Estimating a Poisson Regression Model with R

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The R Programming Language

- R: Popular open-source programming language for statistical analysis.
- Widely used in statistics and econometrics.
- User-friendly and powerful IDE for R: RStudio.
- ▶ Basic functionalities of R can be extended by **packages**.
- Large number of packages available on the Comprehensive R Archive Network.
- ► Goal of this presentation: Illustrate how to use R for the estimation of a Poisson regression model.

Count Data Models

- **Count data** models are used to explain dependent variables that are natural numbers, i.e., positive integers such that $y_i \in \mathbb{N}$, where $\mathbb{N} = \{0, 1, 2, \ldots\}$.
- Count data models are frequently used in economics to study countable events: Number of years of education, number of patent applications filed by companies, number of doctor visits, number of crimes committed in a given city, etc.
- ▶ The **Poisson model** is a popular count data model.

Poisson Regression Model

▶ Given a parameter $\lambda_i > 0$, the **Poisson model** assumes that the probability of observing $Y_i = y_i$, where $y_i \in \mathbb{N}$, is equal to:

$$Prob(Y_i = y_i \mid \lambda_i) = \frac{\lambda_i^{y_i} \exp\{-\lambda_i\}}{y_i!},$$

for $i = 1, \ldots, N$.

▶ The mean and the variance of Y_i are equal to the parameter λ_i :

$$E(Y_i \mid \lambda_i) = V(Y_i \mid \lambda_i) = \lambda_i,$$

implying equi-dispersion of the data.

▶ To control for **observed characteristics**, the parameter λ_i can be parametrized as follows (implying $\lambda_i > 0$):

$$E(Y_i|X_i,\beta) \equiv \lambda_i = \exp\{X_i'\beta\},$$

where X_i is a vector containing the covariates.

Simulating Data

▶ R function simulating data from Poisson regression model:

```
simul_poisson <- function(n, beta) {
  k <- length(beta)  # number of covariates
  x <- replicate(k - 1, rnorm(n)) # simulate covariates
  x <- cbind(1, x)  # for intercept term
  lambda <- exp(x %*% beta)  # individual means
  y <- rpois(n, lambda)  # simulate count
  return(data.frame(y, x))  # return variables
}</pre>
```

Using function to generate data:

```
set.seed(123)
nobs <- 1000
beta <- c(-.5, .4, -.7)
data <- simul_poisson(nobs, beta)</pre>
```

Data Description

Descriptive statistics:

```
# extract variables of interest from data set
y <- data[, 1]
x <- as.matrix(data[, 2:4])
# descriptive statistics
library(psych)
describe(data)
```

```
## y 1 1000 0.76 1.08 0.00 0.54 0.00 0.00 8.00
## X2 3 1000 0.02 0.99 0.01 0.01 0.96 -2.81 3.24 0
## X3 4 1000 0.04 1.01 0.05 0.05 1.05 -3.05 3.39
##
```

n mean sd median trimmed mad min max ra

y 0.03 ## X1 0.00

vars

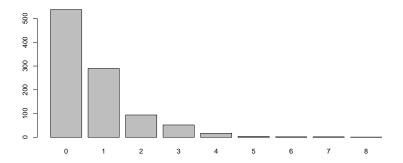
se

##

Data Description

► Histogram of count variable:

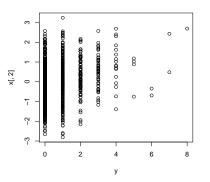
barplot(table(y))

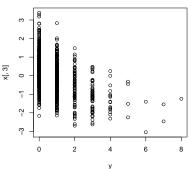


Data Description

▶ Relationship between count variable and covariates:

```
par(mfrow = c(1, 2))
plot(y, x[, 2])
plot(y, x[, 3])
```





Likelihood Function and ML Estimator

▶ Individual contribution to the likelihood function:

$$L_i(\beta; y_i, x_i) = \frac{\exp\{y_i x_i \beta\} \exp\{-\exp\{x_i \beta\}\}}{y_i!}$$

► Individual log-Likelihood function:

$$\ell_i(\beta; y_i, x_i) = \log L_i(\beta; y_i, x_i) = y_i x_i \beta - \exp\{x_i \beta\} - \log(y_i!)$$

► Maximum Likelihood Estimator:

$$\hat{\beta}_{\mathsf{MLE}} = \arg\max_{\beta} \sum_{i=1}^{N} \ell(\beta; y, X)$$

▶ Optimization (using *minimization* of objective function):

$$\hat{eta}_{\mathsf{MLE}} = \arg\min_{eta} \, Q(eta; y, X) \qquad Q(eta; y, X) = -rac{1}{N} \sum_{i=1}^{N} \ell_i(eta; y_i, x_i)$$

Coding the Objective Function

```
# Objective function of Poisson regression model
obj poisson <- function(beta, y, x) {
  lambda \leftarrow x \% *\% beta
  llik <- y*lambda - exp(lambda) - lfactorial(y)</pre>
  return(-mean(llik))
# Evaluating objective function
beta0 <- c(1, 2, 3)
obj_poisson(beta0, y, x)
```

```
## [1] 1757.113
```

Maximizing the Objective Function

Set starting values:

```
beta0 <- rep(0, length(beta))</pre>
```

Optimize using quasi-Newton method (BFGS algorithm):

Show results:

```
## ML estimates: -0.5740286 0.3921569 -0.7231029 ## Objective function: 0.9998689
```

Comparing Results to Built-in Function

```
opt_glm <- glm(y \sim 0 + x, family = poisson)
summary(opt_glm)
##
## Call:
## glm(formula = y \sim 0 + x, family = poisson)
##
## Deviance Residuals:
##
      Min 1Q Median 3Q
                                       Max
## -2.1707 -0.9185 -0.5915 0.5133 3.6768
##
## Coefficients:
## Estimate Std. Error z value Pr(>|z|)
## xX1 -0.57403  0.04602 -12.47  <2e-16 ***
## xX2 0.39216 0.03690 10.63 <2e-16 ***
## xX3 -0.72310 0.03662 -19.74 <2e-16 ***
```

Signif codes: 0 '***' 0 001 '**' 0 05 ' ' 0 '

Comparing Results to Built-in Function

Collect results from the two approaches to compare them:

```
## True MLE GLM
## constant -0.5 -0.5740286 -0.5740289
## x1 0.4 0.3921569 0.3921561
## x2 -0.7 -0.7231029 -0.7231039
```

Question: Our results (MLE) are virtually the same as those obtained with the built-in function GLM, but not identical. Where do the small differences come from?

Empirical Illustration

- Goal: Investigate the determinants of fertility.
- Poisson regression model used to estimate the relationship between explanatory variables and count outcome variable.
- ▶ Both our estimator coded from scratch and R built-in function will be used.

Data

- Source: Botswana's 1988 Demographic and Health Survey.
- ▶ Data set borrowed from Wooldridge:

```
library(wooldridge)
data(fertil2)
```

Outcome variable: Total number of living children:

```
y_lab <- "children"
```

► Explanatory variables: Education, age, marital status, living in urban area, having electricity/TV at home:

Loading data

Selecting variables and removing missing values:

```
data <- fertil2[, c(y_lab, x_lab)]
data <- na.omit(data)</pre>
```

▶ Show first 6 observations on first 8 variables:

```
head(data[, 1:8], n = 6)
```

```
children educ age agesq evermarr urban electric tv
##
## 1
           0
               12 24 576
           3
               13 32 1024
## 2
           1 5 30 900
## 3
                                                  0
           2 4 42 1764
## 4
           2
               11 43
                                                  1
                      1849
           1
               7
                  36
                      1296
                                                  0
```

Descriptive Statitics

library(psych) describe(data)

```
##
                           sd median trimmed
          vars
                  n
                     mean
## children
             1 4358 2.27
                           2.22
## educ
             2 4358 5.86
                            3.93
## age
             3 4358 27.40
                            8.69
           4 4358 826.42 526.96
## agesq
             5 4358
                     0.48
                            0.50
## evermarr
## urban
             6 4358 0.52
                            0.50
## electric
          7 4358 0.14
                           0.35
             8 4358 0.09
                            0.29
## tv
           skew kurtosis
##
                          se
## children
           1.07
                   0.75 0.03
## educ
          -0.03
                  -0.50 0.06
## age
                  -0.490.13
          0.59
           1.10
## agesq
                  0.52 7.98
## evermarr
           0.09
                   -1.990.01
```

mad 1

2.97

4.45

8.90

0.00

0.00

0.00

0.00

1.95

5.79

26.71

0.47

0.52

0.05

0.00

752.87 467.02 3

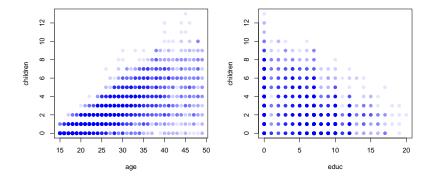
26

676

0

Plot

```
attach(data)
par(mfrow = c(1, 2))
blue_transp <- adjustcolor("blue", alpha.f = 0.1)
plot(age, children, pch = 19, col = blue_transp)
plot(educ, children, pch = 19, col = blue_transp)</pre>
```



MLE of the Poisson Model

Maximum likelihood function using built-in function glm():

Maximum likelihood function using our own function:

MLE of the Poisson Model

- ► Results different from glm()?
- Optimization algorithms are iterative methods that rely on different criteria to dertermine if/when the optimum has been reached.
- ► For example: Change in the objective function, change in the parameter values, change in the gradient, step size, etc.
- [More in Advanced Microeconometrics course].
- ► Try to adjust tuning parameters, for example add control = list(ndeps = rep(1e-8, ncol(x))) to optim() to change step size of gradient approximation.

Summarizing the Empirical Results

summary(mle)

##

##

educ

evermarr

```
## Call:
## glm(formula = children ~ educ + age + agesq + evermarr +
## electric + tv, family = "poisson", data = data)
##
## Deviance Residuals:
## Min 1Q Median 3Q Max
## -3.5620 -0.8116 -0.1091 0.5439 2.8893
##
## Coefficients:
```

Estimate Std. Error z value Pr(>|z|)

0.0244473 12.875 < 2e-16.***

-0.0216645 0.0029131 -7.437 1.03e-13 ***

age 0.3373308 0.0099365 33.949 < 2e-16 ***
agesq -0.0041158 0.0001453 -28.331 < 2e-16 ***

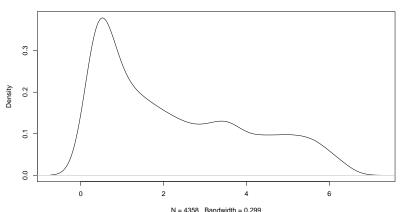
0 3147510

(Intercept) -5.3748294 0.1628671 -33.001 < 2e-16 ***

Fitted Values

```
plot(density(mle$fitted.values),
    main = "Density of fitted mean values")
```





Formatting the results

```
library(xtable)
xtable(mle)
## % latex table generated in R 3.6.0 by xtable 1.8-4 packs
## % Mon May 6 11:06:08 2019
## \begin{table}[ht]
## \centering
## \begin{tabular}{rrrrr}
     \hline
##
    & Estimate & Std. Error & z value & Pr($>$$|$z$|$) \\
##
    \hline
##
   (Intercept) & -5.3748 & 0.1629 & -33.00 & 0.0000 \\
##
     educ & -0.0217 & 0.0029 & -7.44 & 0.0000 \\
##
     age & 0.3373 & 0.0099 & 33.95 & 0.0000 \\
##
```

agesq & -0.0041 & 0.0001 & -28.33 & 0.0000 \\ evermarr & 0.3148 & 0.0244 & 12.87 & 0.0000 \\ ## urban & -0.0861 & 0.0216 & -3.98 & 0.0001 \\ ## ## electric & -0.1205 & 0.0388 & -3.10 & 0.0019 \\