# **Analysis on Student Gaming Preferences Data**

#### Introduction

With gaming becoming an increasingly popular past time activity for students around the world, it is no surprise that we would be interested in studying what factors affect a student's gaming preferences. More specifically, in this study, I hope to answer 2 questions: Does a student's sex plays a role in their preference for games? Does the relationship between a student's preference for games and their sex change depending on their expected grades. In other words, is there some sort of interaction effect going on between a student's expected grade and other variables which might affect their preferences for gaming.

# **Preliminary Analysis**

	Sex	
Game Preference	Male	Female
Likes Games	122	114
Doesn't Like Games	29	134

From the table above, we can tell that the proportion of students who enjoy playing games that are male is  $P_{male} = 122/236$ , and for females that proportion is  $P_{female} = 114/236$ . Similarly, the proportion of male and female students who don't enjoy playing games are  $q_{male} = 29/163$  and  $q_{female} = 134/163$ . Furthermore, using the difference of proportions test, we arrive at a p-value of  $6.704 \times 10^{-12}$ . And using the Fisher test, we arrived at a p-value of  $2.515\times 10^{-12}$ . It should be noted that the chi-square test's p-value was not used since it is the same as the one in the proportions test. Given these 2 p-values at a 10% significance level, we conclude that there is strong evidence that a student's sexual orientation is dependent on their preference on games.

Below are contingency tables for the two grade types:

Grade			the second secon		
	Cov			Sex	
Game Preference	Sex	Female	Game Preference	Male	Female
Likes Games	32	26	Likes Games	90	88
Doesn't Like Games	11	31	Doesn't Like Games	18	103
A+			Grade of NOT A+		

### **Methods**

C ... 1.

For this particular dataset, it contains 399 different observation. The data was collected through a student survey conducted in the class of STA303 by Professor Shivon Sue Chee. In addition, logistic models and Poisson models were fitted to the data in R. Both these models were used in order to show two different approaches of answering the study question. A logistic model was chosen as the data consisted of a discrete categorical variable. It should also be noted that the Poisson model was fitted to data which

consisted of the counts of students who liked and did not like video games (The data used can be found in the *Appendix*).

## Logistic Models:

Let X= grade, W = sex. Where X represents the 2 levels of the variable 'grade' (i.e. 1= student's grade is A+, and 0 = student's grade is not A+), and W represents the 2 levels of the variable 'sex' (i.e. 1= male, 0 = Female). In addition,  $\pi_i$ , represents the probability of a student liking to play video games.

**Model 2.1:** logit(
$$\pi_i$$
) =  $\beta_0 + \beta_1 X + \beta_2 W + \beta_3 XW$  for i = 1,...,399  
= -0.1574 + 1.7668X - 0.0185W - 0.5231XW

**Model 2.2:** logit(
$$\pi_i$$
) =  $\alpha_0 + \alpha_1 X + \alpha_2 W$  for i = 1,...,399  
= -0.1189 + 1.6111X - 0.1871W

#### Poisson Model:

Let X represent variable 'like' (X= 1 for likes game, X=0 otherwise). Let W represent the variable 'sex' (W=1 for male, W=0 otherwise). Let Z represent the variable 'grade' (Z=1 for grade of A+, Z=0 otherwise). Let

$$\begin{aligned} \textbf{Model 3.1:} \ \log(\mu_{ijk}) &= \beta_0 + \beta_1 X + \beta_2 W + \beta_3 Z + \beta_4 X W + \beta_5 X Z + \beta_6 W Z + \beta_7 X W Z \\ &= 3.4340 - 0.1759 X - 1.0361 W + 1.2007 Z + 1.2437 X W + 0.0185 X Z - 0.7083 W Z + 0.5231 X W Z \end{aligned}$$

$$\begin{aligned} \textbf{Model 3.2:} & \log(\mu_{ijk}) = \beta_0 + \beta_1 X + \beta_2 W + \beta_3 Z + \beta_4 XW + \beta_5 XZ + \beta_6 WZ \\ & = 3.413 - 0.361X - 1.2751W + 1.1256Z + 1.6111XW + 0.1871XZ - 0.3547WZ \end{aligned}$$

#### **Model Selection**

In order to choose the best logistic model, I performed both a Likelihood Ratio Test and a Wald's Test on models 2.1 and 2.2. The results from the tests can be seen below:

**Test 1:** The first test is the Likelyhood Ratio Test (LRT).

Null hypothesis (H <sub>0</sub> ):	The reduced model (Model 2.2) is appropriate	
Alternative hypothesis	The full model (Model 2.1) is better	
$(\mathbf{H}_{\mathbf{A}})$ :		
<b>Test Statistics:</b>	$G^2 = 489.37 - 488.41 = 0.96$	
	Following a chi-square distribution with 1 degree of freedom	
P-Value:	0.3272 (see appendix for code)	
1 (title)	(50272 (500 upponum 151 5500)	
Conclusions:	Since we are using a significance level of 0.10, we would fail to reject the null hypothesis since our p-value of 0.3272 is greater than our significance level. In other words, the reduced model (Model 2.2) is an appropriate fit for our data.	

**Test 2:** The second test is a Wald's test

Null hypothesis (H <sub>o</sub> ):	β <sub>3</sub> =0
Alternative hypothesis	$\beta_3 \neq 0$
(H <sub>A</sub> ):	
Test Statistics:	$Z^2 = 0.9741$
	Following a chi-square distribution with 1 degree of freedom
P-Value:	0.3230
Conclusions:	Since we are using a significance level of 0.10, we would still fail to reject the null hypothesis since our p-value of 0.3236 is greater than our significance level. In other words, we have enough evidence saying that grade and sex don't have an effect on the odds of liking games. Thus we should stick to model 2.2.

Regarding the Poisson models, I chose to perform just the Wald's Test on the 2 models. The results indicated that model 3.2 (a reduced model) was the better choice for this study.

## **Conclusion**

In conclusion, from the fitted logistic models above, we see that there is no significant interaction effect between a student's sex and expected grades on their preference for playing video games; this was previously shown by the high p-value of 0.3272 which indicates a lack of an interaction effect. Similarly, the fitted Poisson models also appears to show no signs of an interaction effect between the student's sex and grades on gaming preference. However, when looking at the results from both of the chosen reduced models (i.e. Model 2.2 & Model 3.2), it would appear that a student's sex plays a crucial part in their preference for gaming. That is, if a student is male, there are higher odds of that student preferring to game in contrast to a female student.

# **Appendix**

#### Pre-code:

```
studData <- read.csv("a3data.csv")</pre>
 apSubset <- subset(studData, grade == "1")
 notapSubset <- subset(studData, Grade == "A" | Grade == "B" | Grade == "C")
 Below is code for the 2x2 table:
 gamerMale <- subset(studData, like == "1" & sex == "Male")
 gamerFemale <- subset(studData, like == "1" & sex == "Female")
 notGamerMale <- subset(studData, like == "0" & sex == "Male")
 notGamerFemale <- subset(studData, like == "0" & sex == "Female")
 contTable <- matrix(c(nrow(gamerMale),nrow(gamerFemale), nrow(notGamerMale),
 nrow(notGamerFemale)), nrow= 2, byrow = TRUE)
 dimnames(contTable) <- list(c("Likes Games", "Doesn't Like Games"), c("Male", "Female"))
 names(dimnames(contTable)) <- c("Game Preference", "Sex")</pre>
 Code below for proportion and fisher tests
 diffPropTest <- prop.test(contTable, correct = FALSE</pre>
> diffPropTest
         2-sample test for equality of proportions without continuity correction
data: contTable
X-squared = 47.112, df = 1, p-value = 6.704e-12
alternative hypothesis: two.sided
95 percent confidence interval:
 0.2523654 0.4257047
sample estimates:
   prop 1 prop 2
0.5169492 0.1779141
 fisherTest <- fisher.test(contTable)</pre>
```

#### > fisherTest

```
Fisher's Exact Test for Count Data
data: contTable
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
    Logistic Models
    like <- studData$like
    sex <- studData$sex</pre>
    grades <- studData$grade
    model2.1 <- glm(like ~ sex * grades, family = binomial, data = studData)
    model2.2 <- glm(like ~ sex + grades, family = binomial, data = studData)
    summary1 <- summary(model2.1)</pre>
    > summary1
    glm(formula = like ~ sex * grades, family = binomial, data = studData)
    Deviance Residuals:
        Min
              1Q
                      Median
                                      3Q
                                              Max
    -1.8930 -1.1114
                        0.6039
                                 1.2449
                                           1.2530
    Coefficients:
                    Estimate Std. Error z value Pr(>|z|)
    (Intercept)
                     -0.1574
                                 0.1452 -1.084
                                                    0.278
                                          5.965 2.45e-09 ***
    sexMale
                      1.7668
                                 0.2962
                                          -0.061
                     -0.0185
                                 0.3030
                                                    0.951
    grades
                                 0.5297 -0.987
    sexMale:grades -0.5231
                                                    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
```

```
> pchisq(0.96, 1, lower.tail = FALSE)
[1] 0.3271869
summary2 <- summary(model2.2)</pre>
> summary2
Call:
glm(formula = like \sim sex + grades, family = binomial, data = studData)
Deviance Residuals:
    Min 1Q Median 3Q
                                        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
sexMale 1.6111
grades -0.1871
                         0.2438 6.610 3.85e-11 ***
                         0.2519 -0.743 0.458
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
Poisson Models
Generating Count data to feed into poisson model
count <- c(31,103,11,18,26,88,32,90)
like1 <- as.factor(c("no", "no", "no", "no", "yes", "yes", "yes", "yes"))
sex1 <- as.factor(c("female", "female", "male", "female", "female", "female", "male", "male"))
```

grade1 <- as.factor(c("A+", "not A+", "A+", "not A+", "A+", "not A+", "A+", "not A+"))

pModel1 <- glm(count ~ like1 \* sex1 \* grade1, family = poisson) pModel2 <- glm(count ~ (like1 + sex1 + grade1)^ 2, family = poisson)

```
> summary(pModel1)
 glm(formula = count ~ like1 * sex1 * grade1, family = poisson)
 Deviance Residuals:
 [1] 0 0 0 0 0 0 0 0
 Coefficients:
                              Estimate Std. Error z value Pr(>|z|)
 (Intercept)
                                3.4340
                                          0.1796 19.120 < 2e-16 ***
                                                 -0.661 0.50835
 like1yes
                               -0.1759
                                          0.2659
 sex1male
                               -1.0361
                                          0.3509 -2.952 0.00315 **
                                1.2007
                                                   5.861 4.59e-09 ***
 grade1not A+
                                          0.2049
                                1.2437
                                          0.4392
                                                   2.832 0.00463 **
 likelyes:sex1male
 likelyes:gradelnot A+
                                0.0185
                                          0.3030
                                                   0.061
                                                         0.95131
 sex1male:grade1not A+
                               -0.7083
                                          0.4341
                                                 -1.632
                                                         0.10276
 likelyes:sex1male:grade1not A+
                              0.5231
                                          0.5297
                                                   0.987
                                                         0.32341
 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: 4.6629e-15 on 0 degrees of freedom
 AIC: 59.808
> summary(pModel2)
Call:
glm(formula = count \sim (like1 + sex1 + grade1)^2, family = poisson)
Deviance Residuals:
                          3
                                             5
                                                       6
      1
                    0.5849 -0.4170
                                        0.3672 -0.1935 -0.3171
-0.3220
           0.1812
                                                                     0.1940
Coefficients:
                       Estimate Std. Error z value Pr(>|z|)
                                             21.131 < 2e-16 ***
(Intercept)
                          3.4913
                                     0.1652
like1yes
                         -0.3061
                                      0.2329
                                              -1.314
                                                         0.189
                                              -4.715 2.42e-06 ***
sex1male
                         -1.2751
                                      0.2704
                         1.1256
                                     0.1865
                                               6.034 1.60e-09 ***
grade1not A+
likelyes:sex1male
                         1.6111
                                      0.2438
                                               6.610 3.85e-11 ***
                                     0.2519
                                               0.743
                                                         0.458
likelyes:gradelnot A+
                         0.1871
sex1male:grade1not A+
                        -0.3547
                                     0.2523 - 1.406
                                                         0.160
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
                      0.96302 on 1
Residual deviance:
                                       degrees of freedom
AIC: 58.771
```