```
##
## $contrasts
```

```
##   contrast estimate   SE df lower.CL upper.CL
##   L - M           10.00 3.87 50    2.22    17.8
##   L - H           14.72 3.87 50    6.94    22.5
##   M - H            4.72 3.87 50   -3.06    12.5
```

```
##
## Results are averaged over the levels of: wool
## Confidence level used: 0.95
```

warpbreaks

```
em <- emmeans(m, pairwise ~ wool, adjust = "none")
confint(em)

## $emmeans
##   wool emmean   SE df lower.CL upper.CL
##   A      31.0 2.24 50    26.5    35.5
##   B      25.3 2.24 50    20.8    29.7
##
## Results are averaged over the levels of: tension
## Confidence level used: 0.95
##
## $contrasts
##   contrast estimate   SE df lower.CL upper.CL
##   A - B           5.78 3.16 50   -0.573    12.1
##
## Results are averaged over the levels of: tension
## Confidence level used: 0.95
```

warpbreaks

```
em <- emmeans(m, pairwise ~ tension | wool, adjust = "none")
confint(em)$emmeans

## wool = A:
##   tension emmean   SE df lower.CL upper.CL
##   L         39.3 3.16 50    32.9    45.6
##   M         29.3 3.16 50    22.9    35.6
##   H         24.6 3.16 50    18.2    30.9
##
## wool = B:
##   tension emmean   SE df lower.CL upper.CL
##   L         33.5 3.16 50    27.1    39.9
##   M         23.5 3.16 50    17.1    29.9
##   H         18.8 3.16 50    12.4    25.1
##
## Confidence level used: 0.95
```

warpbreaks

```
confint(em)$contrasts
```

```
## wool = A:
```

```
## contrast estimate SE df lower.CL upper.CL
```

```
## L - M 10.00 3.87 50 2.22 17.8
```

```
## L - H 14.72 3.87 50 6.94 22.5
```

```
## M - H 4.72 3.87 50 -3.06 12.5
```

```
##
```

```
## wool = B:
```

```
## contrast estimate SE df lower.CL upper.CL
```

```
## L - M 10.00 3.87 50 2.22 17.8
```

```
## L - H 14.72 3.87 50 6.94 22.5
```

```
## M - H 4.72 3.87 50 -3.06 12.5
```

```
##
```

```
## Confidence level used: 0.95
```


warpbreaks

```
em <- emmeans(m, pairwise ~ wool | tension, adjust = "none")
confint(em)$emmeans
```

```
## tension = L:
## wool emmean SE df lower.CL upper.CL
## A      39.3 3.16 50     32.9    45.6
## B      33.5 3.16 50     27.1    39.9
##
## tension = M:
## wool emmean SE df lower.CL upper.CL
## A      29.3 3.16 50     22.9    35.6
## B      23.5 3.16 50     17.1    29.9
##
## tension = H:
## wool emmean SE df lower.CL upper.CL
## A      24.6 3.16 50     18.2    30.9
## B      18.8 3.16 50     12.4    25.1
##
## Confidence level used: 0.95
```

warpbreaks

```
confint(em)$contrasts

## tension = L:
## contrast estimate SE df lower.CL upper.CL
## A - B          5.78 3.16 50  -0.573    12.1
##
## tension = M:
## contrast estimate SE df lower.CL upper.CL
## A - B          5.78 3.16 50  -0.573    12.1
##
## tension = H:
## contrast estimate SE df lower.CL upper.CL
## A - B          5.78 3.16 50  -0.573    12.1
##
## Confidence level used: 0.95
```