STA 312f2012 Quiz 10

- 1. For homework, you examined a log-linear model for a 3×2 contingency table of race by belief in life after death
 - (a) (1 point) Write a saturated linear model for $\log \mu_{ij}$ using the λ notation. Race is X and Belief in afterlife is Y.

(b) (1 point) Write a saturated linear model for $\log \mu_{ij}$ using a regression model with effect coding; this is the β notation. You do not have to say how the dummy variables are defined, but the dummy variables for race should be named r-something and the dummy variables for belief should be named b-something so we know what they are.

(c) (1 point) In terms of the λ notation, what is the null hypothesis when you are testing for independence?

$$H_{o}$$
; $\gamma_{ij}^{xy} = 0$

(d) (1 point) In terms of the λ notation, how many quantities (not unique quantities) are you setting to zero under H_0 ? The answer is a number.

(e) (1 point) In terms of the β notation, what is the null hypothesis when you are testing for independence?

(f) (1 point) In terms of the β notation, how many quantities are you setting to zero under H_0 ? The answer is a number.

- 2. For the table relating Alcohol use (A), Cigarette use (B) and Marijuana use (M), you used R fit the model (AC)(AM)(CM).
 - (a) (1 point) Under the model, what is the estimated expected number of students who do not drink alcohol, do not smoke cigarettes, and do not use marijuana? The answer is a number from your printout.

(b) (1 point) What is the Likelihood Ratio test statistic G^2 for the test of model fit? The answer is a number from your printout.

(c) (2 points) You tested whether Alcohol and cigarette use were conditionally independent given Cigarette use. Give the Likelihood Ratio test statistic G^2 . The answer is a number from your printout.

Attach your printout for Question 2 (Homework Question 3). Make sure your name is written on the printout.

Quiz 10 Printout

```
> # Question 3, Data of Table 7.3 on p. 209.
> drugs = read.table(stdin())
      Alcohol Cigarette
                         Marijuana
                                       Freq
1: 1
                 Yes
        Yes
                            Yes
                                        911
2: 2
        Yes
                 Yes
                                        538
                            No
3: 3
        Yes
                            Yes
                                         44
                 No
4: 4
        Yes
                 No
                                        456
                            No
5: 5
                 Yes
                            Yes
                                          3
        No
6: 6
                                         43
        No
                 Yes
                            No
7: 7
        No
                 No
                            Yes
                                          2
8:8
        No
                 No
                            No
                                        279
9:
> # Take a look: See 2 sub-tables
> xtabs(Freq ~ Cigarette+Marijuana+Alcohol,data=drugs)
, , Alcohol = No
         Marijuana
Cigarette No Yes
      No 279
                2
      Yes
          43
                3
, , Alcohol = Yes
         Marijuana
Cigarette No Yes
      No 456 44
      Yes 538 911
```

```
> # Order Yes and No is alphabetical, different from text.
> dtab = xtabs(Freq ~ Alcohol+Cigarette+Marijuana,data=drugs)
> drugmod = loglin(dtab, margin=list(c(1,2),c(1,3),c(2,3)), fit=T, param=T)
5 iterations: deviation 0.03408344
> drugmod
$1rt
[1] <mark>0.3739859</mark>
$pearson
[1] 0.4010998
$df
[1] 1
$margin
$margin[[1]]
[1] "Alcohol"
                 "Cigarette"
$margin[[2]]
[1] "Alcohol"
                 "Marijuana"
$margin[[3]]
[1] "Cigarette" "Marijuana"
$fit
, , Marijuana = No
       Cigarette
Alcohol
                No
                           Yes
    No 279.614402 42.383882
                                    2a
    Yes 455.385598 538.616118
, , Marijuana = Yes
       Cigarette
Alcohol
                 No
                           Yes
    No
          1.383160
                      3.616919
    Yes 44.616840 910.383081
```

```
$param$`(Intercept)`
[1] 4.251537
$param$Alcohol
                Yes
       No
-1.503994 1.503994
$param$Cigarette
        No
                  Yes
-0.2822777 0.2822777
$param$Marijuana
       No
                Yes
 1.196045 -1.196045
$param$Alcohol.Cigarette
       Cigarette
Alcohol
                           Yes
                No
    No
         0.5136255 -0.5136255
    Yes -0.5136255 0.5136255
$param$Alcohol.Marijuana
       Marijuana
Alcohol
               No
                        Yes
         0.746502 -0.746502
    No
    Yes -0.746502 0.746502
$param$Cigarette.Marijuana
         Marijuana
Cigarette
                  No
                             Yes
           0.7119739 -0.7119739
      No
      Yes -0.7119739 0.7119739
> # Test of A*M from Table 7.5 gives G^2 = 91.64
> redAM = loglin(dtab,margin=list(c(1,2),c(2,3))) # (1,3) is missing
2 iterations: deviation 0
> G2 = redAM$1rt-drugmod$1rt; G2 # 91.64437
[1] <mark>91.64437</mark> | 2c
```

\$param