## stat431 a3 q4

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```
# step 0
N <- 1000
x1 <- rnorm(N, 0, 1)
x2 <- rbinom(N, 1, 0.25)

estimates <- matrix(NA,nrow=3, ncol=500)
CIs <- matrix(NA,nrow=6,ncol=500)</pre>
```

## (a)

```
for (i in 1:500) {
  beta0 <- 0.5
  beta1 <- 0.2
  beta2 <- 0.5
  y \leftarrow rpois(N, exp(beta0 + beta1*x1 + beta2*x2))
  mod \leftarrow glm(y \sim x1 + x2, family = poisson)
  est_beta0 <- summary(mod)$coefficients[1,1]</pre>
  se0 <- summary(mod)$coefficients[1,2]</pre>
  est_beta1 <- summary(mod)$coefficients[2,1]</pre>
  se1 <- summary(mod)$coefficients[2,2]</pre>
  est_beta2 <- summary(mod)$coefficients[3,1]</pre>
  se2 <- summary(mod)$coefficients[3,2]</pre>
  estimates[1,i] <- est_beta0</pre>
  estimates[2,i] <- est_beta1</pre>
  estimates[3,i] <- est_beta2</pre>
  c <- 1.96
  CIs[1,i] <- est_beta0 - c*se0
  CIs[2,i] \leftarrow est_beta0 + c*se0
  CIs[3,i] <- est_beta1 - c*se1
  CIs[4,i] <- est_beta1 + c*se1</pre>
  CIs[5,i] <- est_beta2 - c*se2
  CIs[6,i] <- est_beta2 + c*se2
# average estimates
```

```
average_est_beta0 <- mean(estimates[1,])</pre>
average_est_beta1 <- mean(estimates[2,])</pre>
average_est_beta2 <- mean(estimates[3,])</pre>
average_est_beta0
## [1] 0.4987404
average_est_beta1
## [1] 0.1994879
average_est_beta2
## [1] 0.5001897
# proportion of confidence interval coverage
coverage <- matrix(NA, nrow=3, ncol=500)</pre>
for (i in 1:500) {
  L0 \leftarrow CIs[1,i]
  U0 <- CIs[2,i]
  L1 \leftarrow CIs[3,i]
  U1 <- CIs[4,i]
  L2 <- CIs[5,i]
  U2 <- CIs[6,i]
  coverage[1,i] \leftarrow (beta0 >= L0) & (U0 >= beta0)
  coverage[2,i] \leftarrow (beta1 >= L1) & (U1 >= beta1)
  coverage[3,i] \leftarrow (beta2 >= L2) & (U2 >= beta2)
beta0_pro <- mean(coverage[1,])</pre>
beta1_pro <- mean(coverage[2,])</pre>
beta2_pro <- mean(coverage[3,])</pre>
beta0_pro
## [1] 0.962
beta1_pro
## [1] 0.956
beta2_pro
## [1] 0.958
```

Comments: Based on the results, we found that average estimates are very close to the true values; the proportion of confidence intervals that cover true value is very close to 0.95 (since we constructed 95% CI).

(b)

```
N <- 1000
x1 <- rnorm(N, 0, 1)
x2 <- rbinom(N, 1, 0.25)
estimates <- matrix(NA, nrow=3, ncol=500)</pre>
```

```
CIs <- matrix(NA, nrow=6, ncol=500)</pre>
for (i in 1:500) {
  beta0 <- 0.5
  beta1 <- 0.2
  beta2 <- 0.5
  gamma0 <- 0.5
  gamma1 <- 0
  gamma2 <- 0.5
  y_star <- rpois(N, exp(beta0 + beta1*x1 + beta2*x2))</pre>
  pie \leftarrow exp(gamma0 + gamma1*x1 + gamma2*x2) /
    (1 + \exp(\text{gamma0} + \text{gamma1}*x1 + \text{gamma2}*x2))
  w <- rbinom(N, 1, pie)
  y \leftarrow c(NA, N)
  y[w==1] <- 0
  y[w==0] \leftarrow y_star[w==0]
  # fit model
  mod \leftarrow glm(y \sim x1 + x2, family = poisson)
  est_beta0 <- summary(mod)$coefficients[1,1]</pre>
  se0 <- summary(mod)$coefficients[1,2]</pre>
  est_beta1 <- summary(mod)$coefficients[2,1]</pre>
  se1 <- summary(mod)$coefficients[2,2]</pre>
  est_beta2 <- summary(mod)$coefficients[3,1]</pre>
  se2 <- summary(mod)$coefficients[3,2]</pre>
  estimates[1,i] <- est_beta0</pre>
  estimates[2,i] <- est_beta1</pre>
  estimates[3,i] <- est_beta2</pre>
  c <- 1.96
  CIs[1,i] <- est_beta0 - c*se0</pre>
  CIs[2,i] <- est_beta0 + c*se0</pre>
  CIs[3,i] <- est_beta1 - c*se1
  CIs[4,i] <- est_beta1 + c*se1
  CIs[5,i] <- est_beta2 - c*se2</pre>
  CIs[6,i] \leftarrow est_beta2 + c*se2
}
# average estimates
average_est_beta0 <- mean(estimates[1,])</pre>
average_est_beta1 <- mean(estimates[2,])</pre>
average_est_beta2 <- mean(estimates[3,])</pre>
```

```
average_est_beta0
## [1] -0.4781659
average_est_beta1
## [1] 0.1956083
average_est_beta2
## [1] 0.1494656
# proportion of confidence interval coverage
coverage <- matrix(NA, nrow=3, ncol=500)</pre>
for (i in 1:500) {
  L0 \leftarrow CIs[1,i]
  U0 <- CIs[2,i]
  L1 <- CIs[3,i]
  U1 <- CIs[4,i]
  L2 <- CIs[5,i]
  U2 <- CIs[6,i]
  coverage[1,i] \leftarrow (beta0 >= L0) & (U0 >= beta0)
  coverage[2,i] \leftarrow (beta1 >= L1) & (U1 >= beta1)
  coverage[3,i] \leftarrow (beta2 >= L2) & (U2 >= beta2)
}
beta0_pro <- mean(coverage[1,])</pre>
beta1_pro <- mean(coverage[2,])</pre>
beta2_pro <- mean(coverage[3,])</pre>
beta0_pro
## [1] 0
beta1_pro
## [1] 0.792
beta2_pro
```

## [1] 0.108

Comments: Based on the results, we found that average estimates are not close to the true values; the proportion of confidence intervals that cover true value is much lower than 0.95 (we constructed 95% CI).

(c)

```
N <- 1000
x1 <- rnorm(N, 0, 1)
x2 <- rbinom(N, 1, 0.25)
estimates <- matrix(NA,nrow=3, ncol=500)
CIs <- matrix(NA,nrow=6,ncol=500)

for (i in 1:500) {</pre>
```

```
beta0 <- 0.5
  beta1 <- 0.2
  beta2 <- 0.5
  gamma0 <- 0.5
  gamma1 <- 0
  gamma2 <- 0.5
  y_star <- rpois(N, exp(beta0 + beta1*x1 + beta2*x2))</pre>
  pie <- exp(gamma0 + gamma1*x1 + gamma2*x2) /</pre>
    (1 + \exp(\text{gamma0} + \text{gamma1*x1} + \text{gamma2*x2}))
  w <- rbinom(N, 1, pie)
  y \leftarrow c(NA, N)
  y[w==1] <- 0
  y[w==0] <- y_star[w==0]
  # fit model
  require(pscl)
  mod \leftarrow zeroinfl(y \sim x1 + x2)
  est_beta0 <- summary(mod)$coefficients$count[1]</pre>
  se0 <- sqrt(summary(mod)$vcov[1,1])</pre>
  est_beta1 <- summary(mod)$coefficients$count[2]</pre>
  se1 <- sqrt(summary(mod)$vcov[2,2])</pre>
  est_beta2 <- summary(mod)$coefficients$count[3]</pre>
  se2 <- sqrt(summary(mod)$vcov[3,3])</pre>
  estimates[1,i] <- est_beta0</pre>
  estimates[2,i] <- est_beta1</pre>
  estimates[3,i] <- est_beta2</pre>
  c <- 1.96
  CIs[1,i] <- est_beta0 - c*se0
  CIs[2,i] \leftarrow est beta0 + c*se0
  CIs[3,i] <- est_beta1 - c*se1</pre>
  CIs[4,i] <- est_beta1 + c*se1
  CIs[5,i] <- est_beta2 - c*se2</pre>
  CIs[6,i] <- est_beta2 + c*se2</pre>
}
## Loading required package: pscl
## Classes and Methods for R developed in the
## Political Science Computational Laboratory
## Department of Political Science
## Stanford University
## Simon Jackman
## hurdle and zeroinfl functions by Achim Zeileis
```

```
# average estimates
average_est_beta0 <- mean(estimates[1,])</pre>
average_est_beta1 <- mean(estimates[2,])</pre>
average_est_beta2 <- mean(estimates[3,])</pre>
average_est_beta0
## [1] 0.4938926
average_est_beta1
## [1] 0.1998101
average_est_beta2
## [1] 0.502593
# proportion of confidence interval coverage
coverage <- matrix(NA, nrow=3, ncol=500)</pre>
for (i in 1:500) {
  LO <- CIs[1,i]
  U0 <- CIs[2,i]
  L1 <- CIs[3,i]
  U1 <- CIs[4,i]
  L2 <- CIs[5,i]
  U2 <- CIs[6,i]
  coverage[1,i] \leftarrow (beta0 >= L0) & (U0 >= beta0)
  coverage[2,i] <- (beta1 >= L1) & (U1 >= beta1)
  coverage[3,i] \leftarrow (beta2 >= L2) & (U2 >= beta2)
}
beta0_pro <- mean(coverage[1,])</pre>
beta1_pro <- mean(coverage[2,])</pre>
beta2_pro <- mean(coverage[3,])</pre>
beta0_pro
## [1] 0.952
beta1_pro
## [1] 0.942
beta2_pro
```

```
## [1] 0.946
```

Comments: Based on the results, we found that average estimates are very close to the true values; the proportion of confidence intervals that cover true value is very close to 0.95 (since we constructed 95% CI).

```
(d)
```

```
N <- 1000

x1 <- rnorm(N, 0, 1)

x2 <- rbinom(N, 1, 0.25)
```

```
estimates <- matrix(NA, nrow=3, ncol=500)</pre>
CIs <- matrix(NA, nrow=6, ncol=500)
for (i in 1:500) {
  beta0 <- 0.5
  beta1 <- 0.2
  beta2 <- 0.5
  gamma0 <- 0.5
  gamma1 <- 0
  gamma2 <- 0.5
  y_star \leftarrow rpois(N, exp(beta0 + beta1*x1 + beta2*x2))
  pie <- exp(gamma0 + gamma1*x1 + gamma2*x2) /</pre>
    (1 + \exp(\text{gamma0} + \text{gamma1*x1} + \text{gamma2*x2}))
  w <- rbinom(N, 1, pie)
  y \leftarrow c(NA, N)
  y[w==1] <- 0
  y[w==0] \leftarrow y_star[w==0]
  # fit model and use ad-hoc method
  mod \leftarrow glm(y \sim x1 + x2, family = poisson)
  estimate_dispersion <- summary(mod)$deviance / summary(mod)$df.residual</pre>
  est_beta0 <- summary(mod)$coefficients[1,1]</pre>
  se0_adj <- summary(mod)$coefficients[1,2] * sqrt(estimate_dispersion)</pre>
  est_beta1 <- summary(mod)$coefficients[2,1]</pre>
  se1_adj <- summary(mod)$coefficients[2,2] * sqrt(estimate_dispersion)</pre>
  est_beta2 <- summary(mod)$coefficients[3,1]</pre>
  se2_adj <- summary(mod)$coefficients[3,2] * sqrt(estimate_dispersion)</pre>
  estimates[1,i] <- est_beta0</pre>
  estimates[2,i] <- est_beta1</pre>
  estimates[3,i] <- est_beta2</pre>
  c < -1.96
  CIs[1,i] <- est_beta0 - c*se0_adj
  CIs[2,i] <- est_beta0 + c*se0_adj
  CIs[3,i] <- est_beta1 - c*se1_adj</pre>
  CIs[4,i] <- est_beta1 + c*se1_adj</pre>
  CIs[5,i] <- est_beta2 - c*se2_adj
  CIs[6,i] \leftarrow est_beta2 + c*se2_adj
}
# average estimates
```

```
average_est_beta0 <- mean(estimates[1,])</pre>
average_est_beta1 <- mean(estimates[2,])</pre>
average_est_beta2 <- mean(estimates[3,])</pre>
average_est_beta0
## [1] -0.4761964
average_est_beta1
## [1] 0.1969907
average_est_beta2
## [1] 0.1531694
# proportion of confidence interval coverage
coverage <- matrix(NA, nrow=3, ncol=500)</pre>
for (i in 1:500) {
  LO <- CIs[1,i]
  U0 <- CIs[2,i]
  L1 \leftarrow CIs[3,i]
  U1 <- CIs[4,i]
  L2 <- CIs[5,i]
  U2 <- CIs[6,i]
  coverage[1,i] \leftarrow (beta0 >= L0) & (U0 >= beta0)
  coverage[2,i] \leftarrow (beta1 >= L1) & (U1 >= beta1)
  coverage[3,i] \leftarrow (beta2 >= L2) & (U2 >= beta2)
beta0_pro <- mean(coverage[1,])</pre>
beta1_pro <- mean(coverage[2,])</pre>
beta2_pro <- mean(coverage[3,])</pre>
beta0_pro
## [1] 0
beta1_pro
## [1] 0.904
beta2_pro
```

## [1] 0.218

Comments: By using ad-hoc method, it does not yield valid inference. Since the proportion of confidence intervals that cover the true value is till far less than 0.95 after using estimated dispersion parameter.