

Code ▼

W241 Class Project - Analysis

1. Load libraries, data

Load up the data and do simple analysis. This version uses the complete dataset.

Hide

```
# Libraries
library(lmtest)
library(sandwich)
library(ggplot2)
library(data.table)
library(stargazer)
library(ri)
library(multiwayvcov)
library(AER)
rm(list=ls())
d <- read.csv('~/Documents/mids-w241-final/Analysis/Combined Log.csv')
#d <- read.csv("C:/Users/Chris/OneDrive/Documents/MIDS/WS241/final/mids-w241-final/Analysis/Combined Log.csv")
d <- data.table(d)
# d <- d[complete.cases(d),] drops any row that is incomplete - too stringent since we
  have some cols which are not essential
d <- d[!is.na(no)] # Just drop row with missing values
# d <- d[complete.cases(d),] drops any row that is too incomplete - too stringent since
  we have some cols which are not essential
#d <- d[!is.na(no)] # Just drop row with missing values
# Base data
head(d)
```

n	city	title
<int>	<fctr>	<fctr>
6	seattle	Top Floor 2x2 Park & Lake Side/Corner Home!! USB Plugins!!
87	chicago	836 S Bishop St #G
62	houston	Sophisticated Living 2 BR APT with Excellent Amenities-Receive Up To \$
89	chicago	ALL BRAND NEW OUTSTANDING EAST LAKEVIEW LOCATION
96	chicago	Upgrade your lifestyle. Inexpensive with great character!
26	seattle	LUXURY 2 BEDROOM ***2 Master Suites*Coming Soon! Call for Availability

6 rows | 1-4 of 17 columns

Hide

```

# Convert, transform data for analysis
# Drop some cols
d[,c('title','full_URL', 'reply_email_TO_BE_FILLED_IN_standard','posting_ID','notes')
:=NULL]
# Set gender = 1 for Jane
d[treatment_assignment=='Jane_Control' | treatment_assignment=='Jane_Treat_High' | treat
ment_assignment=='Jane_Treat_Low',gender:=1]
d[treatment_assignment=='John_Control' | treatment_assignment=='John_Treat_High' | trea
tment_assignment=='John_Treat_Low',gender:=0]
# Set treatment variable = 0 for control, 1 for low, 2 for high (treatment here is cont
inuous)
d[treatment_assignment=='Jane_Control' | treatment_assignment=='John_Control', treatmen
t:=0]
d[treatment_assignment=='Jane_Treat_Low' | treatment_assignment=='John_Treat_Low', trea
tment:=1]
d[treatment_assignment=='Jane_Treat_High' | treatment_assignment=='John_Treat_High', tr
eatment:=2]
# Alternatively, treat treatment types as categorical variables instead of continuous
d[treatment_assignment=='Jane_Treat_Low' | treatment_assignment=='John_Treat_Low', low_
treatment:=1]
d[treatment_assignment=='Jane_Treat_High' | treatment_assignment=='John_Treat_High', hi
gh_treatment:=1]
d$low_treatment[is.na(d$low_treatment)] <- 0
d$high_treatment[is.na(d$high_treatment)] <- 0
d[low_treatment==1 | high_treatment==0, assigned:=1]
d$assigned[is.na(d$assigned)] <- 0
# Capture complier
#d[sent!='', compliers:=1]
#d$compliers[is.na(d$compliers)] <- 0
# Labeling data
d$gender <- factor(d$gender,labels = c("Male", "Female"))
d$outcome_f <- factor(d$outcome, labels = c("No Response", "Response"))
d$bedrooms <- factor(d$bedrooms, labels = c("1-bedroom", "2-bedroom"))
d$professional <- factor(d$professional, labels = c("Non-professional",
"Professional"))
d$treatment_f <- factor(d$treatment, labels = c("Control","Low","High"))
head(d)

```

n	city	posting_date	bedrooms	sqft	price	treatment_assignment	professional
<int>	<fctr>	<fctr>	<fctr>	<int>	<int>	<fctr>	<fctr>
6	seattle	3/17/2017 13:32	2-bedroom	974	1791	Jane_Treat_High	Professional
87	chicago	3/17/2017 15:22	2-bedroom	NA	2050	Jane_Treat_Low	Non-professional
62	houston	3/17/2017 15:24	2-bedroom	1141	1598	Jane_Treat_High	Professional
89	chicago	3/17/2017 15:27	1-bedroom	1100	1995	Jane_Treat_Low	Professional
96	chicago	3/17/2017 15:26	1-bedroom	NA	1650	Jane_Control	Non-professional
26	seattle	3/17/2017 13:11	2-bedroom	1234	1655	John_Treat_Low	Professional

6 rows | 1-8 of 19 columns

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NA

2. Check data, do simple tables to check for balance

Recode missing sqft values. It's not necessary but we use this in some of our model specifications.

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```
# Recode missing sqft with mean of cluster (city)
d[, Mean:=mean(sqft, na.rm=TRUE), by=city]
d[is.na(sqft)]$sqft <- d[is.na(sqft)]$Mean
```

Coerced 'double' RHS to 'integer' to match the column's type; may have truncated precision. Either change the target column to 'double' first (by creating a new 'double' vector length 483 (nrows of entire table) and assign that; i.e. 'replace' column), or coerce RHS to 'integer' (e.g. 1L, NA_[real|integer]_, as.*, etc) to make your intent clear and for speed. Or, set the column type correctly up front when you create the table and stick to it, please.

Hide

```
d[,c('Mean') :=NULL]
```

Remove duplicate emails

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```
# If duplicate, only retain first email sent, sets up for exploration later
d <- d[duplicate_email == 0]
```

For the most part it looks like we have a balanced dataset.

Hide

```
cat('Table of Outcomes:')
```

Table of Outcomes:

Hide

```
table(d$outcome_f)
```

No Response	Response
256	207

Hide

```
cat('\nTable of Outcomes (By Gender):')
```

Table of Outcomes (By Gender):

Hide

```
table(d$outcome_f, d$gender)
```

	Male	Female
No Response	131	125
Response	100	107

Hide

```
cat('\nTable of Outcomes (By Treatment):')
```

Table of Outcomes (By Treatment):

Hide

```
table(d$outcome_f, d$treatment_f)
```

	Control	Low	High
No Response	87	87	82
Response	68	68	71

Hide

```
cat('\nTable of Outcomes (By Treatment and Gender):')
```

Table of Outcomes (By Treatment and Gender):

Hide

```
table(d$outcome_f, factor(d$treatment_assignment))
```

	Jane_Control	Jane_Treat_High	Jane_Treat_Low	John_Control	John_Treat_High
John_Treat_Low					
No Response	39	38	48	48	44
Response	38	37	32	30	34

Hide

```
cat('\nTable of Outcomes (By City):')
```

Table of Outcomes (By City):

Hide

```
table(d$outcome_f,factor(d$city))
```

	chicago	houston	sandiego	seattle
No Response	63	78	65	50
Response	61	38	51	57

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```
cat('\nTable of Outcomes (By Rooms):')
```

Table of Outcomes (By Rooms):

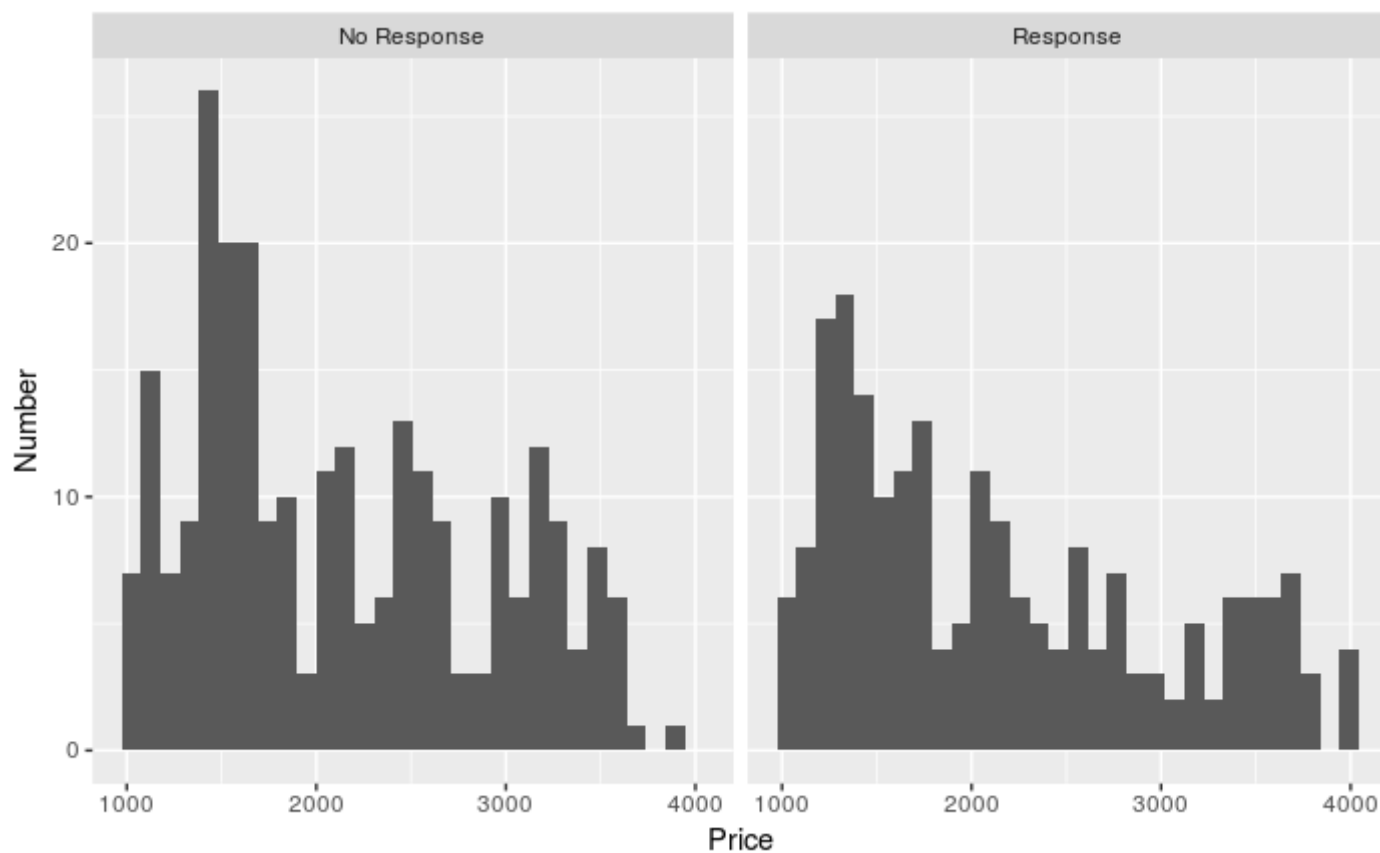
Hide

```
table(d$outcome_f,factor(d$bedrooms))
```

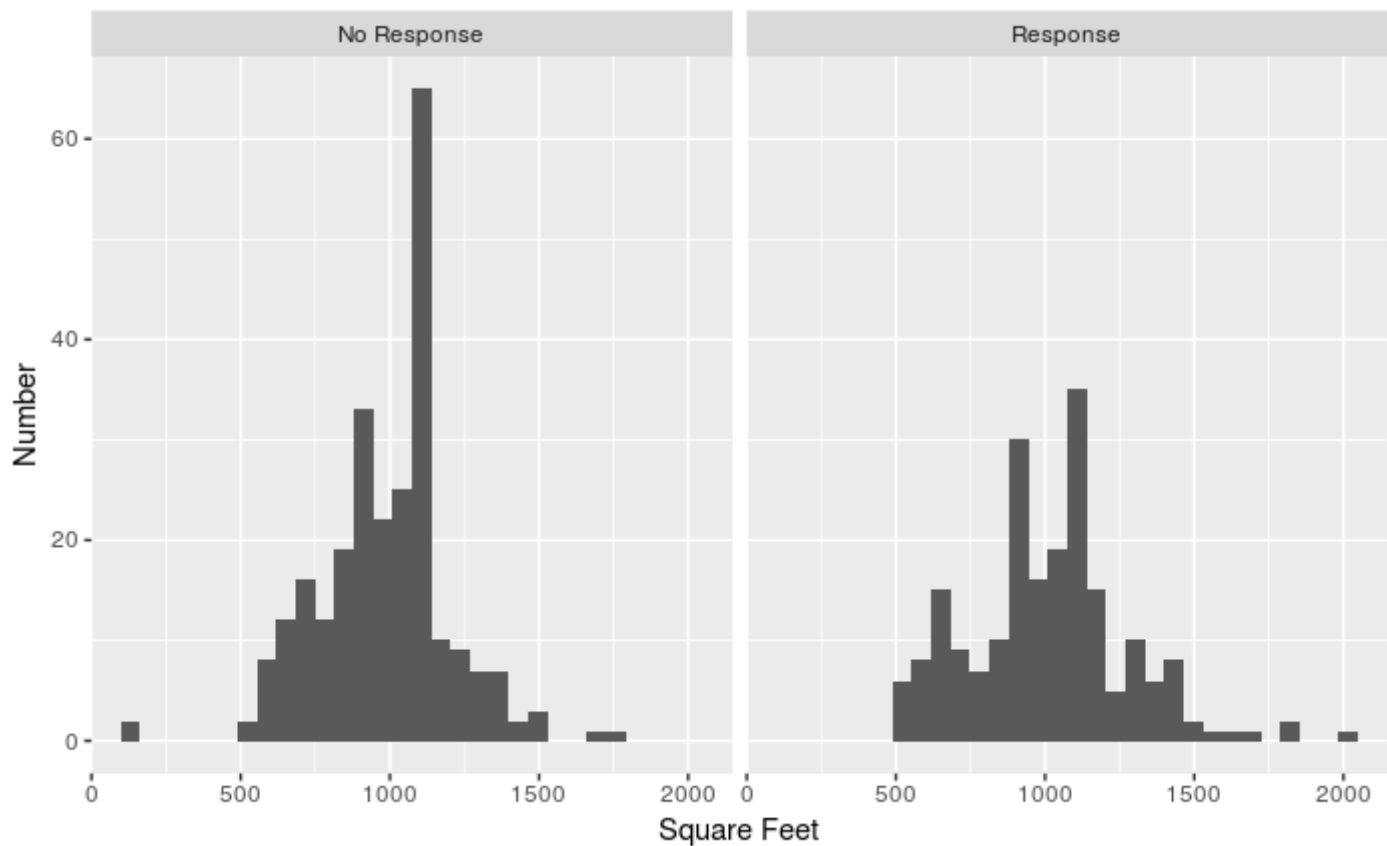
	1-bedroom	2-bedroom
No Response	125	131
Response	96	111

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```
ggplot(d,aes(x=price))+geom_histogram()+facet_grid(~outcome_f)+labs(x="Price",y="Number")
```

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```
# Sqft info has missing values => we can drop all cases (see above) but for now leave this alone or we use the clustering estimate
# Similar but somewhat worse issue for professional, same.email info
ggplot(d,aes(x=sqft))+geom_histogram()+facet_grid(~outcome_f)+labs(x="Square Feet",y="Number")
```



Hide

```
cat('\nTable of Outcomes (By Professional):')
```

Table of Outcomes (By Professional):

Hide

```
table(d$outcome_f,factor(d$professional))
```

	Non-professional	Professional
No Response	55	201
Response	35	172

3. Analysis

Simple Analysis

We do a chi-squared test of independence to see if the observations are independent. We cannot reject the hypothesis that the observations are independent. This is true for even the professional category.

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```
# For Outcome and Gender
tbl <- table(d$outcome_f,d$gender)
tbl
```

	Male	Female
No Response	131	125
Response	100	107

[Hide](#)

```
chisq.test(tbl)
```

Pearson's Chi-squared test with Yates' continuity correction

```
data: tbl
X-squared = 0.26941, df = 1, p-value = 0.6037
```

[Hide](#)

```
# On Outcome and Treatment
tbl <- table(d$outcome_f,d$treatment)
tbl
```

	0	1	2
No Response	87	87	82
Response	68	68	71

[Hide](#)

```
chisq.test(tbl)
```

Pearson's Chi-squared test

```
data: tbl
X-squared = 0.26615, df = 2, p-value = 0.8754
```

[Hide](#)

```
# On Outcome and Treatment Assignment
tbl <- table(d$outcome_f,factor(d$treatment_assignment))
tbl
```


	Jane_Control	Jane_Treat_High	Jane_Treat_Low	John_Control	John_Treat_High
John_Treat_Low					
No Response	39	38	48	48	44
Response	38	37	32	30	34

Hide

```
chisq.test(tbl)
```

Pearson's Chi-squared test

```
data: tbl
X-squared = 3.6372, df = 5, p-value = 0.6027
```

Hide

```
# On Outcome and Professional
tbl <- table(d$outcome_f,d$professional)
tbl
```

	Non-professional	Professional
No Response	55	201
Response	35	172

Hide

```
chisq.test(tbl)
```

Pearson's Chi-squared test with Yates' continuity correction

```
data: tbl
X-squared = 1.2523, df = 1, p-value = 0.2631
```

Regression

We run regression on treatment as a factor (control, low, high) with and without gender as another factor. Other co-variates are added including city, price, bedrooms.

Basic model

```
Outcome variable = alpha + B_high + B_low + gender + covariates
```

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```
# First we treat treatment as a continous variable
# Model 1a - Basic model
m1 <- lm(outcome~treatment,data=d)
stargazer(m1,type='text')
```

```
=====
                        Dependent variable:
                        -----
                        outcome
-----
treatment                0.013
                        (0.028)

Constant                0.434***
                        (0.037)

-----
Observations                463
R2                        0.0004
Adjusted R2                -0.002
Residual Std. Error      0.498 (df = 461)
F Statistic              0.198 (df = 1; 461)
=====
Note:          *p<0.1; **p<0.05; ***p<0.01
```

[Hide](#)

```
coeftest(m1, vcovHC(m1)) # Robust se
```

t test of coefficients:

```
              Estimate Std. Error t value Pr(>|t|)
(Intercept) 0.434495    0.036582 11.8771  <2e-16 ***
treatment   0.012644    0.028502  0.4436  0.6575
---
Signif. codes:  0 '***' 0.001 '**' 0.01 '*' 0.05 '.' 0.1 ' ' 1
```

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```
# Model 2a - Treatment & gender
m2 <- lm(outcome~treatment*gender,data=d)
stargazer(m2,type='text')
```

```

=====
                        Dependent variable:
                        -----
                                outcome
                        -----
treatment                0.026
                        (0.040)

genderFemale              0.054
                        (0.073)

treatment:genderFemale   -0.026
                        (0.057)

Constant                 0.407***
                        (0.052)

-----
Observations              463
R2                        0.002
Adjusted R2              -0.005
Residual Std. Error      0.499 (df = 459)
F Statistic              0.262 (df = 3; 459)
=====
Note:                    *p<0.1; **p<0.05; ***p<0.01

```

[Hide](#)

```
coeftest(m2, vcovHC(m2)) # Robust se
```

```
t test of coefficients:
```

```

                Estimate Std. Error t value Pr(>|t|)
(Intercept)    0.407259   0.051210  7.9527 1.434e-14 ***
treatment      0.025641   0.039798  0.6443  0.5197
genderFemale    0.054454   0.073288  0.7430  0.4579
treatment:genderFemale -0.026152  0.057200 -0.4572  0.6477
---
Signif. codes:  0 '***' 0.001 '**' 0.01 '*' 0.05 '.' 0.1 ' ' 1

```

[Hide](#)

```

# Model 3a - Treatment & gender + covariates
m3 <- lm(outcome~treatment*gender+factor(city)+factor(bedrooms)+price,data=d)
stargazer(m3,type='text')

```

```

=====
                        Dependent variable:
                        -----
                                outcome
                        -----
treatment                0.027
                        (0.040)

genderFemale              0.052
                        (0.073)

factor(city)houston      -0.164**
                        (0.064)

factor(city)sandiego     -0.051
                        (0.064)

factor(city)seattle      0.042
                        (0.066)

factor(bedrooms)2-bedroom 0.027
                        (0.050)

price                    -0.00000
                        (0.00003)

treatment:genderFemale   -0.025
                        (0.057)

Constant                 0.441***
                        (0.088)

-----
Observations              463
R2                        0.026
Adjusted R2              0.009
Residual Std. Error      0.496 (df = 454)
F Statistic              1.516 (df = 8; 454)
=====
Note:                    *p<0.1; **p<0.05; ***p<0.01

```

[Hide](#)

```
coeftest(m3, vcovHC(m3)) # Robust se
```

t test of coefficients:

	Estimate	Std. Error	t value	Pr(> t)	
(Intercept)	4.4134e-01	9.0292e-02	4.8880	1.416e-06	***
treatment	2.7172e-02	3.9686e-02	0.6847	0.49389	
genderFemale	5.2129e-02	7.3091e-02	0.7132	0.47609	
factor(city)houston	-1.6386e-01	6.3923e-02	-2.5633	0.01069	*
factor(city)sandiego	-5.1371e-02	6.5858e-02	-0.7800	0.43578	
factor(city)seattle	4.2490e-02	6.7313e-02	0.6312	0.52821	
factor(bedrooms)2-bedroom	2.7215e-02	4.9288e-02	0.5522	0.58111	
price	-2.4420e-06	3.1488e-05	-0.0776	0.93822	
treatment:genderFemale	-2.4833e-02	5.7221e-02	-0.4340	0.66451	

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

Hide

```
# Model 3a-1 - Treatment & gender + all covariates including less reliable sqft and professional
m3.1 <- lm(outcome~treatment*gender+factor(city)+factor(bedrooms)+price+sqft+professional,data=d)
stargazer(m3.1,type='text')
```

```

=====
                        Dependent variable:
                        -----
                        outcome
                        -----
treatment                0.026
                        (0.040)

genderFemale             0.059
                        (0.073)

factor(city)houston     -0.181***
                        (0.067)

factor(city)sandiego    -0.031
                        (0.068)

factor(city)seattle      0.051
                        (0.071)

factor(bedrooms)2-bedroom -0.009
                        (0.055)

price                   -0.00003
                        (0.00003)

sqft                    0.0002*
                        (0.0001)

professionalProfessional 0.124**
                        (0.062)

treatment:genderFemale  -0.027
                        (0.057)

Constant                0.193
                        (0.144)

-----
Observations              463
R2                        0.039
Adjusted R2               0.018
Residual Std. Error      0.493 (df = 452)
F Statistic               1.832* (df = 10; 452)
=====
Note:                    *p<0.1; **p<0.05; ***p<0.01

```

[Hide](#)

```
coeftest(m3.1, vcovHC(m3.1)) # Robust se
```

t test of coefficients:

	Estimate	Std. Error	t value	Pr(> t)
(Intercept)	1.9334e-01	1.4554e-01	1.3285	0.18470
treatment	2.6110e-02	3.9819e-02	0.6557	0.51235
genderFemale	5.9426e-02	7.2995e-02	0.8141	0.41601
factor(city)houston	-1.8053e-01	6.6516e-02	-2.7141	0.00690 **
factor(city)sandiego	-3.0741e-02	6.9931e-02	-0.4396	0.66044
factor(city)seattle	5.0970e-02	7.2326e-02	0.7047	0.48134
factor(bedrooms)2-bedroom	-8.6209e-03	5.3405e-02	-0.1614	0.87183
price	-3.0499e-05	3.4144e-05	-0.8932	0.37220
sqft	2.2319e-04	1.2288e-04	1.8164	0.06998 .
professionalProfessional	1.2359e-01	6.3672e-02	1.9410	0.05288 .
treatment:genderFemale	-2.6580e-02	5.7256e-02	-0.4642	0.64271

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

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```
# Next we treat treatment as a categorical variable (effect might not be linear)
# Model lb - Basic model
m4 <- lm(outcome~treatment_f,data=d)
stargazer(m4,type='text')
```

```
=====
Dependent variable:
-----
outcome
-----
treatment_fLow      0.000
                    (0.057)

treatment_fHigh     0.025
                    (0.057)

Constant            0.439***
                    (0.040)

-----
Observations        463
R2                  0.001
Adjusted R2         -0.004
Residual Std. Error 0.499 (df = 460)
F Statistic         0.132 (df = 2; 460)
=====
Note:                *p<0.1; **p<0.05; ***p<0.01
```

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```
coeftest(m4, vcovHC(m4)) # Robust se
```

t test of coefficients:

	Estimate	Std. Error	t value	Pr(> t)
(Intercept)	4.3871e-01	4.0117e-02	10.9358	<2e-16 ***
treatment_fLow	2.4600e-16	5.6734e-02	0.0000	1.0000
treatment_fHigh	2.5343e-02	5.7065e-02	0.4441	0.6572

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

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```
# Model 2b - Treatment & gender
m5 <- lm(outcome~treatment_f*gender,data=d)
stargazer(m5,type='text')
```

```
=====
                        Dependent variable:
                        -----
                        outcome
-----
treatment_fLow          0.095
                        (0.081)

treatment_fHigh         0.051
                        (0.080)

genderFemale            0.109
                        (0.080)

treatment_fLow:genderFemale -0.189*
                        (0.113)

treatment_fHigh:genderFemale -0.051
                        (0.114)

Constant                0.385***
                        (0.056)

-----
Observations            463
R2                      0.008
Adjusted R2             -0.003
Residual Std. Error     0.498 (df = 457)
F Statistic              0.724 (df = 5; 457)
=====
Note:                    *p<0.1; **p<0.05; ***p<0.01
```

[Hide](#)

```
coeftest(m5, vcovHC(m5)) # Robust se
```


t test of coefficients:

	Estimate	Std. Error	t value	Pr(> t)	
(Intercept)	0.384615	0.055801	6.8926	1.827e-11	***
treatment_fLow	0.095385	0.080823	1.1802	0.23855	
treatment_fHigh	0.051282	0.079678	0.6436	0.52015	
genderFemale	0.108891	0.080287	1.3563	0.17568	
treatment_fLow:genderFemale	-0.188891	0.113758	-1.6605	0.09751	.
treatment_fHigh:genderFemale	-0.051455	0.114474	-0.4495	0.65329	

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

Hide

```
# Model 3b - Treatment & gender + covariates
m6 <- lm(outcome~treatment_f*gender+factor(city)+factor(bedrooms)+price,data=d)
stargazer(m6,type='text')
```

```

=====
                        Dependent variable:
                        -----
                        outcome
                        -----
treatment_fLow           0.099
                        (0.080)

treatment_fHigh          0.054
                        (0.079)

genderFemale             0.109
                        (0.080)

factor(city)houston      -0.169***
                        (0.064)

factor(city)sandiego     -0.053
                        (0.064)

factor(city)seattle      0.038
                        (0.066)

factor(bedrooms)2-bedroom 0.028
                        (0.050)

price                   -0.00000
                        (0.00003)

treatment_fLow:genderFemale -0.194*
                        (0.113)

treatment_fHigh:genderFemale -0.049
                        (0.113)

Constant                 0.419***
                        (0.092)

-----
Observations              463
R2                        0.033
Adjusted R2               0.011
Residual Std. Error       0.495 (df = 452)
F Statistic               1.527 (df = 10; 452)
=====
Note:                      *p<0.1; **p<0.05; ***p<0.01

```

[Hide](#)

```
coeftest(m6, vcovHC(m6)) # Robust se
```

t test of coefficients:

	Estimate	Std. Error	t value	Pr(> t)	
(Intercept)	4.1906e-01	9.4500e-02	4.4345	1.16e-05	***
treatment_fLow	9.8680e-02	8.0530e-02	1.2254	0.221067	
treatment_fHigh	5.4357e-02	7.9457e-02	0.6841	0.494254	
genderFemale	1.0870e-01	8.0054e-02	1.3578	0.175197	
factor(city)houston	-1.6851e-01	6.3987e-02	-2.6335	0.008742	**
factor(city)sandiego	-5.3147e-02	6.6088e-02	-0.8042	0.421713	
factor(city)seattle	3.8055e-02	6.7536e-02	0.5635	0.573391	
factor(bedrooms)2-bedroom	2.8279e-02	4.9665e-02	0.5694	0.569366	
price	-1.8808e-06	3.1555e-05	-0.0596	0.952499	
treatment_fLow:genderFemale	-1.9405e-01	1.1360e-01	-1.7081	0.088303	.
treatment_fHigh:genderFemale	-4.8976e-02	1.1449e-01	-0.4278	0.669017	

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

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```
# Model 3b-1 - Treatment & gender + all covariates including less reliable sqft and professional
m6.1 <- lm(outcome~treatment*gender+factor(city)+factor(bedrooms)+price+sqft+professional,data=d)
stargazer(m6.1,type='text')
```

```

=====
                        Dependent variable:
                        -----
                        outcome
                        -----
treatment                0.026
                        (0.040)

genderFemale              0.059
                        (0.073)

factor(city)houston      -0.181***
                        (0.067)

factor(city)sandiego     -0.031
                        (0.068)

factor(city)seattle      0.051
                        (0.071)

factor(bedrooms)2-bedroom -0.009
                        (0.055)

price                    -0.00003
                        (0.00003)

sqft                     0.0002*
                        (0.0001)

professionalProfessional  0.124**
                        (0.062)

treatment:genderFemale   -0.027
                        (0.057)

Constant                 0.193
                        (0.144)

-----
Observations              463
R2                        0.039
Adjusted R2               0.018
Residual Std. Error       0.493 (df = 452)
F Statistic               1.832* (df = 10; 452)
=====
Note:                    *p<0.1; **p<0.05; ***p<0.01

```

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```
coeftest(m6.1, vcovHC(m6.1)) # Robust se
```

t test of coefficients:

	Estimate	Std. Error	t value	Pr(> t)							
(Intercept)	1.9334e-01	1.4554e-01	1.3285	0.18470							
treatment	2.6110e-02	3.9819e-02	0.6557	0.51235							
genderFemale	5.9426e-02	7.2995e-02	0.8141	0.41601							
factor(city)houston	-1.8053e-01	6.6516e-02	-2.7141	0.00690	**						
factor(city)sandiego	-3.0741e-02	6.9931e-02	-0.4396	0.66044							
factor(city)seattle	5.0970e-02	7.2326e-02	0.7047	0.48134							
factor(bedrooms)2-bedroom	-8.6209e-03	5.3405e-02	-0.1614	0.87183							
price	-3.0499e-05	3.4144e-05	-0.8932	0.37220							
sqft	2.2319e-04	1.2288e-04	1.8164	0.06998	.						
professionalProfessional	1.2359e-01	6.3672e-02	1.9410	0.05288	.						
treatment:genderFemale	-2.6580e-02	5.7256e-02	-0.4642	0.64271							

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

In all models, the coefficients on treatment, whether continuous or as a factor, are not statistically significant. If we add gender, there is also no evidence of a the interaction term being statistically significant. Thus, there is no evidence that exclamation points have influenced the likelihood of receiving a response.

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```
# We try an alternative specification for treatment (as dummy variables)

# Model 1c - Basic model
m7 <- lm(outcome ~ low_treatment + high_treatment, data=d)

stargazer(m7, type='text')
coeftest(m7, vcovHC(m7)) # Robust se

# Model 2c - Treatment & gender
m8 <- lm(outcome ~ low_treatment + high_treatment*gender, data=d)
stargazer(m8, type='text')
coeftest(m8, vcovHC(m8)) # Robust se

# Model 3c - Treatment & gender + covariates
m9 <- lm(outcome ~ low_treatment + high_treatment + gender + factor(city) + factor(bedrooms) + price, data=d)
stargazer(m9, type='text')
coeftest(m9, vcovHC(m9)) # Robust se
```

The coefficients on treatment are also statistically insignificant. There is no evidence that exclamation points have an effect.

Randomization Inference

Next we use randomization inference (assuming a Sharp Null of No Effect) to understand if our observation is consistent with an empirical null distribution. For this, we combine low and high treatment into treatment (since we have not learned more complex fixes for heterogenous effects).

Hide

```
# Combining treatments
di <- d
di[treatment==2,treatment:=1]

# Define distributions
y <- di$outcome
Z <- di$treatment
blk1 <- as.numeric(di$gender) # We block by gender
blk2 <- as.numeric(di$city) # Block by city
blk3 <- as.numeric(di$bedrooms)

# By gender
perms <- genperms(Z, clustvar = NULL, blockvar = blk1)
probs <- genprobexact(Z, clustvar = NULL, blockvar = blk1) # probability of treatment
ate <- estate(y,Z,prob=probs) # estimate the ATE

Ys <- genouts(y,Z,ate=0) # generate potential outcomes under sharp null of no effect
distout <- gendist(Ys,perms, prob=probs) # generate sampling dist. under sharp null
dispdist(distout, ate, quantiles = c(0.025, 0.975), display.plot = TRUE) # display characteristics of sampling dist. for inference

# By city
perms <- genperms(Z, clustvar = NULL, blockvar = blk2)
probs <- genprobexact(Z, clustvar = NULL, blockvar = blk2) # probability of treatment
ate <- estate(y,Z,prob=probs) # estimate the ATE

Ys <- genouts(y,Z,ate=0) # generate potential outcomes under sharp null of no effect
distout <- gendist(Ys,perms, prob=probs) # generate sampling dist. under sharp null
dispdist(distout, ate, quantiles = c(0.025, 0.975), display.plot = TRUE) # display characteristics of sampling dist. for inference

# By bedroom
perms <- genperms(Z, clustvar = NULL, blockvar = blk3)
probs <- genprobexact(Z, clustvar = NULL, blockvar = blk3) # probability of treatment
ate <- estate(y,Z,prob=probs) # estimate the ATE

Ys <- genouts(y,Z,ate=0) # generate potential outcomes under sharp null of no effect
distout <- gendist(Ys,perms, prob=probs) # generate sampling dist. under sharp null
dispdist(distout, ate, quantiles = c(0.025, 0.975), display.plot = TRUE) # display characteristics of sampling dist. for inference

#P-value for actual data
p.val.actual = sum(abs(distout) > ate) / length(distout)
p.val.actual

#get response rate by treatment or control
actual.response.rate.by.treatment <- di[, mean(outcome), by = c("treatment")]
actual.response.rate.by.treatment

di[, sum(outcome > -100), by = c("treatment")]
```

Once again, we cannot reject the null hypothesis of no effect.

Other Analysis

1. Although not a significant issue for this experiment we estimate the CACE. For this we define non-compliers as those for who we sent emails but did not received them - and we know this because we received a “bounced” email message.

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```
# We calculate the CACE manually

# Manually compute CACE
itt <- mean(d$outcome[d$treatment != 0]) - mean(d$outcome[d$treatment == 0])
prop_treated <- 481/483

sprintf("\nThe estimated CACE is: %.5f", itt / prop_treated)

# or 2SLS
#itt_fit <- ivreg(outcome ~treatment,~compliers,data=d)
#stargazer(itt_fit, type='text')
```

2. We also did some work Work to find treatment response rate required to reject null. That is, how much does the

[Hide](#)

```
sprintf("Which is about %0.f times more than what we see currently", 2/z[6])
```

```
[1] "Which is about 5 times more than what we see currently"
```

4. Conclusion

Despite running a few different models, we find no evidence that the number of exclamation points affected response rates to our email.