R09 - Poisson Regression

HCI/PSYCH 522 Iowa State University

April 14, 2022

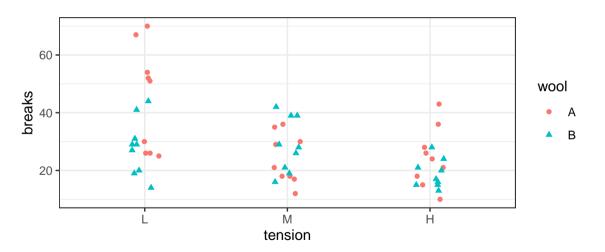
Overview

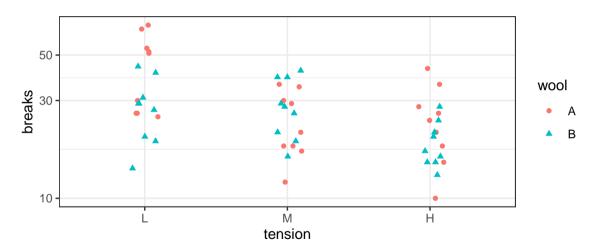
- Linear regression (with logarithm of breaks)
 - Tension
 - + Wool Type
- Poisson regression
 - Tension
 - + Wool Type

warpbreaks				
##		breaks	wool	tension
##	1	26	Α	L
##	2	30	Α	L
##	3	54	A	L
##	4	25	A	L
##	5	70	Α	L
##	6	52	A	L
##	7	51	A	L
##	8	26	A	L
##	9	67	Α	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

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
```





Linear regression with log of dependent variable

Regression with

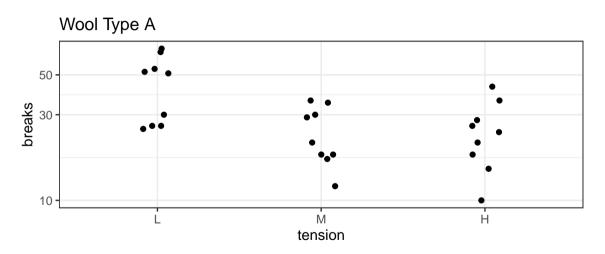
$$\log(Y_i) \stackrel{ind}{\sim} N(\mu_i, \sigma^2), \quad \mu_i = \beta_0 + \beta_1 X_{i,1} + \dots + \beta_p X_{i,p}$$

where, for observation i,

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

Interpretation

- ullet e^{eta_0} is median of the dependent variable when all independent variables (X's) are 0
- $100(e^{\beta_p}-1)$ for $p\neq 0$, is the percent increase in the median of the dependent variable for each unit increase in the associated independent variable
- The coefficient of determination, R^2 , is the proportion of variability in the logarithm of the dependent variable explained by the model.



```
##
## Call:
## lm(formula = log(breaks) ~ tension, data = warpbreaks %>% filter(wool ==
##
      "A"))
##
## Residuals:
##
       Min 10 Median 30
                                        Max
## -0.81504 -0.30014 0.06043 0.27774 0.64358
##
## Coefficients:
##
             Estimate Std. Error t value Pr(>|t|)
## (Intercept) 3.7179 0.1394 26.678 < 2e-16 ***
## tensionM -0.6012 0.1971 -3.050 0.00550 **
## tensionH -0.6003 0.1971 -3.046 0.00556 **
## ---
## Signif. codes: 0 '***' 0.001 '**' 0.05 '.' 0.1 ' ' 1
##
## Residual standard error: 0.4181 on 24 degrees of freedom
## Multiple R-squared: 0.3404, Adjusted R-squared: 0.2855
## F-statistic: 6.194 on 2 and 24 DF, p-value: 0.006777
```

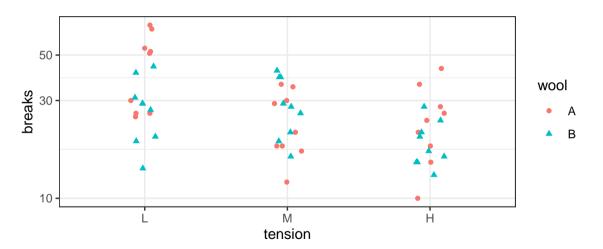
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```
em <- emmeans(m, pairwise ~ tension)</pre>
ci <- confint(em, type = "response", adjust = "none")</pre>
ci
## $emmeans
  tension response SE df lower.CL upper.CL
              41.2 5.74 24
                              30.9
##
                                      54.9
           22.6 3.15 24 16.9 30.1
## M
          22.6 3.15 24 16.9 30.1
##
  H
##
## Confidence level used: 0.95
## Intervals are back-transformed from the log scale
##
## $contrasts
   contrast ratio SE df lower.CL upper.CL
   L / M 1.824 0.360 24 1.215
                                 2.74
  L / H 1.823 0.359 24 1.214 2.74
##
##
   M / H 0.999 0.197 24 0.665 1.50
##
## Confidence level used: 0.95
## Intervals are back-transformed from the log scale
```

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- For wool type A, the median number of breaks when
 - tension is low is 41 (31,55),
 - tension is medium is 23 (17,30),
 - tension is high is 23 (17,30),
- For wool type A, tension
 - low compared to medium causes an 82% (21,174) increase in median number of breaks,
 - low compared to high causes an 82% (21,174) increase in median number of breaks, and
 - medium compared to high causes an 0% (-33,50) increase in median number of breaks.



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warpbreaks

```
##
## Call:
## lm(formula = log(breaks) ~ tension + wool, data = warpbreaks)
##
## Residuals:
##
       Min
                10 Median 30
                                        Max
## -0.80421 -0.29975 -0.01627 0.28367 0.67424
##
## Coefficients:
##
             Estimate Std. Error t value Pr(>|t|)
## (Intercept) 3.5762 0.1063 33.644 < 2e-16 ***
## tensionM -0.2871 0.1302 -2.205 0.032048 *
## tensionH -0.4893 0.1302 -3.758 0.000448 ***
## woolB
         -0.1522 0.1063 -1.431 0.158540
## ---
## Signif. codes: 0 '***' 0.001 '**' 0.05 '.' 0.1 ' ' 1
##
## Residual standard error: 0.3906 on 50 degrees of freedom
## Multiple R-squared: 0.246, Adjusted R-squared: 0.2008
## F-statistic: 5.438 on 3 and 50 DF, p-value: 0.002576
```

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```
em_tension <- emmeans(m, pairwise ~ tension)</pre>
ci tension <- confint(em tension, type = "response", adjust = "none")
ci tension
## $emmeans
  tension response SE df lower.CL upper.CL
              33.1 3.05 50
                              27.5
                                       39.8
##
              24.9 2.29 50 20.7 29.9
##
              20.3 1.87 50 16.9 24.4
##
  H
##
## Results are averaged over the levels of: wool
## Confidence level used: 0.95
## Intervals are back-transformed from the log scale
##
## $contrasts
   contrast ratio
                  SE df lower.CL upper.CL
   L / M 1.33 0.173 50
                          1.026
##
                                     1.73
   L / H 1.63 0.212 50 1.256 2.12
##
## M / H 1.22 0.159 50 0.942
                                  1.59
##
## Results are averaged over the levels of: wool
```

Confidence level used: 0.95

```
em_wool <- emmeans(m, pairwise ~ wool)</pre>
ci wool <- confint(em wool, type = "response", adjust = "none")
ci wool
## $emmeans
   wool response SE df lower.CL upper.CL
            27.6 2.07 50 23.7
                                     32.1
##
  B 23.7 1.78 50 20.4 27.6
##
##
## Results are averaged over the levels of: tension
## Confidence level used: 0.95
## Intervals are back-transformed from the log scale
##
## $contrasts
## contrast ratio SE df lower.CL upper.CL
  A / B 1.16 0.124 50 0.94 1.44
##
## Results are averaged over the levels of: tension
## Confidence level used: 0.95
## Intervals are back-transformed from the log scale
```

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```
em_tension_by_wool <- emmeans(m, pairwise ~ tension | wool)</pre>
ci_tension_by_wool <- confint(em_tension_by_wool, type = "response", adjust = "none")</pre>
ci_tension_by_wool$emmeans
## wool = A:
  tension response SE df lower.CL upper.CL
## L
             35.7 3.80 50 28.9 44.2
## M
         26.8 2.85 50 21.7 33.2
         21.9 2.33 50 17.7 27.1
## H
##
## wool = B:
   tension response SE df lower.CL upper.CL
##
          30.7 3.26 50
                             24.8
                                     38.0
         23.0 2.45 50 18.6 28.5
## M
## H
           18.8 2.00 50 15.2 23.3
##
## Confidence level used: 0.95
## Intervals are back-transformed from the log scale
```

```
ci_tension_by_wool$contrasts
## wool = A:
   contrast ratio SE df lower.CL upper.CL
   L / M 1.33 0.173 50
                        1.026
                                  1.73
   L / H 1.63 0.212 50 1.256 2.12
  M / H 1.22 0.159 50 0.942 1.59
##
## wool = B:
   contrast ratio SE df lower.CL upper.CL
   L / M 1.33 0.173 50
                        1.026
                                1.73
  L / H 1.63 0.212 50 1.256 2.12
   M / H 1.22 0.159 50 0.942
                                  1.59
##
## Confidence level used: 0.95
## Intervals are back-transformed from the log scale
```

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warpbreaks

```
em_wool_by_tension <- emmeans(m, pairwise ~ wool | tension)</pre>
ci_wool_by_tension <- confint(em_wool_by_tension, type = "response", adjust = "none")</pre>
ci_wool_by_tension$emmeans
## tension = I.:
   wool response SE df lower.CL upper.CL
           35.7 3.80 50
                           28.9
                                   44.2
##
##
  B 30.7 3.26 50 24.8 38.0
##
  tension = M:
   wool response SE df lower.CL upper.CL
##
   A 26.8 2.85 50 21.7
                                   33.2
##
         23.0 2.45 50 18.6 28.5
##
  tension = H:
##
   wool response SE df lower.CL upper.CL
           21.9 2.33 50 17.7
                                   27.1
##
##
  B 18.8 2.00 50 15.2 23.3
##
## Confidence level used: 0.95
## Intervals are back-transformed from the log scale
```

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```
ci_wool_by_tension$contrasts
## tension = L:
  contrast ratio SE df lower.CL upper.CL
  A / B 1.16 0.124 50 0.94 1.44
##
## tension = M:
  contrast ratio SE df lower.CL upper.CL
  A / B 1.16 0.124 50 0.94 1.44
##
## tension = H:
  contrast ratio SE df lower.CL upper.CL
   A / B 1.16 0.124 50 0.94 1.44
##
## Confidence level used: 0.95
## Intervals are back-transformed from the log scale
```

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warpbreaks

The estimated median number of breaks is

```
pm <- ci_tension_by_wool$emmeans %>%
 as.data.frame() %>%
 mutate(mean_with_ci = paste0(
   round(response), " (", round(lower.CL), ", ", round(upper.CL), ")")
 ) %>%
 tidyr::pivot_wider(id_cols = tension, names_from = wool, values_from = mean_with_ci)
pm
  # A tibble: 3 \times 3
   tension A
   <fct> <chr> <chr>
##
      36 (29, 44) 31 (25, 38)
## 2 M 27 (22, 33) 23 (19, 29)
## 3 H 22 (18, 27) 19 (15, 23)
```

While holding wool type constant, the percent change in median number of breaks is

```
ci_tension$contrasts %>% as.data.frame %>%
 mutate(
   change = 100*(ratio-1),
   lower = 100*(lower.CL-1),
   upper = 100*(upper.CL-1).
   change_with_ci = paste0(
     round(change), " (", round(lower), ", ", round(upper), ")"
   )) %>%
 select(contrast, change_with_ci)
##
    contrast change with ci
## 1
    L / M 33 (3, 73)
## 2 L / H 63 (26, 112)
## 3 M / H 22 (-6, 59)
```

Poisson regression

Poisson regression with

$$Y_i \stackrel{ind}{\sim} Po(\lambda_i), \quad \log(\lambda_i) = \beta_0 + \beta_1 X_{i,1} + \dots + \beta_p X_{i,p}$$

where, for observation i,

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

Interpretation

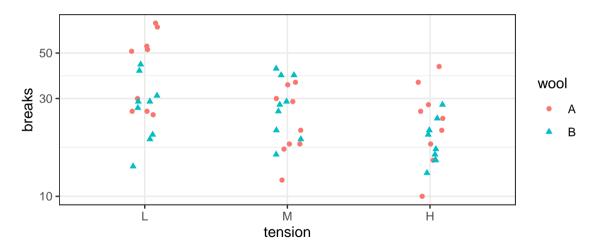
- ullet e^{eta_0} is mean of the dependent variable when all independent variables (X's) are 0
- $100(e^{\beta_p}-1)$ for $p \neq 0$, is the percent increase in the mean of the dependent variable for each unit increase in the associated independent variable.

```
##
## Call:
## glm(formula = breaks ~ tension, family = poisson, data = warpbreaks %>%
     filter(wool == "A"))
##
##
## Deviance Residuals:
##
      Min
                1Ω
                    Median
                                 30
                                         Max
## -3.3383 -1.7940 -0.1125 1.2736
                                      3.5153
##
## Coefficients:
##
              Estimate Std. Error z value Pr(>|z|)
## (Intercept) 3.79674 0.04994 76.030 < 2e-16 ***
## tensionM -0.61868 0.08440 -7.330 2.30e-13 ***
## tensionH -0.59580 0.08378 -7.112 1.15e-12 ***
## ---
## Signif. codes: 0 '***' 0.001 '**' 0.05 '.' 0.1 ' ' 1
##
   (Dispersion parameter for poisson family taken to be 1)
##
##
      Null deviance: 194.97 on 26 degrees of freedom
## Residual deviance: 119.62 on 24 degrees of freedom
## ATC: 264 99
```

```
em <- emmeans(m, pairwise ~ tension)</pre>
ci <- confint(em, type = "response", adjust = "none")</pre>
ci
## $emmeans
  tension rate SE df asymp.LCL asymp.UCL
          44.6 2.22 Inf 40.4
##
                                    49.1
          24.0 1.63 Inf 21.0 27.4
## M
          24.6 1.65 Inf 21.5 28.0
## H
##
## Confidence level used: 0.95
## Intervals are back-transformed from the log scale
##
## $contrasts
   contrast ratio SE df asymp.LCL asymp.UCL
   L / M 1.856 0.1567 Inf
                              1.57 2.19
##
   L / H 1.814 0.1520 Inf 1.54 2.14
##
##
   M / H 0.977 0.0935 Inf 0.81 1.18
##
## Confidence level used: 0.95
## Intervals are back-transformed from the log scale
```

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- For wool type A, the mean number of breaks when
 - tension is low is 45 (40,49),
 - tension is medium is 24 (21,27),
 - tension is high is 25 (22,28),
- For wool type A, tension
 - low compared to medium causes an 86% (57,119) increase in mean number of breaks,
 - low compared to high causes an 81% (54,114) increase in mean number of breaks, and
 - medium compared to high causes an -2% (-19,18) increase in mean number of breaks.



```
##
## Call:
## glm(formula = breaks ~ tension + wool, family = poisson, data = warpbreaks)
##
## Deviance Residuals:
##
      Min
               10 Median
                                 30
                                        Max
## -3.6871 -1.6503 -0.4269 1.1902
                                     4.2616
##
## Coefficients:
              Estimate Std. Error z value Pr(>|z|)
##
## (Intercept) 3.69196 0.04541 81.302 < 2e-16 ***
## tensionM -0.32132 0.06027 -5.332 9.73e-08 ***
## tensionH -0.51849 0.06396 -8.107 5.21e-16 ***
## woolB
              -0.20599 0.05157 -3.994 6.49e-05 ***
## ---
## Signif. codes: 0 '***' 0.001 '**' 0.05 '.' 0.1 ' ' 1
##
  (Dispersion parameter for poisson family taken to be 1)
##
      Null deviance: 297.37 on 53 degrees of freedom
##
## Residual deviance: 210.39 on 50 degrees of freedom
## ATC: 493 06
```

```
em_tension <- emmeans(m, pairwise ~ tension)</pre>
ci tension <- confint(em tension, type = "response", adjust = "none")
ci tension
## $emmeans
   tension rate SE df asymp.LCL asymp.UCL
           36.2 1.42 Inf
                            33.5
##
                                     39.1
          26.2 1.21 Inf
                            24.0
                                    28.7
##
  M
          21.6 1.09 Inf 19.5
## H
                                     23.8
##
## Results are averaged over the levels of: wool
## Confidence level used: 0.95
## Intervals are back-transformed from the log scale
##
## $contrasts
   contrast ratio
                     SE df asymp.LCL asymp.UCL
  L / M 1.38 0.0831 Inf
                               1.23
                                         1.55
##
   L / H 1.68 0.1074 Inf
                               1.48 1.90
##
## M / H 1.22 0.0832 Inf
                               1.07
                                        1.39
##
## Results are averaged over the levels of: wool
```

Confidence level used: 0.95

```
em_wool <- emmeans(m, pairwise ~ wool)</pre>
ci wool <- confint(em wool, type = "response", adjust = "none")
ci wool
## $emmeans
   wool rate SE df asymp.LCL asymp.UCL
        30.3 1.061 Inf
                           28.3 32.5
##
## B 24.7 0.955 Inf 22.9 26.6
##
## Results are averaged over the levels of: tension
## Confidence level used: 0.95
## Intervals are back-transformed from the log scale
##
## $contrasts
## contrast ratio SE df asymp.LCL asymp.UCL
  A / B 1.23 0.0634 Inf 1.11
                                          1.36
##
## Results are averaged over the levels of: tension
## Confidence level used: 0.95
## Intervals are back-transformed from the log scale
```

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```
em_tension_by_wool <- emmeans(m, pairwise ~ tension | wool)</pre>
ci_tension_by_wool <- confint(em_tension_by_wool, type = "response", adjust = "none")</pre>
ci_tension_by_wool$emmeans
## wool = A:
  tension rate SE df asymp.LCL asymp.UCL
          40.1 1.82 Inf 36.7 43.9
## M
          29.1 1.50 Inf 26.3 32.2
          23.9 1.33 Inf 21.4 26.6
## H
##
## wool = B:
   tension rate SE df asymp.LCL asymp.UCL
##
          32.7 1.58 Inf
                           29.7
                                   35.9
          23.7 1.28 Inf 21.3 26.3
## M
## H
          19.4 1.13 Inf 17.4 21.8
##
## Confidence level used: 0.95
## Intervals are back-transformed from the log scale
```

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```
ci_tension_by_wool$contrasts
## wool = A:
   contrast ratio SE df asymp.LCL asymp.UCL
  L / M 1.38 0.0831 Inf
                            1.23
                                    1.55
  L / H 1.68 0.1074 Inf 1.48 1.90
## M / H 1.22 0.0832 Inf 1.07 1.39
##
## wool = B:
   contrast ratio SE df asymp.LCL asymp.UCL
  L / M 1.38 0.0831 Inf
                            1.23
                                    1.55
  L / H 1.68 0.1074 Inf 1.48 1.90
  M / H 1.22 0.0832 Inf 1.07 1.39
##
## Confidence level used: 0.95
## Intervals are back-transformed from the log scale
```

```
em_wool_by_tension <- emmeans(m, pairwise " wool | tension)</pre>
ci_wool_by_tension <- confint(em_wool_by_tension, type = "response", adjust = "none")</pre>
ci_wool_by_tension$emmeans
## tension = I.:
   wool rate SE df asymp.LCL asymp.UCL
        40.1 1.82 Inf
                          36.7
##
                                   43.9
##
        32.7 1.58 Inf 29.7
                                   35.9
   В
##
  tension = M:
   wool rate SE df asymp.LCL asymp.UCL
##
        29.1 1.50 Inf
                          26.3
                                   32.2
##
   В
        23.7 1.28 Inf 21.3
                                   26.3
##
  tension = H:
   wool rate SE df asymp.LCL asymp.UCL
        23.9 1.33 Inf
                          21.4
                                   26.6
##
##
   В
        19.4 1.13 Inf 17.4
                                   21.8
##
## Confidence level used: 0.95
## Intervals are back-transformed from the log scale
```

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```
ci_wool_by_tension$contrasts
## tension = L:
  contrast ratio SE df asymp.LCL asymp.UCL
  A / B 1.23 0.0634 Inf 1.11 1.36
##
## tension = M:
  contrast ratio SE df asymp.LCL asymp.UCL
  A / B 1.23 0.0634 Inf 1.11 1.36
##
## tension = H:
## contrast ratio SE df asymp.LCL asymp.UCL
   A / B 1.23 0.0634 Inf 1.11
                                      1.36
##
## Confidence level used: 0.95
## Intervals are back-transformed from the log scale
```

The estimated mean number of breaks is

```
pm <- ci_tension_by_wool$emmeans %>%
 as.data.frame() %>%
 mutate(mean_with_ci = paste0(
   round(rate), " (", round(asymp.LCL), ", ", round(asymp.UCL), ")")
 ) %>%
 tidyr::pivot_wider(id_cols = tension, names_from = wool, values_from = mean_with_ci)
pm
## # A tibble: 3 x 3
##
   tension A
                       В
   <fct> <chr> <chr>
## 1 L.
      40 (37, 44) 33 (30, 36)
## 2 M 29 (26, 32) 24 (21, 26)
## 3 H 24 (21, 27) 19 (17, 22)
```

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While holding wool type constant, the percent change in mean number of breaks is

```
ci_tension$contrasts %>% as.data.frame %>%
 mutate(
   change = 100*(ratio-1),
   lower = 100*(asymp.LCL-1),
   upper = 100*(asymp.UCL-1),
   change_with_ci = paste0(
     round(change), " (", round(lower), ", ", round(upper), ")"
   )) %>%
 select(contrast, change_with_ci)
##
    contrast change with ci
## 1 L / M 38 (23, 55)
## 2 L / H 68 (48, 90)
## 3 M / H 22 (7, 39)
```

Summary

With count data (with no upper limit), a Poisson regression model is appropriate.

