Contigency table, Logistic Regression and Poisson Regression

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- 1. Analysis comparing proportions and using contingency tables:
 - a. The 2 by 2 table of sex by like is shown below, as t:

```
like
sex 0 1
Female 134 114
Male 29 122
```

There are evidence that in fact, sex is not independent of a student's preference for playing video games. The p-value from both prop.test(t, correct = FALSE) and fisher.test(t) gives significant evidence to reject the hypothesis that sex is independent of a student's preference for playing video games. Equals 6.704e-12 and 2.515e-12 respectively. In practical terms, male tends to like playing video games than female at a statistical significant level, given the odd ratio from fisher.test(t), the odds of preference of playing video games for a male person is 4.9247 times that for a female person.

- b. Construct two tables with different expected grade for male and female respectively,
 - > expect_Aplus

0 1 Female 31 26 Male 11 32

For students expect grade of A+, from fisher.test(expect_Aplus) and chisq.test(expect_Aplus, correct = FALSE), both of p-value significant being 0.004462, 0.003861. Indicating that there are significant evidence that sex is not independent of preference. With an odd ratio of 3.4237, indicating that for students expecting grade of A+, the odds of preference of playing video games for a male person is 3.4237 times that for a female person.

> expectOther

0 1
Female 103 88
Male 18 90

For students expect grade others, from fisher.test(expect_Aplus) and chisq.test(expect_Aplus, correct = FALSE), both of p-value significant being 1.048e-10, 2.877e-10. Indicating that there are significant evidence that sex is not independent of preference. With an odd ratio of 5.8175, indicating that for students expecting other grades, the odds of preference of playing video games for a male person is 5.8175 times that for a female person.

By calculating the proportion of preference of video games for male and female with expectation of grade A+ is 0.7442 and 0.4561 respectively, and with expectation of other grades, the proportion for male and female likes video games is 0.8333 and 0.4607 respectively. There is an increase of proportion of liking video games for both male and female when do not expect a grade of A+, although not really obvious for female.

2. Analysis using Logistic Regression:

a. For model 2.1, $\log(\frac{\hat{\pi}}{1-\hat{\pi}}) = -0.1574 + 1.7668\mathbb{I}_{\texttt{sex_f}} - 0.0185\mathbb{I}_{\texttt{grade_f}} - 0.5231\mathbb{I}_{\texttt{sex_f}} * \mathbb{I}_{\texttt{grade_f}}$ For model 2.2 without interaction term: $\log(\frac{\hat{\pi}}{1-\hat{\pi}}) = -0.1189 + 1.6111\mathbb{I}_{\texttt{sex_f}} - 0.1871\mathbb{I}_{\texttt{grade_f}}$ where π is is the estimated probability that likes playing video games $\mathbb{I}_{\texttt{sex_f}}$:1 if being male and 0 otherwise.

 \mathbb{I}_{grade_f} : 1 if the expecting grade of A+ and 0 otherwise.

Conduct LRT and Wald test to see if the additive model is better than interaction model. For LRT: H_0 : additive model is adequate, and H_1 : interaction model is better.

Observe the test statistic $G^2=489.37-488.41=0.96\sim\chi_1^2$, with $P(\chi_1^2>0.96)>0.10$ non significant, with p-value in Appendix.

Wald test: with $H_0: \gamma^3=0$ and $H_1: \gamma^3\neq 0$, observe the test statistic $Z=\frac{-0.5231}{0.5297}=-0.9887, Z^2\approx 0.9775\sim \chi_1^2$ with $P(Z^2>0.9775)>0.10$, non significant, with p-value in Appendix.

Since the p-value is large, we fail to reject the null hypothesis and conclude that the data are consistent with the coefficient of the interaction term being 0. Therefore, the interaction does not contributes in a statistically significant way to the explanation of the odds of likes playing video games. In conclusion, mod2.2 without interaction term should be used.

b. From estimated coefficient of $\mathbb{I}_{\texttt{sexMale}}$, $\beta_2 = 1.6111$, the odds ratio of which compares to to female, is $\exp(1.6111) \approx 5.0083$, which is similar to the conclusion drawn from fisher.test() in part a.

From estimated coefficient of \mathbb{I}_{gradeA^+} , $\beta_3 = -0.1871$, the odds ratio of which compares to whom that do not expect a grade of A+, is $\exp(-0.1871) \approx 0.8294$, which is agrees with the conclusion drawn from part 1 that the lower standards of expectation of grades does increase the proportion of liking video games.

3. Analysis using Poisson Regression:

a. First, construct the table, then construct mod3.1 and mod3.2 as shown in Appendix. The models are:

mod3.1

$$\begin{split} \log(\mathbb{E}(y_{ijk})) &= 4.6347 - 1.2007 \mathbb{I}_{\texttt{grade.p}} - 1.7444 \mathbb{I}_{\texttt{sex.p}} - 0.1574 \mathbb{I}_{\texttt{like.p}} + 0.7083 \mathbb{I}_{\texttt{grade.p}} * \mathbb{I}_{\texttt{sex.p}} - 0.01850 \mathbb{I}_{\texttt{grade.p}} * \mathbb{I}_{\texttt{like.p}} + 1.7668 \mathbb{I}_{\texttt{sex.p}} * \mathbb{I}_{\texttt{like.p}} - 0.5231 \mathbb{I}_{\texttt{grade.p}} * \mathbb{I}_{\texttt{sex.p}} * \mathbb{I}_{\texttt{like.p}} \\ & \texttt{mod 3.2:} \end{split}$$

 $\log(\mathbb{E}(y_{ijk})) = 4.6168 - 1.1256\mathbb{I}_{\texttt{grade_p}} - 1.6298\mathbb{I}_{\texttt{sex_p}} - 0.1189\mathbb{I}_{\texttt{like_p}} + 0.3547\mathbb{I}_{\texttt{grade_p}} * \mathbb{I}_{\texttt{sex_p}} - 0.1871\mathbb{I}_{\texttt{grade_p}} * \mathbb{I}_{\texttt{like_p}} + 1.6111\mathbb{I}_{\texttt{sex_p}} * \mathbb{I}_{\texttt{like_p}}$ where $\mathbb{E}(y_{ijk})$ is the counts classified by i, j, kth status.

 \mathbb{I}_{grade_p} : 1 if expecting grade of A+ and 0 otherwise.

 \mathbb{I}_{sex_p} : 1 if sex is male and 0 otherwise.

 \mathbb{I}_{like_p} : 1 if ith likes video games and 0 otherwise.

b. Comparison of results by:

- **Deviance**: The deviance between model3.1 and model3.2 is the same as the deviance between model2.1 and model2.2, which both result around 0.96
- -Wald tests: the Wald tests matches between:
 - i. interaction of sex and like in model3.2 and sex effect in model 2.1, with same p-value of 2.45e-09
- ii. interaction between grade, sex and like in model 3.1 and grade and sex in model 2.1, with p-value of 0.3230

- iii. interaction between grade and like in model 3.1 and grade effect in model 2.1, with p-value of 0.9510
- interpretation: Logistic model predicts the log-odd of like with $\log(\frac{\hat{\pi}_i}{1-\hat{\pi}_i})$ while the Poisson model predicts log of mean value of the counts, $\log(\mathbb{E}(y_{ijk}))$. Meanwhile the response variable is clear in Logistic model, a model is built for counts in the Poisson model.

Appendix

```
Q1:
> t = table(sex, like)
        like
           0
              1
sex
 Female 134 114
 Male 29 122
> fisher.test(t)
        Fisher's Exact Test for Count Data
data: t
p-value = 2.515e-12
alternative hypothesis: true odds ratio is not equal to 1
95 percent confidence interval:
3.008412 8.248768
sample estimates:
odds ratio
  4.924757
> chisq.test(t, correct = FALSE)
        Pearson's Chi-squared test
data: t
X-squared = 47.112, df = 1, p-value = 6.704e-12
> expect_Aplus <- table(sex[grade==1], like[grade==1])</pre>
> expect_Aplus
  Female 31 26
 Male 11 32
> expectOther = table(sex[grade==0], like[grade==0])
> expectOther
           0
               1
  Female 103 88
 Male
          18 90
> fisher.test(expect_Aplus)
```

```
data: expect_Aplus
p-value = 0.004462
alternative hypothesis: true odds ratio is not equal to 1
95 percent confidence interval:
1.36147 9.09891
sample estimates:
odds ratio
 3.423749
> fisher.test(expectOther)
       Fisher's Exact Test for Count Data
data: expectOther
p-value = 1.048e-10
alternative hypothesis: true odds ratio is not equal to 1
95 percent confidence interval:
 3.185573 11.085730
sample estimates:
odds ratio
 5.817501
> chisq.test(expect_Aplus, correct = FALSE)
       Pearson's Chi-squared test
data: expect_Aplus
X-squared = 8.3481, df = 1, p-value = 0.003861
> chisq.test(expectOther, correct = FALSE)
       Pearson's Chi-squared test
data: expectOther
X-squared = 39.757, df = 1, p-value = 2.877e-10
Q2:
> grade_f = as.factor(grade)
> sex_f = as.factor(sex)
> mod2.1 = glm(like ~ grade_f * sex_f, family = binomial)
> summary(mod2.1)
Call:
glm(formula = like ~ grade_f * sex_f, family = binomial)
Deviance Residuals:
             1Q Median
                               3Q
                                       Max
-1.8930 -1.1114 0.6039 1.2449
                                   1.2530
```

Fisher's Exact Test for Count Data

```
Coefficients:
                Estimate Std. Error z value Pr(>|z|)
                 (Intercept)
grade_f1
                 -0.0185
                            0.3030 -0.061
                                             0.951
sex_fMale
                  grade_f1:sex_fMale -0.5231
                           0.5297 -0.987 0.323
Signif. codes: 0 '***, 0.001 '**, 0.01 '*, 0.05 '., 0.1 ', 1
(Dispersion parameter for binomial family taken to be 1)
   Null deviance: 539.70 on 398 degrees of freedom
Residual deviance: 488.41 on 395 degrees of freedom
AIC: 496.41
Number of Fisher Scoring iterations: 4
> 1 - pchisq(0.96, 1) # p-value
[1] 0.3271869
> mod2.2 = glm(like ~ grade_f + sex_f, family = binomial)
> summary(mod2.2)
Call:
glm(formula = like ~ grade_f + sex_f, family = binomial)
Deviance Residuals:
       1Q Median
                             3Q
   Min
                                    Max
-1.8412 -1.1273 0.6369 1.2283
                                1.3098
Coefficients:
          Estimate Std. Error z value Pr(>|z|)
(Intercept) -0.1189
                   0.1397 -0.851 0.395
                      0.2519 -0.743
                                     0.458
grade_f1
           -0.1871
sex_fMale
           1.6111
                      0.2438 6.610 3.85e-11 ***
Signif. codes: 0 '*** 0.001 '** 0.01 '* 0.05 '.' 0.1 ' 1
(Dispersion parameter for binomial family taken to be 1)
   Null deviance: 539.70 on 398 degrees of freedom
Residual deviance: 489.37 on 396 degrees of freedom
AIC: 495.37
Number of Fisher Scoring iterations: 4
> 1-pchisq(0.9775,1)
[1] 0.3228168
> n = 399
> count = rep(1,n)
> new = aggregate(count ~ grade + sex + like,data=a3data, FUN = sum)
```

> new

```
grade
        sex like count
   0 Female 0 103
2
    1 Female 0 31
    0 Male 0 18
3
4
    1 Male 0 11
   0 Female 1 88
   1 Female 1 26
6
7
    0 Male 1 90
    1 Male 1 32
> grade_p = as.factor(new$grade)
> sex_p = as.factor(new$sex)
> like_p = as.factor(new$like)
> mod3.1 = glm(counts ~ grade_p*sex_p*like_p, family = poisson)
> summary(mod3.1)
Call:
glm(formula = counts ~ grade_p * sex_p * like_p, family = poisson)
Deviance Residuals:
[1] 0 0 0 0 0 0 0 0
Coefficients:
                     Estimate Std. Error z value Pr(>|z|)
                     (Intercept)
grade_p1
                     sex_pMale
                    -1.74436 0.25547 -6.828 8.61e-12 ***
                    -0.15739 0.14516 -1.084 0.278
like_p1
                    0.70827 0.43409 1.632 0.103
grade_p1:sex_pMale
                     -0.01850 0.30297 -0.061 0.951
grade_p1:like_p1
                sex_pMale:like_p1
Signif. codes: 0 '***, 0.001 '**, 0.01 '*, 0.05 '., 0.1 ', 1
(Dispersion parameter for poisson family taken to be 1)
   Null deviance: 1.9388e+02 on 7 degrees of freedom
Residual deviance: -1.4655e-14 on 0 degrees of freedom
AIC: 59.808
Number of Fisher Scoring iterations: 3
> mod3.2 = glm(counts ~ grade_p + sex_p + like_p + grade_p:sex_p + grade_p:like_p + like_p:sex_p, famil
> summary(mod3.2)
Call:
glm(formula = counts ~ grade_p + sex_p + like_p + grade_p:sex_p +
   grade_p:like_p + like_p:sex_p, family = poisson)
Deviance Residuals:
                                 5
                                               7
                   3
                          4
                                         6
    1
           2
```

```
0.1812 \quad -0.3220 \quad -0.4170 \quad 0.5849 \quad -0.1935 \quad 0.3672 \quad 0.1940 \quad -0.3171
```

Coefficients:

Estimate Std. Error z value Pr(>|z|)(Intercept) 4.61683 0.09773 47.241 < 2e-16 *** grade_p1 -1.12555 0.18653 -6.034 1.60e-09 *** -1.62975 0.21888 -7.446 9.64e-14 *** sex_pMale -0.11893 0.13968 -0.851 0.395 like_p1 grade_p1:sex_pMale 0.35467 0.25229 1.406 0.160 grade_p1:like_p1 -0.18713 0.25189 -0.743 0.458 sex_pMale:like_p1 1.61115 0.24375 6.610 3.85e-11 ***

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

(Dispersion parameter for poisson family taken to be 1)

Null deviance: 193.87673 on 7 degrees of freedom Residual deviance: 0.96302 on 1 degrees of freedom

AIC: 58.771

Number of Fisher Scoring iterations: 4