Count data and the Poisson distribution

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Counts as extensions of binary data

- · Counts can be thought of as repeated binary trials
- $\sum y_i$ where y is equal to 1 or 0 provides a count
- Generally, we could treat sum(y==1) + sum(y==0) or nrow(y) as the exposure, or denominator for a rate. Why?

An example of count data

```
load("./data/fieldplayer_overall_season_stats.rda")
load("./data/player_rda")

nwsl_stats<-fieldplayer_overall_season_stats
nwsl_players<-player

head(nwsl_players)</pre>
```

```
## # A tibble: 6 x 5
##
    person_id player
                              nation pos
                                          name_other
                              <chr> <chr> <chr> <chr>
##
        <dhl> <chr>>
## 1
          342 Marisa Abegg
                              USA
                                     DF
                                           <NA>
          117 Danesha Adams
                              USA FW,MF <NA>
## 2
## 3
            6 Adriana
                               FSP
                                     FW
                                            <NA>
          300 Leigh Ann Brown USA
                                      DF,MF <NA>
## 4
## 5
          202 Jazmyne Avant
                              USA
                                     DF
                                          <NA>
## 6
           28 Amv Barczuk
                              IISΔ
                                     DF
                                         < N A >
```

make a joined table with players names

```
### attaching names
dat<-nwsl stats %>%
  left join(nwsl players %>%
              select(person id, player))
### check to ensure that the dimensions are what we want
nrow(dat) == nrow(nwsl stats)
## [1] TRUE
### if we have multiple positions for players
dat<-nwsl stats %>%
  left join(nwsl players)
```

Approaches to modeling count data

The Poisson model

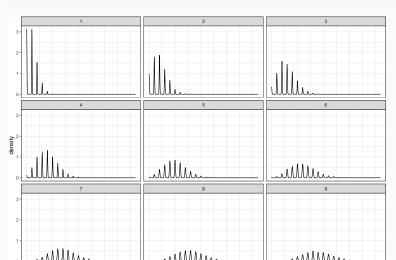
Where y is a non-negative integer (count)

$$y \sim Poisson(\lambda)$$
 $E(y) = \bar{y} = \lambda$
 $Var(y) = \lambda$
 $Pr(y = k) = \frac{\lambda^k e^{-\lambda}}{k!}$

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Shape of the Poisson distribution

```
ggplot(pois_demo, aes(x=count)) +
  geom_density(adjust = 1/4) +
  facet_wrap(~lambda)
```



Let's look at each Poisson variable

```
## # A tibble: 9 x 3
##
    lambda mean variance
## *
     <int> <dbl>
                   <dbl>
## 1
         1 1.01 1.01
         2 2.00
## 2
                 1.99
## 3
         3 2.97
                    2.94
         4 4.00
## 4
                    3.99
## 5
         5 5.02
                    5.04
## 6
         6 6.02
                    6.27
## 7
         7 7.00
                    6.93
         8 7.98
                    8.11
## 8
## 9
         9 8.98
                    9.20
```

Poisson models as a GLM

For a count variable y, we can specify a Poisson GLM with a log link function

$$y \sim Poisson(\lambda)$$
 $\lambda = e^{eta X} = e^{eta_0 + eta_1 x_1 \cdots eta_n x_n}$ $E(y|x) = e^{\lambda}$ $\log(E(y|x)) = \lambda = eta X$

```
m0 mp < -glm(mp ~
             1.
           data = dat,
           family = "poisson")
### check to see if the data looks similar to the poisson
data sim<-data.frame(</pre>
  y_{pred} = rpois(nrow(dat), 13.54815),
 mp = dat mp
## evaluate the distributions
ggplot(data_sim,
       aes(x = y_pred)) +
  geom histogram(fill = "red", alpha = 0.5) +
  geom_histogram(aes(x = mp), fill = "blue", alpha = 0.5)
```

Modeling assists

```
assist 0<-glm(ast ~ 1,
              data = dat,
              family = "poisson")
sim_dat<-data.frame(</pre>
  v pred = rpois(nrow(dat), 0.9782119),
  ast = dat$ast
ggplot(sim_dat,
       aes(x = v pred)) +
  geom_histogram(fill = "red", alpha =0.5) +
  geom_histogram(aes(x = ast), fill = "blue", alpha = 0.5)
```



let's generate predictions for players

```
### take our two predictors
### define reasonable ranges to predict over
mp<-1:24
gls<-0:5
### generate expected values
fake_data<- expand_grid(mp, gls)</pre>
fake data<-fake data%>%
  mutate(expected_assists =
           predict(assist 3, fake data,
                   type = "response"))
ggplot(fake_data,
       aes(x = mp, y = expected assists,
           color = factor(gls))) +
  geom_line()
```

Advantages of the Poisson distribution for regression

- 1. Constrained to non-negative integers
- 2. Variance scales with the expectation of y
- 3. Relatively simple to interpret

However:

$$\lambda = E(y|x) = var(y)$$

Homework

- Visualize the distribution of goals across players for the 2019 season (your choice on geom)
- 2. Define a linear predictor for goals made during a season, where the players' position is the only predictor.
- 3. Estimate this model with a Normal likelihood (OLS)
- 4. Estimate this model with a Poisson likelihood (family = "poisson")
- 5. Generate predictions for each position for both models
- 6. Compare the predictions. Which model do you prefer? Why?