Count outcomes, Poisson GLMs

Brian Caffo, Jeffrey Leek, Roger Peng

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Key ideas

- Many data take the form of counts
- Calls to a call center
- Number of flu cases in an area
- Number of cars that cross a bridge
- Data may also be in the form of rates
- Percent of children passing a test
- Percent of hits to a website from a country
- ► Linear regression with transformation is an option

Poisson distribution

- ▶ The Poisson distribution is a useful model for counts and rates
- Here a rate is count per some monitoring time
- Some examples uses of the Poisson distribution
 - Modeling web traffic hits
 - Incidence rates
 - Approximating binomial probabilities with small p and large n
 - Analyzing contigency table data

The Poisson mass function

▶ $X \sim Poisson(t\lambda)$ if

$$P(X = x) = \frac{(t\lambda)^x e^{-t\lambda}}{x!}$$

For x = 0, 1, ...

- ▶ The mean of the Poisson is $E[X] = t\lambda$, thus $E[X/t] = \lambda$
- ▶ The variance of the Poisson is $Var(X) = t\lambda$.
- ▶ The Poisson tends to a normal as $t\lambda$ gets large.

```
par(mfrow = c(1, 3))
plot(0 : 10, dpois(0 : 10, lambda = 2), type = "h", frame = plot(0 : 20, dpois(0 : 20, lambda = 10), type = "h", frame plot(0 : 200, dpois(0 : 200, lambda = 100), type = "h", frame plot(0 : 200, dpois(0 : 200, lambda = 100), type = "h", frame plot(0 : 200, dpois(0 : 200, lambda = 100), type = "h", frame plot(0 : 200, dpois(0 : 200, lambda = 100), type = "h", frame plot(0 : 200, dpois(0 : 200, lambda = 100), type = "h", frame plot(0 : 200, dpois(0 : 200, lambda = 100), type = "h", frame plot(0 : 200, dpois(0 : 200, lambda = 100), type = "h", frame plot(0 : 200, dpois(0 : 200, lambda = 100), type = "h", frame plot(0 : 200, dpois(0 : 200, lambda = 100), type = "h", frame plot(0 : 200, dpois(0 : 200, lambda = 100), type = "h", frame plot(0 : 200, dpois(0 : 200, lambda = 100), type = "h", frame plot(0 : 200, dpois(0 : 200, lambda = 100), type = "h", frame plot(0 : 200, dpois(0 : 200, lambda = 100), type = "h", frame plot(0 : 200, dpois(0 : 200, lambda = 100), type = "h", frame plot(0 : 200, dpois(0 : 200, lambda = 100), type = "h", frame plot(0 : 200, dpois(0 : 200, lambda = 100), type = "h", frame plot(0 : 200, dpois(0 : 200, lambda = 100), type = "h", frame plot(0 : 200, dpois(0 : 200, lambda = 100), type = "h", frame plot(0 : 200, lambda = 100), type = "h", frame plot(0 : 200, lambda = 100), type = "h", frame plot(0 : 200, lambda = 100), type = "h", frame plot(0 : 200, lambda = 100), type = "h", frame plot(0 : 200, lambda = 100), type = "h", frame plot(0 : 200, lambda = 100), type = "h", frame plot(0 : 200, lambda = 100), type = "h", frame plot(0 : 200, lambda = 100), type = "h", frame plot(0 : 200, lambda = 100), type = "h", frame plot(0 : 200, lambda = 100), type = "h", frame plot(0 : 200, lambda = 100), type = "h", frame plot(0 : 200, lambda = 100), type = "h", frame plot(0 : 200, lambda = 100), type = "h", frame plot(0 : 200, lambda = 100), type = "h", frame plot(0 : 200, lambda = 100), type = "h", frame plot(0 : 200, lambda = 100), type = "h", frame plot(0 : 200, lambda = 100), type
```



Poisson distribution

Sort of, showing that the mean and variance are equal

```
x <- 0 : 10000; lambda = 3
mu <- sum(x * dpois(x, lambda = lambda))
sigmasq <- sum((x - mu)^2 * dpois(x, lambda = lambda))
c(mu, sigmasq)</pre>
```

```
## [1] 3 3
```

Example: Leek Group Website Traffic

Consider the daily counts to Jeff Leek's web site

http://biostat.jhsph.edu/~jleek/

Since the unit of time is always one day, set t=1 and then the Poisson mean is interpretted as web hits per day. (If we set t=24, it would be web hits per hour).

Website data

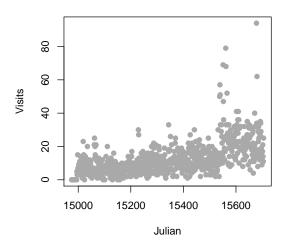
```
download.file("https://dl.dropboxusercontent.com/u/7710864,
load("./data/gaData.rda")
gaData$julian <- julian(gaData$date)
head(gaData)</pre>
```

```
## date visits simplystats julian
## 1 2011-01-01 0 0 14975
## 2 2011-01-02 0 0 14976
## 3 2011-01-03 0 0 14977
## 4 2011-01-04 0 0 14978
## 5 2011-01-05 0 0 14979
## 6 2011-01-06 0 0 14980
```

http://skardhamar.github.com/rga/

Plot data

plot(gaData\$julian,gaData\$visits,pch=19,col="darkgrey",xlat



Linear regression

$$NH_i = b_0 + b_1 JD_i + e_i$$

 NH_i - number of hits to the website

 JD_i - day of the year (Julian day)

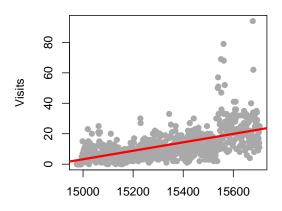
 b_0 - number of hits on Julian day 0 (1970-01-01)

 b_1 - increase in number of hits per unit day

 e_i - variation due to everything we didn't measure

Linear regression line

```
plot(gaData$julian,gaData$visits,pch=19,col="darkgrey",xlab
lm1 <- lm(gaData$visits ~ gaData$julian)
abline(lm1,col="red",lwd=3)</pre>
```



Aside, taking the log of the outcome

- ► Taking the natural log of the outcome has a specific interpretation.
- Consider the model

$$\log(NH_i) = b_0 + b_1 JD_i + e_i$$

 NH_i - number of hits to the website

 JD_i - day of the year (Julian day)

 b_0 - log number of hits on Julian day 0 (1970-01-01)

 b_1 - increase in log number of hits per unit day

 e_i - variation due to everything we didn't measure

Exponentiating coefficients

- $e^{E[\log(Y)]}$ geometric mean of Y.
 - With no covariates, this is estimated by $e^{\frac{1}{n}\sum_{i=1}^{n}\log(y_i)}=(\prod_{i=1}^{n}y_i)^{1/n}$
- When you take the natural log of outcomes and fit a regression model, your exponentiated coefficients estimate things about geometric means.
- $ightharpoonup e^{eta_0}$ estimated geometric mean hits on day 0
- e^{β_1} estimated relative increase or decrease in geometric mean hits per day
- ► There's a problem with logs with you have zero counts, adding a constant works

```
round(exp(coef(lm(I(log(gaData$visits + 1)) ~ gaData$julian
```

```
## (Intercept) gaData$julian
## 0.00000 1.00231
```

Linear vs. Poisson regression

Linear

$$NH_i = b_0 + b_1 JD_i + e_i$$

or

$$E[NH_i|JD_i, b_0, b_1] = b_0 + b_1JD_i$$

Poisson/log-linear

$$\log (E[NH_i|JD_i, b_0, b_1]) = b_0 + b_1 JD_i$$

or

$$E[NH_i|JD_i, b_0, b_1] = \exp(b_0 + b_1JD_i)$$

Multiplicative differences

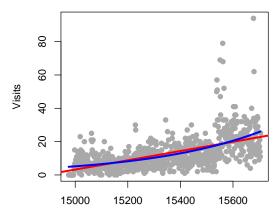
$$E[NH_i|JD_i, b_0, b_1] = \exp(b_0 + b_1JD_i)$$

$$E[NH_i|JD_i,b_0,b_1] = \exp(b_0)\exp(b_1JD_i)$$

If JD_i is increased by one unit, $E[NH_i|JD_i,b_0,b_1]$ is multiplied by $\exp{(b_1)}$

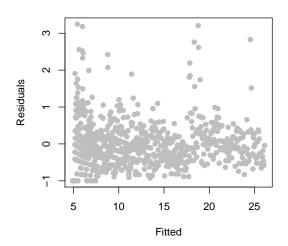
Poisson regression in R

```
plot(gaData$julian,gaData$visits,pch=19,col="darkgrey",xlal
glm1 <- glm(gaData$visits ~ gaData$julian,family="poisson")
abline(lm1,col="red",lwd=3); lines(gaData$julian,glm1$fitte</pre>
```



Mean-variance relationship?

plot(glm1\$fitted,glm1\$residuals,pch=19,col="grey",ylab="Residuals")



Model agnostic standard errors

```
library(sandwich)
confint.agnostic <- function (object, parm, level = 0.95,
    cf <- coef(object); pnames <- names(cf)</pre>
    if (missing(parm))
        parm <- pnames
    else if (is.numeric(parm))
        parm <- pnames[parm]</pre>
    a \leftarrow (1 - level)/2; a \leftarrow c(a, 1 - a)
    pct <- stats:::format.perc(a, 3)</pre>
    fac <- qnorm(a)
    ci <- array(NA, dim = c(length(parm), 2L), dimnames = 1
    ses <- sqrt(diag(sandwich::vcovHC(object)))[parm]</pre>
    ci[] <- cf[parm] + ses %0% fac
    сi
```

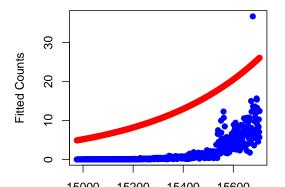
Estimating confidence intervals

```
confint(glm1)
## Waiting for profiling to be done...
                       2.5 % 97.5 %
##
## (Intercept) -34.346577587 -31.159715656
## gaData$julian 0.002190043 0.002396461
confint.agnostic(glm1)
##
                       2.5 % 97.5 %
  (Intercept) -36.362674594 -29.136997254
## gaData$julian 0.002058147 0.002527955
```

Rates

$$E[NHSS_i|JD_i, b_0, b_1]/NH_i = \exp(b_0 + b_1JD_i)$$
 $\log(E[NHSS_i|JD_i, b_0, b_1]) - \log(NH_i) = b_0 + b_1JD_i$
 $\log(E[NHSS_i|JD_i, b_0, b_1]) = \log(NH_i) + b_0 + b_1JD_i$

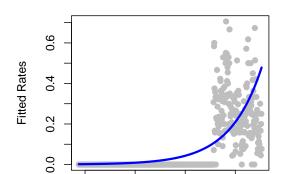
Fitting rates in R



Fitting rates in R

```
family="poisson",data=gaData)
plot(julian(gaData$date),gaData$simplystats/(gaData$visits-
ylab="Fitted Rates",pch=19)
lines(julian(gaData$date),glm2$fitted/(gaData$visits+1),col
```

glm2 <- glm(gaData\$simplystats ~ julian(gaData\$date),offset</pre>



More information

- Log-linear models and multiway tables
- Wikipedia on Poisson regression, Wikipedia on overdispersion
- ▶ Regression models for count data in R
- pscl package the function zeroinfl fits zero inflated models.