1. For 
$$X \sim f(x; \theta)$$
, show that

$$E\left\{\frac{\partial}{\partial \theta}\log f(x;\theta)\right\} = 0, \text{ and } -E\left\{\frac{\partial^2}{\partial \theta^2}\log f(x;\theta)\right\} = E\left[\left\{\frac{\partial}{\partial \theta}\log f(x;\theta)\right\}^2\right].$$

(i) 
$$E\left[\frac{1}{9}\sqrt{\frac{1}{9}}\sqrt{\frac{1}{9}}\right] = E\left[\frac{1}{9}\sqrt{\frac{1}{9}}\sqrt{\frac{1}{9}}\right] = \int \frac{1}{9}\sqrt{\frac{1}{9}}\sqrt$$

$$\begin{bmatrix}
\frac{4\sigma}{\sigma_{1}} + \sigma(x) & \frac{4\sigma}{\sigma_{2}} & \frac{4\sigma}{\sigma_{2}} & \frac{4\sigma(x)}{\sigma_{2}} \\
\frac{4\sigma}{\sigma_{2}} + \sigma(x) & \frac{4\sigma}{\sigma_{2}} &$$

$$\frac{1}{2} - \mathbb{E}\left[\frac{\partial^2}{\partial \theta^2} \mathbb{A}f[\theta](\theta)\right] = \mathbb{E}\left[\left(\frac{\partial}{\partial \theta} \mathbb{A}f[\theta](\theta)\right)^{\frac{1}{2}}\right]$$

- 2. Suppose we have  $y_i \stackrel{\text{ind}}{\sim} Poisson(\mu_i), i = 1, \dots, n$ .
  - (a) Show that the Poisson distribution belongs to the exponential dispersion family to identify  $\theta_i$ ,  $b(\theta_i)$ ,  $a(\phi)$  under the exponential dispersion family form given in Lecture note.
  - (b) Derive the canonical link function  $g(\mu_i)$  to be modeled via  $\eta_i = \mathbf{x}_i \boldsymbol{\beta}, i = 1, \dots, n$  for the Poisson regression.

$$\frac{\left\{\left(y_{i}\right) = \frac{e^{-\mu_{i}} \mu_{i} b_{i}}{y_{i} b_{i}} \quad y_{i} = e^{-\mu_{i}} \mu_{i} b_{i}}{y_{i} b_{i}} \quad y_{i} = e^{-\mu_{i}} \mu_{i} b_{i}}$$

$$\frac{(\alpha)}{(A^i)} = \frac{A^i \beta}{C_{M^i} M^i \beta^i} = \exp\left(\frac{A}{C_{M^i} M^i \beta^i}\right) = \exp\left(-M^i + A^i \beta M^i - \lambda^i A^i \beta^i\right)$$

# ST509\_HW3\_Group 2

3. Write your own code to estimate the parameters  $\beta$  in the Poisson regression based on Newton-Raphson method. Namely, we have the following model for  $(y_i, \mathbf{x_i})$ ,  $i = 1, \dots, n$ :

$$y_i \mid \mathbf{x_i} \sim \text{Poisson}(\mu_i), \quad i = 1, \cdots, n$$
 
$$g(\mu_i) = \eta_i = \mathbf{x_i} \beta$$

with g being the canonical link function obtained in Problem 2-(b).

#### in R

where

```
my_poi <- function(X, y, init = NULL, max_iter = 1000, eps = 1e-5, offset = NULL) {
    # Scaling
    y_sd \leftarrow sd(y); X_sd \leftarrow apply(X, 2, sd)
    y \leftarrow y /y_sd; X \leftarrow t(t(X)/X_sd)
    # Add intercept term
    X \leftarrow cbind(1, X) ; n \leftarrow nrow(X); p \leftarrow ncol(X)
    if (is.null(init)) init <- rep(0, p); beta <- init</pre>
    # Offset
    if (is.null(offset)) offset <- rep(0, n)</pre>
    # Iteration
    for (iter in 1:max_iter) {
        eta <- X %*% beta
        w <- exp(eta + offset) # Mean = Variance
        z \leftarrow eta + (y-w)/w
         # IRLS
        X_tilde <- diag(c(sqrt(w))) %*% X</pre>
         z_tilde <- diag(c(sqrt(w))) %*% z</pre>
         qr_obj <- qr(X_tilde)</pre>
        new_beta <- backsolve(qr_obj$qr, qr.qty(qr_obj, z_tilde))</pre>
         # Check Convergence
         if (max(abs(new_beta - beta))/ max(abs(beta)) < eps) break</pre>
        beta <- new_beta
    }
    # Warning
    if (iter == max_iter) warning("Algorithm may not have converged!")
    # Restore beta coef
    beta <- c(beta) * c(1, 1/X_sd) + c(log(y_sd), rep(0, p-1))
    # Result
    list(est = t(beta), iterations = iter)
}
```

## in Python

```
import numpy as np
import pandas as pd
import statsmodels.api as sm
from statsmodels.genmod.families import Poisson
from scipy.linalg import solve_triangular
def my_poi(X, y, init=None, max_iter=1000, eps=1e-5, offset=None):
    # Scaling
   y_sd = np.std(y); X_sd = np.std(X, axis=0)
   y = y / y_sd; X = X / X_sd[np.newaxis, :]
   # Add intercept term
   X = np.column_stack((np.ones(len(X)), X))
   n, p = X.shape
    # INIT
   if init is None:
       init = np.zeros(p)
   beta = init
    # Offset
   if offset is None:
       offset = np.zeros(n)
    # Iteration
   for iter in range(max_iter):
        eta = X @ beta
       w = np.exp(eta + offset)
        z = eta + (y - w)/w
        # IRLS
       X_tilde = np.sqrt(w)[:, np.newaxis] * X
        z_tilde = np.sqrt(w) * z
       Q, R = np.linalg.qr(X_tilde)
       new_beta = solve_triangular(R, Q.T @ z_tilde, lower=False)
        # Check convergence
        if np.max(np.abs(new_beta - beta)) / np.max(np.abs(beta)) < eps :</pre>
            break
       beta = new_beta
    # Warning
    if iter == max_iter - 1:
       print("Algorithm may not have converged!")
    # Restore beta coefficients
   beta = beta * np.concatenate(([1], 1/X_sd)) + np.concatenate(([np.log(y_sd)], np.zeros(p-1)))
   return beta, iter
```

# I. Scaling

exp term으로 인해 무한대로 발산하는 경우를 방지하기 위해, Data에 스케일링을 진행하여 회귀 계수를 예측하고 이를 복워하는 과정을 거친다.

$$\log \left( E \left[ \frac{\mathbf{Y}}{\sigma_Y} \right] \right) = \left[ 1 \mid \frac{X_1}{\sigma_1} \mid \dots \mid \frac{X_p}{\sigma_p} \right] \begin{bmatrix} \beta_0 \\ \beta_1 \\ \vdots \\ \beta_p \end{bmatrix}$$

which means

$$\log \begin{bmatrix} E(y_1/\sigma_Y) \\ E(y_2/\sigma_Y) \\ \vdots \\ E(y_n/\sigma_Y) \end{bmatrix} = \left[ 1 \mid \frac{X_1}{\sigma_1} \mid \dots \mid \frac{X_p}{\sigma_p} \right] \begin{bmatrix} \beta_0 \\ \beta_1 \\ \vdots \\ \beta_p \end{bmatrix}$$

equal to

$$\log (E[\mathbf{Y}]) = \begin{bmatrix} 1 \mid X_1 \mid \cdots \mid X_p \end{bmatrix} \begin{bmatrix} \beta_0^* \\ \beta_1^* \\ \vdots \\ \beta_p^* \end{bmatrix} + \log (\sigma_Y)$$

#### II. Offset

A response variable Y has an index  $t_i$  s.t. its expected value is proportional to  $t_i$ , e.g. an amount of time or spatial area over which the count is made.

Sample rate =  $y_i/t_i$  and  $E(Y_i/t_i) = \mu_i/t_i$ 

Poisson regression model for rate date will be

$$\log(\mu_i/t_i) = \log(\mu_i) - \log(t_i) = \beta_0 + \beta_1 x_{1i} + \dots + \beta_n x_{ni}$$

$$\log(\mu_i) = \beta_0 + \beta_1 x_{1i} + \dots + \beta_p x_{pi} + \beta_{p+1} x_{(p+1)i}$$

where  $\beta_{p+1} = 1$  and  $x_{(p+1)i} = \log(t_i)$ : offset

예시) 병원에서 일정 기간 동안 발생한 질병의 건수를 예측하는 모델을 만든다고 가정. 이때 각 병원의 환자 수는 예측에 중요한 요소가 된다. 만약 한 병원에서 한 달 동안 질병이 10건 발생했고, 다른 병원에서는 같은 기간 동안 20건 발생했다면, 이 차이는 단순히 두 번째 병원에서 질병이 더 많이 발생하기 때문일 수도 있지만, 환자수가 더 많아서일 수도 있다. 이 경우, 환자 수를 오프셋으로 사용하여 모델을 조정할 수 있다. 즉, 모델이 각병원의 환자 수에 따라 조정되어, 환자 한 명당 질병 발생 건수를 더 잘 예측할 수 있게 된다.

4. Apply your code and report your parameter estimates of the Poisson regression to "Smoking and Lung Cancer data".

해당 문제는 일반적인 포아송 회귀를 적용하여 사망자 수를 예측할 수도 있지만, offset을 이용해서 인구수를 조정하여 사망 비율을 예측할 수도 있다. 또한, 범주형 변수는 홈페이지의 설명대로 label encoding을 진행하여 적합하였다.

#### in R

```
setwd("C:/Users/jhmok/OneDrive/바탕 화면/학교/2024-1/통계계산방법론/과제/3")
dat <- read.table("smoking.dat")</pre>
dat$age <- as.numeric(factor(dat$age))</pre>
dat$smoke <- as.numeric(factor(dat$smoke, levels=c("no", "cigarPipeOnly", "cigarrettePlus", "cigarrette
X <- as.matrix(dat[,-4]); y <- as.matrix(dat[,4])</pre>
X_offset <- as.matrix(dat[,1:2]); pop <-as.matrix(dat[,3])</pre>
# Model (w/o offset)
my_poi(X, y)
## $est
##
                       age
                                smoke
                                               pop
## [1,] 2.425885 0.2618248 0.2689611 0.0004433254
##
## $iterations
## [1] 6
glm(dead ~ age + smoke + pop, data = dat, family = "poisson")
##
## Call: glm(formula = dead ~ age + smoke + pop, family = "poisson", data = dat)
##
## Coefficients:
## (Intercept)
                                    smoke
                        age
                                                   pop
                               0.2689611
                                             0.0004433
##
     2.4258846
                  0.2618248
## Degrees of Freedom: 35 Total (i.e. Null); 32 Residual
## Null Deviance:
## Residual Deviance: 1304 AIC: 1552
```

```
# Model (with offset)
my_poi(X_offset,y, offset=log(pop))
## $est
##
                      age
                             smoke
## [1,] -3.967813 0.331192 0.1638844
## $iterations
## [1] 12
glm(dead ~ age + smoke, data=dat, family="poisson", offset = log(pop))
## Call: glm(formula = dead ~ age + smoke, family = "poisson", data = dat,
## offset = log(pop))
##
## Coefficients:
## (Intercept)
                     age
                                 smoke
##
      -3.9679
                 0.3312
                                0.1639
##
## Degrees of Freedom: 35 Total (i.e. Null); 33 Residual
## Null Deviance: 4056
## Residual Deviance: 85.97 AIC: 332
```

## in Python

```
file_path = "C:/Users/jhmok/OneDrive/바탕 화면/학교/2024-1/통계계산방법론/과제/3/smoking.dat"
dat = pd.read_csv(file_path, sep='\s+')
dat['age'] = pd.Categorical(dat['age']).codes + 1
dat['smoke'] = pd.Categorical(dat['smoke'], categories=["no", "cigarPipeOnly", "cigarrettePlus", "cigar
X = dat.iloc[:, 0:3].values; y = dat.iloc[:, 3].values
X_offset = dat.iloc[:, 0:2]; pop = dat.iloc[:, 2].values
# Model (w/o offset)
beta_no_offset, iter_no_offset = my_poi(X, y)
coef_formatted = [f"{coef:.4f}" for coef in beta_no_offset]
print("coef :", coef_formatted, "iter :", iter_no_offset+1)
## coef : ['2.4259', '0.2618', '0.2690', '0.0004'] iter : 6
glm_no_offset = sm.GLM(y, sm.add_constant(X), family=Poisson(), offset=None).fit()
print(glm_no_offset.summary())
                Generalized Linear Model Regression Results
## Dep. Variable:
                                y No. Observations:
                                                                 36
                              GLM Df Residuals:
## Model:
                                                                 32
                           Poisson Df Model:
## Model Family:
                                                                  3
## Link Function:
                              Log Scale:
                                                             1.0000
                             IRLS Log-Likelihood:
## Method:
                                                             -771.89
## Date:
                       월, 01 4 2024 Deviance:
                                                              1303.8
## Time:
                          00:33:01 Pearson chi2:
                                                            1.12e+03
                               5 Pseudo R-squ. (CS):
## No. Iterations:
                                                              1.000
## Covariance Type: nonrobust
coef std err z P>|z|
                                                    [0.025
##

      0.057
      42.777
      0.000

      0.006
      45.608
      0.000

              2.4259
## const
                                                     2.315
                                                               2.537
             0.2618
## x1
                                                     0.251
                                                              0.273
             0.2690 0.012 22.191
## x2
                                         0.000
                                                     0.245
                                                              0.293
            0.0004 6.29e-06 70.484 0.000
## x3
                                                     0.000
                                                              0.000
```

```
# Model (with offset)
beta_offset, iter_offset = my_poi(X_offset, y, offset = np.log(pop))
coef_formatted2 = [f"{coef:.4f}" for coef in beta_offset]
print("coef :", coef_formatted2, "iter :", iter_no_offset+1)
## coef : ['-3.9678', '0.3312', '0.1639'] iter : 6
glm_offset = sm.GLM(y, sm.add_constant(X_offset), family=Poisson(), offset=np.log(pop)).fit()
print(glm_offset.summary())
              Generalized Linear Model Regression Results
y No. Observations:
## Dep. Variable:
                                                          36
## Model:
                           GLM Df Residuals:
                                                          33
## Model Family:
                       Poisson Df Model:
                                                           2
## Link Function:
                           Log Scale:
                                                       1.0000
## Method:
                          IRLS Log-Likelihood:
                                                      -163.00
                    월, 01 4 2024 Deviance:
## Date:
                                                       85.970
## Time:
                       00:33:01 Pearson chi2:
                                                        82.9
## No. Iterations:
                            6 Pseudo R-squ. (CS):
                                                        1.000
## Covariance Type:
                      nonrobust
##
          coef std err z P>|z| [0.025
                                                       0.975]
## const
          -3.9679 0.055 -71.593
                                      0.000
                                              -4.077
                                                       -3.859
## age 0.3312 0.005 60.423 0.000
## smoke 0.1639 0.012 13.398 0.000
                                              0.320
                                                       0.342
                                               0.140
                                                       0.188
```