Quiz 3

GLM Theory

library(tidyverse)

```
-- Attaching core tidyverse packages ----- tidyverse 2.0.0 --
v dplyr
           1.1.4
                      v readr
                                  2.1.5
v forcats
            1.0.0
                                  1.5.1
                      v stringr
            3.5.1
                                  3.2.1
v ggplot2
                      v tibble
v lubridate 1.9.3
                      v tidyr
                                  1.3.1
            1.0.2
v purrr
-- Conflicts ----- tidyverse_conflicts() --
x dplyr::filter() masks stats::filter()
x dplyr::lag()
                 masks stats::lag()
i Use the conflicted package (<a href="http://conflicted.r-lib.org/">http://conflicted.r-lib.org/</a>) to force all conflicts to become
```

Question 1

Is it acceptable to use the quasi-poisson model when the data actually follow the Poisson distribution? Briefly explain why or why not.

It is not acceptable to use a quasi-poisson model when the data actually follows a Poisson distribution. In fact, the quasi-poisson exists for the case when the data doesn't satisfy assumptions of Poisson data, specifically the assumption that the mean and variance are equal and quasi-poisson will inflate the standard errors to handle variance greater than we would expect in Poisson data. If our data follows a Poisson distribution, then the mean=variance assumption should be satisfied, and quasi-poisson just takes away from the theoretical strength of our analysis.

The support of the distribution is the integers where y>0 or equivalently $[1,2,...,\infty)$.

We start off with the PMF of the truncated Poisson, which is

$$P(Y = y) = \frac{e^{-\lambda} \lambda^y}{(1 - e^{-\lambda})y!}$$

Note that our support is not dependent on an unknown parameter, meaning we can attempt to write this in exponential family form to see if the truncated Poisson is a member of the exponential family.

We will exponentiate the logarithm of the PMF.

$$\begin{split} \exp(\log\frac{e^{-\lambda}\lambda^y}{(1-e^{-\lambda})y!}) \\ exp(\log(e^{-\lambda}) + \log(\lambda^y) - \log(1-e^{-\lambda}) - \log(y!)) \\ \\ \frac{1}{y!}exp(\log(\frac{e^{-\lambda}}{1-e^{-\lambda}}) + ylog(\lambda)) \end{split}$$

We notice that this belongs to the exponential family.

The exponential family in canonical form can be written as:

$$f(y \mid \lambda) = h(y)exp(\eta(\lambda)T(y) - \psi(\lambda))$$

where

$$h(y) = \frac{1}{y!}$$

$$\psi(\lambda) = -log(\frac{e^{-\lambda}}{1 - e^{-\lambda}})$$

$$\eta = log(\lambda)$$

$$T(y) = y$$

Recall that the log partition function can be derived from the η and ψ components of our exponential family structure, specifically reparameterizing ψ in terms of η .

We found that $\eta = log(\lambda)$ or equivalently that $e^{\eta} = \lambda$.

From $\psi = -log(\frac{e^{-\lambda}}{1-e^{-\lambda}})$, we write the log partition function:

$$A = -log(\frac{e^{-e^{\eta}}}{1 - e^{-e^{\eta}}})$$

By logarithm rules, we can say

$$A = e^{\eta} + \log(1 - e^{-e^{\eta}})$$

From the log partition function, we can we aponize the following fact to find ${\cal E}(Y)$

$$E(Y) = \frac{\partial A}{\partial \eta}$$

$$E(Y) = e^{\eta} + \frac{e^{-e^{\eta}}e^{\eta}}{1 - e^{-e^{\eta}}}$$

This is equivalent to

$$E(Y) = \frac{e^{\eta}}{1 - e^{-e^{\eta}}}$$

Since $e^{\eta} = \lambda$,

we can say

$$E(Y) = \frac{\lambda}{1 - e^{-\lambda}} = \frac{\lambda e^{\lambda}}{e^{\lambda} - 1}$$

The canonical link can be viewed by η which connects our response to the predictors. Since $\eta = log(\lambda), \ X\beta = log(\lambda)$ is the link function.

To find the log likelihood of the truncated Poisson, I will first write out the likelihood by taking the product of PMF from n=1, ..., n.

We define the PMF of the truncated Poisson as

$$f_Y = \frac{e^{-\lambda} \lambda^y}{(1 - e^{-\lambda})y!}$$

The likelihood is then

$$\begin{split} L(\lambda) &= \prod_{i=1}^n \frac{e^{-\lambda} \lambda^{y_i}}{(1-e^{-\lambda}) y_i!} \\ L(\lambda) &= \frac{e^{-n\lambda} \lambda^{\sum_{i=1}^n y_i}}{(1-e^{-\lambda})^n \prod_{i=1}^n y_i!} \end{split}$$

In attempt to find the log likelihood, we will take the logarithm.

$$log L(\lambda) = log(\frac{e^{-n\lambda}\lambda^{\sum_{i=1}^{n}y_i}}{(1 - e^{-\lambda})^n \prod_{i=1}^{n}y_i!})$$

By logarithm properties

$$log~L(\lambda) = -n\lambda + \sum_{i=1}^n y_i log\lambda - nlog(1-e^{-\lambda}) - \sum_{i=1}^n log(y_i!)$$

Using our link $log(\lambda) = X\beta$, we know $\lambda = e^{X\beta}$, so we can equivalently write our log likelihood as

$$logL(\beta) = -ne^{X\beta} + \sum_{i=1}^n y_i X\beta - nlog(1 - e^{-e^{X\beta}}) - \sum_{i=1}^n log(y_i!)$$

This is equivalently

$$=\sum_{i=1}^n[-e^{X\beta}+y_iX\beta-\log(1-e^{-e^{X\beta}})-\log(y_i!)]$$

The vectorized form can be written as

$$logL(\beta) = \sum_{i=1}^n [-e^{X_i^T\beta} + y_iXi^T\beta - log(1-e^{-e^{X_i^T\beta}}) - log(y_i!)]$$

This matches the stated log likelihood.

To find the score function, I will take the partial derivative of the log likelihood with respect to β .

$$\begin{split} S(\beta) &= \frac{\partial \; logL(\beta)}{\partial \beta} = \sum_{i=1}^{n} [-e^{X_{i}^{T}\beta}X_{i} + y_{i}X_{i} - \frac{-e^{-e^{X_{i}^{T}\beta}} \times -e^{X_{i}^{T}\beta}X_{i}}{1 - e^{-e^{X_{i}^{T}\beta}}}] \\ S(\beta) &= \frac{\partial \; logL(\beta)}{\partial \beta} = \sum_{i=1}^{n} [-e^{X_{i}^{T}\beta}X_{i} + y_{i}X_{i} - \frac{e^{-e^{X_{i}^{T}\beta}} \times e^{X_{i}^{T}\beta}X_{i}}{1 - e^{-e^{X_{i}^{T}\beta}}}] \\ S(\beta) &= \frac{\partial \; logL(\beta)}{\partial \beta} = \sum_{i=1}^{n} [y_{i}X_{i} - \frac{e^{-e^{X_{i}^{T}\beta}} \times e^{X_{i}^{T}\beta}X_{i} + e^{X_{i}^{T}}X_{i}(1 - e^{-e^{X_{i}^{T}\beta}})}{1 - e^{-e^{X_{i}^{T}\beta}}}] \end{split}$$

This allows us to simplify our score function to

$$S(\beta) = \sum_{i=1}^n [y_i X_i - \frac{e^{X_i^T \beta}}{1 - e^{-e^{X_i^T \beta}}} X_i]$$

The score function in matrix form is:

$$S(\beta) = X^T (y - \frac{e^{X\beta}}{1 - e^{-e^{X\beta}}})$$

The hessian function is found by taking the second partial of the log likelihood with respect to β . This is also found by taking the partial derivative of the score function with respect to β

$$\begin{split} H(\beta) &= \frac{\partial}{\partial \beta} S(\beta) = \frac{\partial}{\partial \beta} [\sum_{i=1}^n [y_i X_i - \frac{e^{X_i^T \beta}}{1 - e^{-e^{X_i^T \beta}}} X_i]] \\ &= -\sum_{i=1}^n \frac{\partial}{\partial \beta} (\frac{e^{X_i^T \beta}}{1 - e^{-e^{X_i^T \beta}}} X_i) \end{split}$$

We can use quotient rule to find this derivative

$$\frac{\partial}{\partial \beta}(\frac{e^{X_{i}^{T}\beta}}{1-e^{-e^{X_{i}^{T}\beta}}}X_{i}) = \frac{(1-e^{-e^{X_{i}^{T}\beta}})e^{X_{i}^{T}\beta}X_{i}X_{i}^{T} - e^{X_{i}^{T}}e^{-e^{X_{i}^{T}\beta}}e^{X_{i}^{T}\beta}X_{i}}{(1-e^{-e^{X_{i}^{T}\beta}})^{2}}$$

We use this to write the Hessian

$$H(\beta) = -\sum_{i=1}^n \frac{(e^{X_i^T\beta} - e^{-e^{X_i^T\beta}}e^{X_i^T\beta} - e^{2X_i^T\beta}e^{-e^{X_i^T\beta}})}{(1 - e^{-e^{X_i^T\beta}})^2} X_i X_i^T$$

In matrix form, we can write the Hessian as:

$$H(\beta) = -X^T W X$$

where
$$W = \frac{e^{X\beta}-e^{-e^{X\beta}}e^{X\beta}-e^{2X\beta}e^{-e^{X\beta}}}{(1-e^{-e^{X\beta}})^2}$$

For simplicity, I use $m = exp(X\beta)$ and $g = \frac{m}{(1 - exp(-m))}$

```
trunc_pois_Newton_Raphson <- function(x, y, b_init = rep(1, ncol(x)),</pre>
                                        tol = 1*10^{(-8)}
{
  change <- Inf
  b_old <- b_init</pre>
  while(change>tol)
    eta <- x %*% b_old
    em <- exp(eta)
    w \leftarrow diag(as.vector((em-(em * exp(-em))-(em^2)*exp(-em))/(1-exp(-em)^2)),
                nrow = nrow(x),
               ncol = nrow(x)
    g \leftarrow em/(1-exp(-em))
    b_{new} \leftarrow b_{old} + solve(t(x)) %% w %% x) %% t(x) %% (y-g)
    change <- sqrt(sum((b_new - b_old)^2))</pre>
    b old <- b new
  b_new
```

I will test my algorithm using the college dataset we used on a previous assignment.

```
college <- read_csv("data/college-data.csv")</pre>
```

```
trunc_pois_Newton_Raphson(x, y)
```

```
[,1]
[1,] 3.749873567
[2,] 0.001541439
[3,] -0.015000046
```

Note that this result appears relatively similar to the result for the zero truncated glm function I found online. Citation (https://stats.oarc.ucla.edu/r/dae/zero-truncated-poisson/.)

Cross Check with Zero Truncated GLM

```
library(VGAM)
```

Loading required package: stats4

Loading required package: splines

```
college <- college |>
  mutate(
    out_tuition = out_of_state_tuition/2000
)

model <- vglm(total ~ out_tuition + type, family = pospoisson(), data = college)
summary(model)</pre>
```

Call:

```
vglm(formula = total ~ out_tuition + type, family = pospoisson(),
    data = college)
```

Coefficients:

Estimate Std. Error z value Pr(>|z|) (Intercept) 2.851016 0.027530 103.560 <2e-16 *** out_tuition 0.052979 0.001289 41.096 <2e-16 *** typePublic 0.002543 0.018548 0.137 0.891

Signif. codes: 0 '***' 0.001 '**' 0.01 '*' 0.05 '.' 0.1 ' ' 1

Name of linear predictor: loglink(lambda)

Log-likelihood: -1580.193 on 545 degrees of freedom

Number of Fisher scoring iterations: 4

No Hauck-Donner effect found in any of the estimates

By default, glm() calls iteratively re-weighted least squares.

No, the pareto 1 distribution is not in the exponential family.