

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

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

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

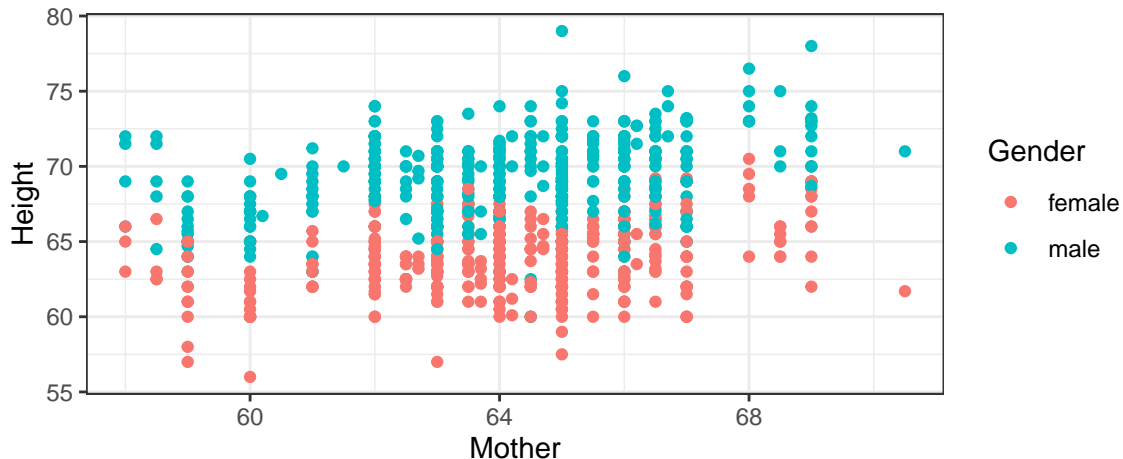
heights

```
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

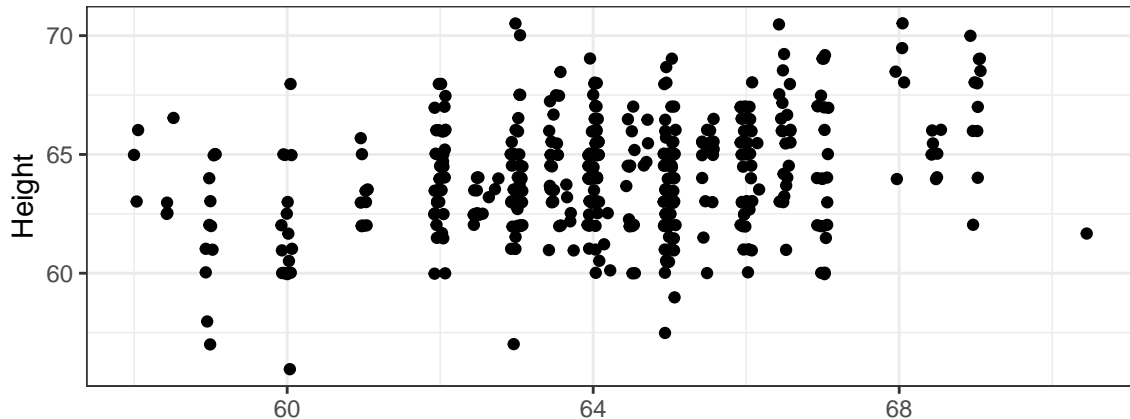
heights

```
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

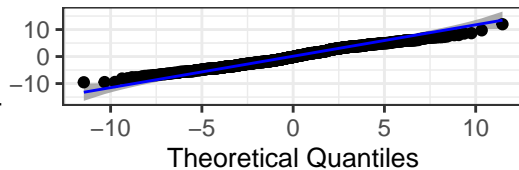
```


heights

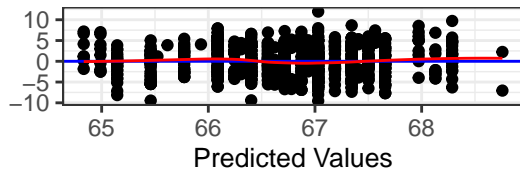
```
m <- lm(Height ~ I(Mother-64), data = ex0726)
```

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

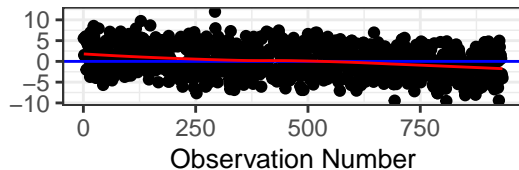
Sample Quantiles

Q-Q Plot

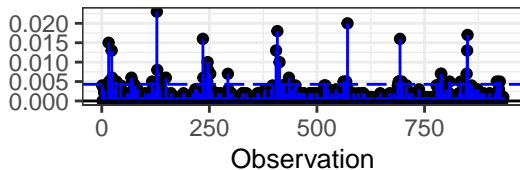
Residuals

Residual Plot

Residuals

Index Plot

COOK's D

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

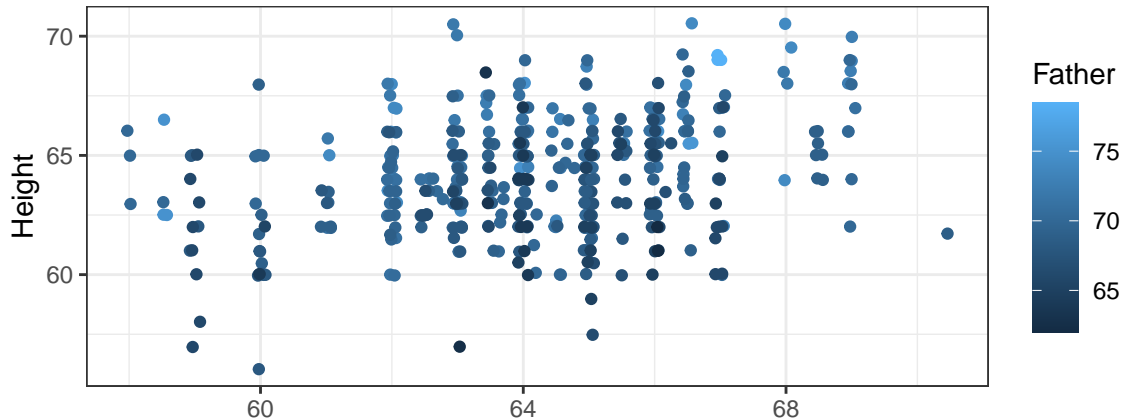
##              2.5 %      97.5 %
## (Intercept)  66.490468 66.9418112
## I(Mother - 64) 0.215519 0.4125328
```

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

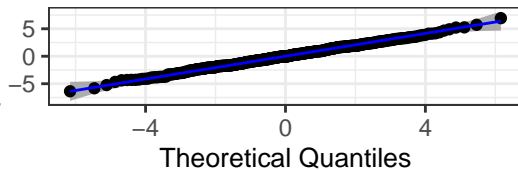
```
m <- lm(Height ~ I(Mother-64) + I(Father-69), data = ex0726 %>% filter(Gender == "female"))
```



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

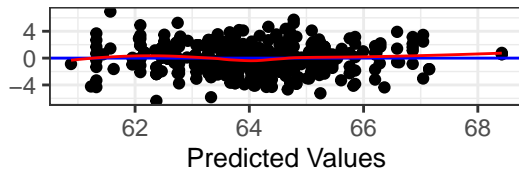
Sample Quantiles

Q-Q Plot



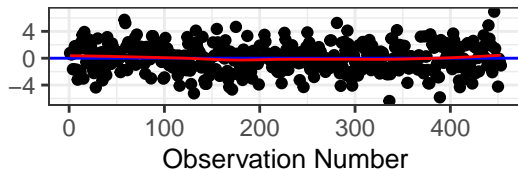
Residuals

Residual Plot



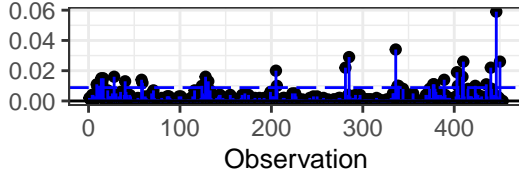
Residuals

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
```

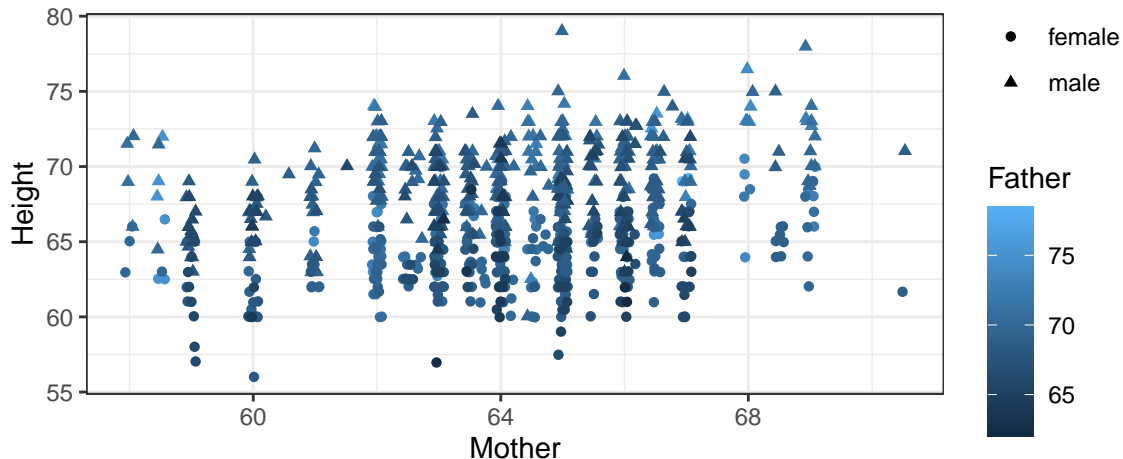
```
## I(Father - 69) 0.3029841 0.4440717
```

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
```

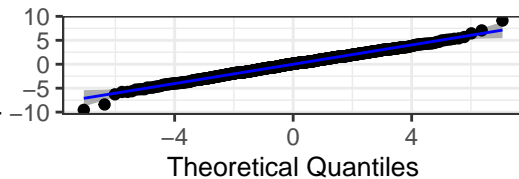
heights

```
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)
```

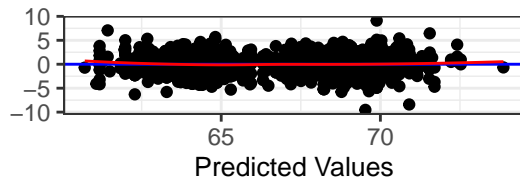
Sample Quantiles

Q-Q Plot



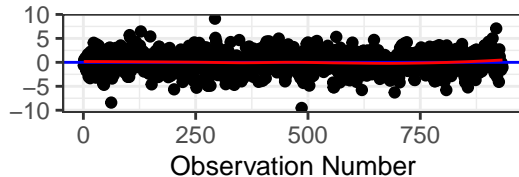
Residuals

Residual Plot



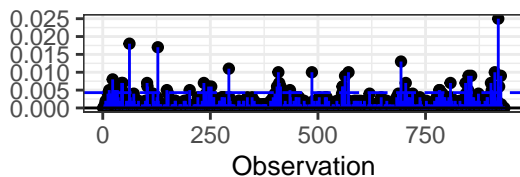
Residuals

Index Plot



COOK's D

COOK's D Plot




```
summary(m)

##
## Call:
## lm(formula = Height ~ I(Mother - 64) + I(Father - 69) + 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 daughter'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 daughter's height increases by 0.39 (0.34,0.45) inches while holding mother's height and gender constant.
- Male children are, on average, 0.39 (0.34,0.45) 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 children's height.

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

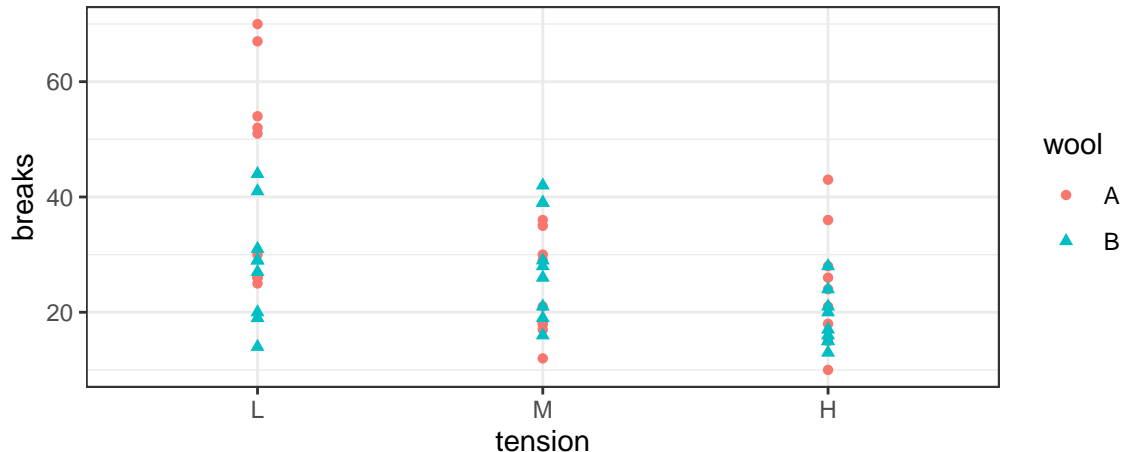
warpbreaks

```
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
```

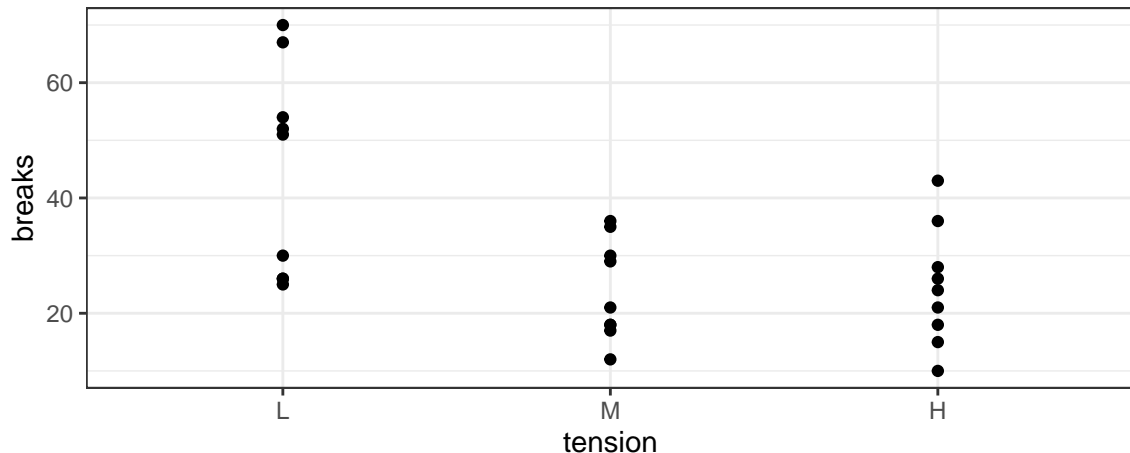
warpbreaks

```
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

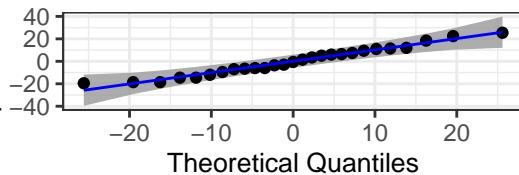
```
m <- lm(breaks ~ tension, data = warpbreaks %>% filter(wool == "A"))
```



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

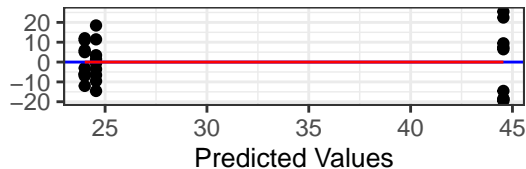
Sample Quantiles

Q-Q Plot



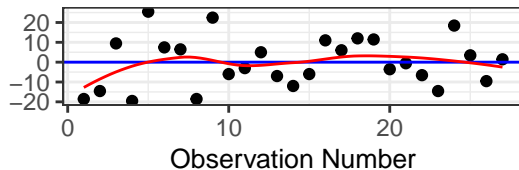
Residuals

Residual Plot



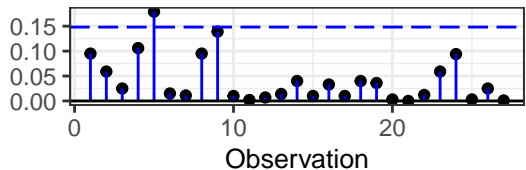
Residuals

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

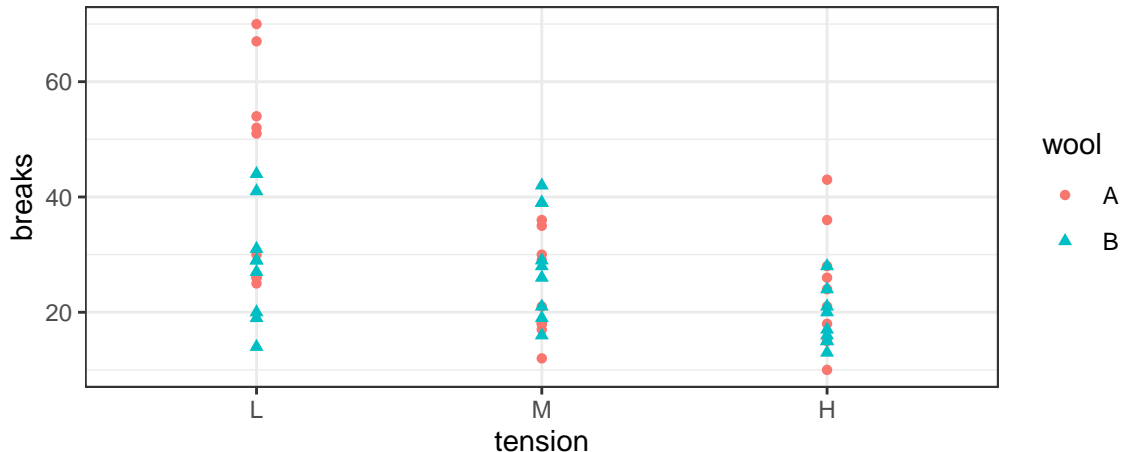
warpbreaks

```
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()
```

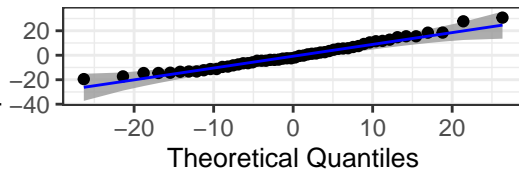


warpbreaks

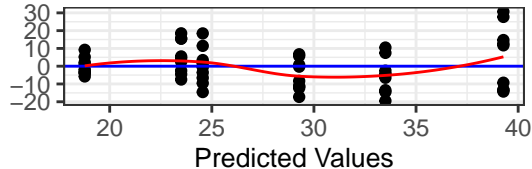
```
m <- lm(breaks ~ tension + wool, data = warpbreaks)
```

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

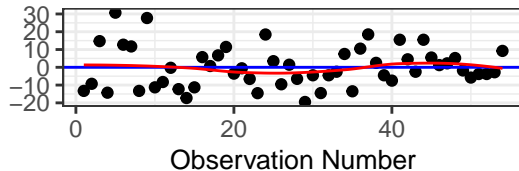
Sample Quantiles

Q-Q Plot

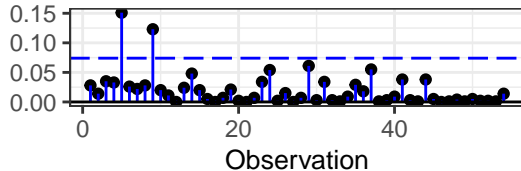
Residuals

Residual Plot

Residuals

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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 when tension is low, the mean number of breaks is 39 in (33,46).
- When moving from tension low to tension medium, the mean number of breaks is 10 (2,18) lower while holding wool type constant.
- When moving from tension low to tension high, the mean number of breaks is 15 (7,23) lower while holding wool type constant.
- 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
```