

## Homework #8

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## Problem 8.1:

```

1 tab <- matrix(c(159,22,8,14),nrow = 2,ncol = 2)
2 mcnemar.test(tab, correct = FALSE)
3 # McNemar's Chi-squared test
4 #
5 # data:  tab
6 # McNemar's chi-squared = 6.5333, df = 1, p-value = 0.01059

```

From the results above, the McNemar chi-squared = 6.53 ( $P$ -value = 0.011), indicating that low birth weight (cases) than normal birth weight (controls) are more likely to be smokers.

## Problem 8.3:

```

1 Opinions <- read.table("http://users.stat.ufl.edu/~aa/cat/data/Opinions.dat", header=TRUE)
2 fit_marginal <- gee(y ~ question, id=person, family=binomial(link=logit), data=Opinions)
3 print(summary(fit_marginal))
4 # Coefficients:
5 #               Estimate Naive S.E.   Naive z Robust S.E.   Robust z
6 # (Intercept) -0.8858933 0.06505597 -13.617401  0.06502753 -13.623357
7 # question     0.1035319 0.09107816  1.136737  0.06397794  1.618244
8 fit_subject <- glmer(y ~ (1|person) + question, family=binomial, nAGQ=50, data=Opinions)
9 print(summary(fit_subject))
10 # Random effects:
11 #   Groups Name      Variance Std.Dev.
12 #   person (Intercept) 8.143    2.854
13 # Number of obs: 2288, groups:  person, 1144
14 #
15 # Fixed effects:
16 #               Estimate Std. Error z value Pr(>|z|)
17 # (Intercept)  -1.8343      0.1624 -11.295  <2e-16 ***
18 # question      0.2100      0.1301  1.614   0.106

```

Marginal model focuses on the marginal distributions of the two responses, and its  $\beta$  is the log odds ratio comparing the marginal distributions (i.e. the log odds ratio for the combined sample). Subject-specific model focuses on the partial table of the two responses conditional on the subject, and its  $\beta$  is the log odds ratio conditional on the subject. To illustrate that, using the environment data, we can find the above results where for the marginal model has  $\hat{\beta} = \log((359/785)/(334/810)) = 0.104$ . and the subject-specific model has  $\hat{\beta} = \log(132/107) = 0.21$ .

**Problem 8.7:**

```

1 Survey <- data.frame(I=c(1,1,1,0,0,0), II=c(-1,0,0,1,1,0), III=c(0,-1,0,-1,0,1), IV=c(0,0,-1,0,-1,-1),
  nij=c(49,3,288,2,186,12), nji=c(129,1,81,0,9,4))
2 symm <- glm(nij/(nij+nji) ~ -1, family=binomial, weights=nij+nji, data=Survey)
3 print(summary(symm))
4 # Residual deviance: 365.81 on 6 degrees of freedom
5 res <- rstandard(symm, type="pearson")
6 #      1      2      3      4      5      6
7 # -5.996254  1.000000 10.775990  1.414214 12.675233  2.000000
8 QS <- glm(nij/(nij+nji) ~ -1 + I + II + III + IV, family=binomial, weights=nij+nji, data=Survey)
9 print(summary(QS)) # quasi-symmetry model
10 # Residual deviance:  5.5347 on 3 degrees of freedom

```

For (a), Symmetry model has residual deviance = 365.81 ( $df = 6$ ) which is such a bad fit. From the results we can observe that if one can do more shifts from II to I (resstandard = 6.0), I to IV (resstandard = 10.8), II to IV (resstandard = 12.7) then it would be symmetry.

For (b), QS model has residual deviance = 5.53 ( $df = 3$ ) which indicating fits well. QS compares to symmetry model resulting in difference =  $365.81 - 5.53 = 360.3$  ( $df = 6 - 3 = 3$  and  $P\text{-value} < .001$ ), providing strong evidence against marginal homogeneity. There are some reasons that includes changes in II (Catholic) or IV (none or other) and large sample size.

**Problem 8.11:**

```

1 coffee <- data.frame(first=c(1,1,1,1,1,2,2,2,2,2,3,3,3,3,3,4,4,4,4,4,5,5,5,5,5), second=c
  (1,2,3,4,5,1,2,3,4,5,1,2,3,4,5,1,2,3,4,5,1,2,3,4,5), count=c
  (93,17,44,7,10,9,46,11,0,9,17,11,155,9,12,6,4,9,15,2,10,4,12,2,27), diag=c
  (1,0,0,0,0,0,2,0,0,0,0,0,3,0,0,0,0,0,4,0,0,0,0,0,5))
2 ordinary <- glm(count~factor(first)+factor(second), family=poisson, data=coffee)
3 print(summary(ordinary))
4 # Residual deviance: 346.38 on 16 degrees of freedom
5 quasi <- glm(count~factor(first)+factor(second)+factor(diag), family=poisson, data=coffee)
6 print(summary(quasi))
7 # Residual deviance:  13.786 on 11 degrees of freedom

```

For the ordinary model, we find that model fits bad with r.d. = 346.4 ( $df = 16$ ). Yet, for the quasi-independence model, we find that model fits quite well with r.d. = 13.8 ( $df = 11$ ). Hence, we can say that the second choice of coffe brand is independent to the first one, under quasi-independence model..