ASSIGNEMNT 3 - DAVID VEITCH 1004657917

1. Selected Field of Interest

Education.

2. Reference to the article

Dhuey, E. & Smith, J (2018). How school principals influence student learning. *Empir Econ* (2018) 54: 851. https://doiorg.myaccess.library.utoronto.ca/10.1007/s00181-017-1259-9

3. UofT Department

Elizabeth Dhuey – Department of Management – University of Toronto Scarborough

4. Soft-Copy

https://doiorg.myaccess.library.utoronto.ca/10.1007/s00181-017-1259-9 http://rdcu.be/JymP

5. Statistical Software

The author does not specify which statistical software was used.

6. Data Source

The data comes from an observational study, namely "administrative records of the North Carolina Department of Public Instruction" which includes data on schools, teachers, students, and principals. The authors also used census data in this study. The authors have a cleaned dataset of 5,388,543 observations of student test scores between 1998 and 2010.

7. Summary Statistics/Plots

Yes the article presents summary statistics in two ways. First it presents summary descriptive statistics of the data used (e.g. student test scores) and then it presents tables of its results (i.e. principal effect on student achievement). Two examples below:

now	school	principals	innuence	student	icaming

Table 7 Relationship between principal quality and education

Table 1 Descriptive statistics		.,	en.		Math	Reading
for students in analysis sample		Mean	SD		(1)	(2)
	Math scores			Panel A: Graduate degrees		
	Grade 3	0.067	0.960		0.0034	0.0109**
	Grade 4	0.038	0.986	Graduate degree (doctorate or advanced certificate)		
	Grade 5	0.041	0.986		(0.0059)	(0.0044)
	Grade 6	0.054	0.983	Adjusted R-squared	0.0001	0.0014
	Grade 7	0.055	0.986	Panel B: Doctorate or advanced certificate		
	Grade 8	0.052	0.984	Doctorate	0.0094	0.0068
	Reading scores				(0.0091)	(0.0069)
	Grade 3	0.063	0.958	Advanced certificate	-0.0005	0.0120**
	Grade 4	0.029	0.987		(0.0063)	(0.0047)
	Grade 5	0.032	0.983	Adjusted R-squared	0.0002	0.0021
	Grade 6	0.047	0.975	Panel C: Competitive or non-competitive		
	Grade 7	0.048	0.975	Competitive master's school	-0.0061	0.0016
	Grade 8	0.046	0.973		(0.0083)	(0.0058)
	Student demographic characteristics			Competitive doctorate school	-0.0169	-0.0089
	Male	0.501	0.500		(0.0177)	(0.0128)
	Black	0.286	0.452	Non-competitive doctorate school	0.0201**	0.0122
	Hispanic	0.054	0.227		(0.0102)	(0.0079)
	White	0.606	0.489	Competitive advanced certificate school	0.0062	0.0220**
	Other race	0.053	0.224		(0.0131)	(0.0107)
Test scores are standardized to	Special education	0.092	0.289	Non-competitive advanced certificate school	-0.0024	0.0103**
have a mean of zero and	Gifted	0.140	0.347		(0.0068)	(0.0051)
standard deviation of 1 in the	Learning disabled in math	0.020	0.140	Adjusted R-squared	0.0012	0.0026
population of test takers within a subject, grade, and year, prior to	Learning disabled in reading	0.041	0.199	Number of observations	4289	4289
sample exclusions. Other race	Number of students	1,664,158		Education variables are indicator variables equal to 1 if the pr		Ct 1 1
includes all races except the	Number of observations	5,388,543		errors are robust to heteroskedasticity	merpar ever nau the particul	ii degree. Standare

*** p < 0.01; ** p < 0.05; * p < 0.1

8. Test Statistics/P-Values/Confidence Intervals

The authors conduct a regression analysis of a number of factors principals have versus student achievement. In presenting the results as shown above the authors highlight factors which are statistically significant (having a P-Value less than 0.10, 0.05, and 0.01).

9. How many decimal places were values recorded.

Regression coefficients were reported to four decimal places. Summary statistics (e.g. mean/standard deviation) were reported to three decimal places.

10. Statistical Method to Analyze Data

The authors use ordinary least squares regression to estimate the effect principals have on students. Description of the model below.

4 Empirical specifications

4.1 Estimating principal effects

To estimate the principal effects, we use the following value-added model of students' test scores:

$$y_{it} = \beta_0 + y_{it-1}\beta_1 + x'_{it}\beta_2 + z'_{s(i,t)t}\beta_3 + p'_{p(i,t)t}\beta_4 + \delta_{p(i,t)} + \varphi_{s(i,t)} + \eta_t + \varepsilon_{it},$$
(1)

where y_{it} is the math or reading score for student i at time t; y_{it-1} is the student's 1-year lagged math or reading score; x'_{it} is a vector of student-level demographic

⁴ While principals possess the power of evaluating teachers, the superintendent or the board of directors (for regional schools) have the power to employ and contract with teachers along with the duty to maintain personnel files and participate in the firing and demoting.



856 E. Dhuey, J. Smith

characteristics; $z'_{s(i,t)t}$ is a vector of school-level demographic characteristics for the school that student i attends at time t; $p'_{p(i,t)t}$ is a vector of principal-level, time-changing characteristics for student i's principal at time t; $\delta_{p(i,t)}$, $\varphi_{s(i,t)}$, and η_t are time-invariant principal, school, and year effects; and ε_{it} is an idiosyncratic error term.

PART 2 CONTINGNEGY TABLES

1a.

Contingency table: sex, like

There is evidence that sex is not independent of a student's preference for playing video games. This is evident from the p-values obtained from a difference of proportions test (p-value 0.06797) and the Fisher test (p-value 0.07824). These p-values are below the 10% level of significance suggesting there is evidence sex is not independent of preference for playing video games.

In practical terms this means if a given subject is male they are much more likely to like playing video games than a female. In this case 84% of males liked playing video games compared to only 68% of females).

1b.

It does appear that the sex and like relationship changes with grade expected.

For students expecting an A in a statistics course there appears to be a relationship between sex and preference for playing video games (males much more likely to like playing video games). This can be seen from the difference of proportion test (p-value 0.001102) and Fisher test (p-value 0.003879). This relationship does not seem to exist for students not expecting an A; the difference of proportion tests (p-value 0.9421) and Fisher test (p-value 1) appear to suggest no relationship.

2a.

We fit two models to the data, Model 2_1 with an interaction term, and Model 2_2 without interaction.

$$\begin{aligned} & \text{Model 2_1} - \textbf{logit(like)} = \beta_{intercept} + I_{male} * \beta_{sexmale} + I_{gradenA} * \beta_{gradenA} + I_{sexmale:gradenA} * \beta_{sexmale:gradenA} \\ & \text{Model 2_2} - \textbf{logit(like)} = \beta_{intercept} + I_{male} * \beta_{sexmale} + I_{gradenA} * \beta_{gradenA} \\ \end{aligned}$$

Model Term	Description
logit(like)	predicted log odds that individual likes video games
$\beta_{intercept}$	predicted log odds for sex = female, grade = A
$\beta_{sexmale}$	change in log odds if sex = male
$\beta_{gradenA}$	change in log odds if $grade = nA$
$\beta_{sexmale:gradenA}$	change in log odds if $sex = male$ and $grade = nA$
I_{male}	indicator variable, 1 if $sex = male$, 0 if $sex = female$, 0 otherwise
$I_{gradenA}$	indicator variable, 1 if grade = nA , 0 if grade = A , 0 otherwise
I _{sexmale:gradenA}	indicator variable, 1 if grade = nA and $sex = male$, 0 otherwise

Model 2_1, the model with interaction, appears to be the better fit the data.

First we conduct a likelihood ratio test between the two models, comparing their residual deviance. From a chi-square distribution of one degree of freedom (the models only differ by one parameter) we get a p-value of 0.0087 suggesting the model with interaction is a better fit.

Second we conduct a Wald chi-square test on the $\beta_{sexmale:gradenA}$ term in Model 2_1. Observing the p-value (0.01848) from the summary output of the logistic regression model we can conclude that the model with $\beta_{sexmale:gradenA}$ better fits the data.

2b.

The model selected in part 2a), the model with interaction, agrees with our analysis in question 1b). The practical implications of this is that when modelling whether a student likes video games one must take into account that males with high grade expectations have a different preference for video games than males with low expectations, and females with high grade expectations have a different preference for video games than females with low expectations.. Therefore to properly model this relationship one should have an interaction term in the model.

3a.

$$\begin{aligned} &\text{Model 3_1} - \textbf{log(Freq)} = \beta_{intercept} + I_{male} * \beta_{sexmale} + I_{gradenA} * \beta_{gradenA} + I_{likeyes} * \beta_{likeyes} + I_{sexmale:gradenA} \\ &\beta_{sexmale:gradenA} + I_{sexmale:likeyes} * \beta_{sexmale:likeyes} + I_{gradenA:likeyes} * \beta_{gradenA:likeyes} + I_{sexmale:gradenA:likeyes} * \beta_{sexmale:gradenA:likeyes} \\ &\text{Model 3_2} - \textbf{log(Freq)} = \beta_{intercept} + I_{male} * \beta_{sexmale} + I_{gradenA} * \beta_{gradenA} + I_{likeyes} * \beta_{likeyes} + I_{sexmale:gradenA} \\ &\beta_{sexmale:gradenA} + I_{sexmale:likeyes} * \beta_{sexmale:likeyes} + I_{gradenA:likeyes} * \beta_{gradenA:likeyes} \\ \end{aligned}$$

Model Term	Description
log(Freq)	predicted log(Freq) where Freq represents the number of individuals in a category
$\beta_{intercept}$	predicted frequency for sex = female, grade = A, like = no
$\beta_{sexmale}$	change in log(Freq) if sex = male
$\beta_{gradenA}$	change in $log(Freq)$ if grade = nA
$\beta_{likeyes}$	change in log(Freq) if like = yes
$\beta_{sexmale:gradenA}$	change in $log(Freq)$ if $sex = male$ and $grade = nA$
$\beta_{sexmale:likeyes}$	change in log(Freq) if sex = male and like = yes
$\beta_{gradenA:likeyes}$	change in log(Freq) if grade = nA and like = yes
$\beta_{sexmale:gradenA:likeyes}$	change in log(Freq) if sex = male and grade = nA and like = yes
I_{male}	indicator variable, 1 if sex = male, 0 if sex = female
$I_{gradenA}$	indicator variable, 1 if grade = nA, 0 if grade = A
$I_{likeyes}$	indicator variable, 1 if like = yes, 0 if like = no
I _{sexmale:gradenA}	indicator variable, 1 if grade = nA and $sex = male$, 0 otherwise
I _{sexmale:likeyes}	indicator variable, 1 if sex = male and like = yes, 0 otherwise
I _{gradenA:likeyes}	indicator variable, 1 if grade = nA and like = yes, 0 otherwise
I _{sexmale:gradenA:likeyes}	indicator variable, 1 if sex = male and grade = nA and like = yes, 0 otherwise

Deviance

Model	Residual Deviance
2_1	85.152
2_2	92.031
3_1	0
3_2	6.8788

The deviance of Model 3_1 is 0. This is unsurprising given it is a saturated model and has one predictor variable for every category so it perfectly fits the data. Model 3_2 has a much lower residual deviance than model 2_1 and 2_2 which is unsurprising given Model 3_2 has seven β predictors (including intercept) as opposed to 2_1 which has four β predictors and 2_2 with three β predictors.

Wald Tests

The p-values of all non-intercept predictors in 2_1 are statistically significant (p-value below 0.1), while in 2_2 only one of two predictors is statistically significant ($\beta_{sexmale}$).

This contrasts with 3_1 and 3_2 which have a number of statistically insignificant predictors (according to a Wald test). Only three of seven predictors in 3_1 have p-values below 0.1, and only three of six predictors in 3_2 have p-values below 0.1. Given this, it may be worthwhile to drop some insignificant predictors from 3_1 and 3_2 in order to simplify the model and improve interpretability.

Interpretation

In interpreting model 2_1 and 2_2 we are able to see that there is a definite change in the relationship between sex and liking video games when grade is taken into account.

Comparing model 3_1 to 2_1 (the models with the lowest residuals for each type of regression) in terms of residual deviance we see that model 3_1 is statistically significantly (p-value of 0 on a deviance goodness of fit test) more accurate. Interpreting this would seem to suggest that using a three-way interaction is a superior way to model the relationship between grade, likes video games, and sex. This suggests there are many interactions occurring at once, and it would be too simplistic to use a model with only one interaction (such as model 2_1).

R CODE APPENDIX - Assn3R.R

David

Thu Mar 22 11:10:18 2018

```
# Assignment 3 David Veitch 1004657917
workdir<-("C:/Users/David/Google Drive/Documents/UofT/NonDegree/303/Assignmen</pre>
t3")
setwd(workdir)
dat <- read.csv("video.csv")</pre>
# 1A
# Create contingency table
cont_tab <- xtabs(~like+sex, data=dat)</pre>
cont_tab
##
        sex
## like female male
             12
     no
##
             26
                  44
     yes
# Run difference of proportions test and chi-sq test
prop.test(cont_tab, correct=FALSE)
##
  2-sample test for equality of proportions without continuity
##
## correction
##
## data: cont tab
## X-squared = 3.3314, df = 1, p-value = 0.06797
## alternative hypothesis: two.sided
## 95 percent confidence interval:
## -0.01414205 0.47128491
## sample estimates:
      prop 1 prop 2
## 0.6000000 0.3714286
fisher.test(cont_tab)
##
   Fisher's Exact Test for Count Data
##
##
## data: cont_tab
## p-value = 0.07824
## alternative hypothesis: true odds ratio is not equal to 1
## 95 percent confidence interval:
## 0.8195672 8.1070182
## sample estimates:
```

```
## odds ratio
##
     2.511179
# 1B
# Create contingency tables for different grades
# Contingency tables & testsfor grade == A
cont_tab_grade_A <- xtabs(~like+sex, data=subset(dat, grade=="A"))</pre>
cont tab grade A
##
        sex
## like female male
     no
                  21
##
              4
    yes
prop.test(cont tab grade A, correct=FALSE)
## Warning in prop.test(cont_tab_grade_A, correct = FALSE): Chi-squared
## approximation may be incorrect
##
## 2-sample test for equality of proportions without continuity
## correction
##
## data: cont tab grade A
## X-squared = 10.648, df = 1, p-value = 0.001102
## alternative hypothesis: two.sided
## 95 percent confidence interval:
## 0.3423132 1.0000000
## sample estimates:
##
      prop 1
                prop 2
## 0.8333333 0.1600000
fisher.test(cont_tab_grade_A)
##
## Fisher's Exact Test for Count Data
##
## data: cont_tab_grade_A
## p-value = 0.003879
## alternative hypothesis: true odds ratio is not equal to 1
## 95 percent confidence interval:
       1.877541 1284.874786
## sample estimates:
## odds ratio
     22.44903
##
# Contingency tables & tests for grade == nA
cont_tab_grade_nA <- xtabs(~like+sex, data=subset(dat, grade=="nA"))</pre>
cont_tab_grade_nA
```

```
sex
## like female male
              7
##
                   7
     no
             22
##
                  23
     yes
prop.test(cont_tab_grade_nA, correct=FALSE)
##
## 2-sample test for equality of proportions without continuity
## correction
##
## data: cont_tab_grade_nA
## X-squared = 0.0052746, df = 1, p-value = 0.9421
## alternative hypothesis: two.sided
## 95 percent confidence interval:
## -0.2887695 0.3109917
## sample estimates:
##
      prop 1
                prop 2
## 0.5000000 0.4888889
fisher.test(cont_tab_grade_nA)
##
## Fisher's Exact Test for Count Data
## data: cont_tab_grade_nA
## p-value = 1
## alternative hypothesis: true odds ratio is not equal to 1
## 95 percent confidence interval:
## 0.2637887 4.1391671
## sample estimates:
## odds ratio
     1.044688
##
# 2 A
# Create logistic regression models
# Model 2.1 to include interaction between sex and grade
model_2_1 <- glm(like~sex+grade+sex*grade,family=binomial, data=dat)</pre>
summary(model_2_1)
##
## Call:
## glm(formula = like ~ sex + grade + sex * grade, family = binomial,
       data = dat)
##
## Deviance Residuals:
       Min
                      Median
##
                 10
                                   3Q
                                            Max
## -2.4864
             0.3050
                      0.7290
                               0.7433
                                        1.2735
##
```

```
## Coefficients:
##
                   Estimate Std. Error z value Pr(>|z|)
## (Intercept)
                    -0.2231
                                0.6708
                                       -0.333
                                                0.73940
                     3.2677
                                               0.00758 **
## sexmale
                                1.2237
                                         2.670
## gradenA
                     1.3683
                                0.7989
                                         1.713
                                                0.08679 .
## sexmale:gradenA -3.2232
                                1.3682 -2.356 0.01848 *
## Signif. codes: 0 '***' 0.001 '**' 0.05 '.' 0.1 ' ' 1
## (Dispersion parameter for binomial family taken to be 1)
##
##
       Null deviance: 95.347 on 89 degrees of freedom
## Residual deviance: 85.152 on 86 degrees of freedom
## AIC: 93.152
##
## Number of Fisher Scoring iterations: 5
# Model 2.2 to not include interaction between sex and grade
model_2_2 <- glm(like~sex+grade,family=binomial, data=dat)</pre>
summary(model_2_2)
##
## Call:
## glm(formula = like ~ sex + grade, family = binomial, data = dat)
##
## Deviance Residuals:
##
       Min
                 1Q
                      Median
                                   3Q
                                           Max
                                        0.8774
## -1.9533
           0.5668
                      0.5861
                               0.8512
##
## Coefficients:
               Estimate Std. Error z value Pr(>|z|)
##
## (Intercept) 0.8288
                            0.5586
                                     1.484
                                             0.1379
                                     1.736
                            0.5291
                                             0.0826 .
## sexmale
                 0.9183
## gradenA
                -0.0727
                            0.5679 -0.128
                                             0.8981
## ---
## Signif. codes: 0 '***' 0.001 '**' 0.01 '*' 0.05 '.' 0.1 ' ' 1
## (Dispersion parameter for binomial family taken to be 1)
##
       Null deviance: 95.347 on 89 degrees of freedom
##
## Residual deviance: 92.031 on 87 degrees of freedom
## AIC: 98.031
##
## Number of Fisher Scoring iterations: 4
# Conduct Likelihood Ratio test using residual deviance
# Test is chi-squre with one degree of freedom
p_val = (1-pchisq(model_2_2$deviance - model_2_1$deviance,1))
p_val
## [1] 0.008722588
```

```
# 3 A
# Create a 3-way contingency table and convert to a dataframe
cont_tab_3_way <- xtabs(~like+sex+grade, data=dat)</pre>
df_3_way <- as.data.frame(cont_tab_3_way)</pre>
df_3_way
##
     like
             sex grade Freq
## 1
       no female
                     Α
                          5
## 2 yes female
                     Α
                          4
                     Α
## 3
            male
                          1
       no
## 4 yes
            male
                     Α
                         21
## 5
      no female
                    nΑ
                         7
## 6 yes female
                    nΑ
                         22
## 7
            male
                    nΑ
                          7
      no
                         23
## 8 yes
            male
                    nΑ
# Create models with counts as Poisson variables
# Model 3.1 three two-way interaction terms and one three-way interaction
model 3 1 <- glm(Freq~sex+grade+like+sex*grade*like,family=poisson, data=df 3
_way)
summary(model_3_1)
##
## Call:
## glm(formula = Freq ~ sex + grade + like + sex * grade * like,
##
       family = poisson, data = df_3_way)
##
## Deviance Residuals:
## [1] 0 0 0 0 0 0 0 0
##
## Coefficients:
##
                           Estimate Std. Error z value Pr(>|z|)
                                        0.4472
                                                 3.599 0.00032 ***
## (Intercept)
                             1.6094
## sexmale
                            -1.6094
                                        1.0954
                                                -1.469 0.14177
## gradenA
                             0.3365
                                        0.5855
                                                 0.575
                                                        0.56554
## likeyes
                            -0.2231
                                        0.6708
                                                -0.333
                                                        0.73940
## sexmale:gradenA
                             1.6094
                                        1.2189
                                                 1.320
                                                        0.18670
## sexmale:likeyes
                                                        0.00758 **
                             3.2677
                                        1.2238
                                                 2.670
## gradenA:likeyes
                                        0.7989
                                                 1.713
                             1.3683
                                                         0.08679
## sexmale:gradenA:likeyes -3.2232
                                        1.3683
                                                -2.356 0.01849 *
## ---
## Signif. codes: 0 '***' 0.001 '**' 0.05 '.' 0.1 ' ' 1
##
## (Dispersion parameter for poisson family taken to be 1)
##
       Null deviance: 5.4112e+01 on 7
                                         degrees of freedom
## Residual deviance: -5.7732e-15 on 0 degrees of freedom
## AIC: 47.169
```

```
##
## Number of Fisher Scoring iterations: 3
# Model 3.2 three two-way interaction terms, no three-way interaction
model 3 2 <- glm(Freq~sex+grade+like+ sex:grade + sex:like + grade:like,famil</pre>
y=poisson, data=df_3_way)
summary(model_3_2)
##
## Call:
## glm(formula = Freq ~ sex + grade + like + sex:grade + sex:like +
       grade:like, family = poisson, data = df_3_way)
##
##
## Deviance Residuals:
                                             5
                           3
                                    4
                                                      6
##
  1.2260
           -0.9698
                    -1.4709
                               0.5132
                                      -0.7780
                                                 0.5005
                                                           0.9711
                                                                   -0.4576
##
## Coefficients:
                   Estimate Std. Error z value Pr(>|z|)
##
## (Intercept)
                     1.0061
                                0.5122
                                         1.964
                                                 0.0495 *
## sexmale
                     0.1771
                                0.5684
                                         0.312
                                                 0.7553
## gradenA
                     1.2201
                                0.5480
                                         2.227
                                                 0.0260 *
## likeyes
                     0.8288
                                0.5586
                                         1.484
                                                 0.1379
                                       -1.761
## sexmale:gradenA -0.8484
                                0.4819
                                                 0.0783 .
## sexmale:likeyes
                     0.9183
                                0.5291
                                         1.736
                                                 0.0826 .
                                0.5679 -0.128
                                                 0.8981
## gradenA:likeyes -0.0727
## ---
## Signif. codes: 0 '***' 0.001 '**' 0.05 '.' 0.1 ' ' 1
## (Dispersion parameter for poisson family taken to be 1)
##
##
       Null deviance: 54.1125 on 7 degrees of freedom
## Residual deviance: 6.8788 on 1 degrees of freedom
## AIC: 52.048
##
## Number of Fisher Scoring iterations: 5
# 3 B
# Compare models 2_2 and 3_1, chisq with 4 df
# Model 3 statistically significantly more accurate
p_val = (1-pchisq(model 2 1$deviance - model 3 1$deviance, 4))
p_val
## [1] 0
```