

ImpactDataViz 241 Final Project

```
# load packages
library(data.table)
library(foreign)
library(sandwich)
library(stargazer)

##
## Please cite as:

## Hlavac, Marek (2018). stargazer: Well-Formatted Regression and Summary Statistics Tables.

## R package version 5.2.2. https://CRAN.R-project.org/package=stargazer

library(lmtest)

## Loading required package: zoo

## Warning: package 'zoo' was built under R version 3.6.2

##
## Attaching package: 'zoo'

## The following objects are masked from 'package:base':
##
##      as.Date, as.Date.numeric

library(tidyr)

## Warning: package 'tidyr' was built under R version 3.6.2

library(knitr)
library('ggplot2')
library(fastDummies)
```

Common functions

```

# function to return confidence intervals with robust se
get_confint_robust = function(model, vcovCL) {
  t<-qt(.975, model$df.residual)
  ct<-coeftest(model, vcovCL)
  est<-cbind(ct[,1], ct[,1]-t*ct[,2], ct[,1]+t*ct[,2], ct[,4])
  colnames(est)<-c("Estimate", "LowerCI", "UpperCI", "pValue")
  return(est)
}

# parse out the regression results using robust standard errors
get_regression_results_robust_se = function(model, df, variable_names, showAsTibble) {
  model$vcovHC = vcovHC(model, type="HC1")

  robust_se_all <- sqrt(diag(model$vcovHC))

  est = get_confint_robust(model, model$vcovHC)

  robust_se = c(rep(0, length(variable_names)))
  i = 1
  for (variable_name in variable_names) {
    robust_se_single <- sqrt(diag(model$vcovHC))[variable_name]
    robust_se[i] = robust_se_single
    i = i + 1
  }

  coef = est[variable_names, 'Estimate']
  ci_lower_robust = est[variable_names, 'LowerCI']
  ci_upper_robust = est[variable_names, 'UpperCI']
  p_value = est[variable_names, 'pValue']
  results = data.table(id = variable_names)
  results[, coef := round(coef, 4)]
  results[, ci_lower := round(ci_lower_robust, 4)]
  results[, ci_upper := round(ci_upper_robust, 4)]
  results[, p_value := signif(p_value, 5)]
  results[, robust_se := round(robust_se, 4)]

  if (showAsTibble) {
    print(as_tibble(results))
  }
  return( list('estimates'=results, 'robust_se_all'=robust_se_all))
}

# Multiple plot function
#
# ggplot objects can be passed in ..., or to plotlist (as a list of ggplot objects)
# - cols: Number of columns in layout
# - layout: A matrix specifying the layout. If present, 'cols' is ignored.
#
# If the layout is something like matrix(c(1,2,3,3), nrow=2, byrow=TRUE),
# then plot 1 will go in the upper left, 2 will go in the upper right, and
# 3 will go all the way across the bottom.
#
multiplot <- function(..., plotlist=NULL, file, cols=1, layout=NULL) {

```

```

library(grid)

# Make a list from the ... arguments and plotlist
plots <- c(list(...), plotlist)

numPlots = length(plots)

# If layout is NULL, then use 'cols' to determine layout
if (is.null(layout)) {
  # Make the panel
  # ncol: Number of columns of plots
  # nrow: Number of rows needed, calculated from # of cols
  layout <- matrix(seq(1, cols * ceiling(numPlots/cols)),
                    ncol = cols, nrow = ceiling(numPlots/cols))
}

if (numPlots==1) {
  print(plots[[1]])
} else {
  # Set up the page
  grid.newpage()
  pushViewport(viewport(layout = grid.layout(nrow(layout), ncol(layout))))

  # Make each plot, in the correct location
  for (i in 1:numPlots) {
    # Get the i,j matrix positions of the regions that contain this subplot
    matchidx <- as.data.frame(which(layout == i, arr.ind = TRUE))

    print(plots[[i]], vp = viewport(layout.pos.row = matchidx$row,
                                     layout.pos.col = matchidx$col))
  }
}

```

1. Parse the survey data into a data.table

```

parse_survey_data = function(filename, treatment_only=FALSE) {
  cat(filename, "\n")
  raw <- fread(filename)

  # covariates
  setnames(raw, 'Q3', 'state')
  setnames(raw, 'Q4', 'gender')
  setnames(raw, 'Q5', 'age')
  setnames(raw, 'Q6', 'ethnicity_multi')
  setnames(raw, 'Q7', 'political_party')
  setnames(raw, 'Q8', 'education')
  setnames(raw, 'Q9', 'covid_sick')
  setnames(raw, 'Q10', 'covid_hospitalized')
  setnames(raw, 'Q11', 'covid_died')

```

```

# duration of survey time
setnames(raw, 'Duration (in seconds)', 'duration_of_survey')

# which block was active? (did the user see the treatment or
# control data viz)
setnames(raw, 'Q15', 'treatment_viz_is_accurate')
if (!treatment_only) {
  setnames(raw, 'Q17', 'control_viz_is_accurate')
}

# outcome questions about COVID attitudes
setnames(raw, 'Q18', 'outcome_spread')
setnames(raw, 'Q19', 'outcome_death')

# which block was active determines if
# subject received treatment data viz or control
# data viz
cat(" number of responses", nrow(raw), '\n')
raw[, treatment := ifelse(is.na(treatment_viz_is_accurate), 0, 1)]
cleaned = raw[!is.na(outcome_spread) & !is.na(outcome_death),]
cat(" number of responses after dropping na", nrow(cleaned), '\n')

# ethnicity allows for multiple choice
# for covariates, just grab the first one
ethnicity_single = rep(0, nrow(cleaned))
i = 1
for (eth_entry in cleaned[, ethnicity_multi]) {
  eth_tokens = unlist(strsplit(eth_entry, ","))
  ethnicity_single[i] = as.numeric(eth_tokens[1])
  i = i + 1
}
cleaned[, ethnicity := ethnicity_single ]

# counts in control vs treatment
n_control = nrow(cleaned[treatment == 0, ])
n_treatment = nrow(cleaned[treatment == 1, ])

cat(" number in treatment", n_treatment, "\n")
cat(" number in control", n_control, "\n\n")

return(cleaned)
}

```

```

# A large run of 260 subjects run on 7/21
run1 <- parse_survey_data("data/run1.2020.07.21.csv")

```

```

## data/run1.2020.07.21.csv
## number of responses 265
## number of responses after dropping na 259
## number in treatment 131
## number in control 128

```

```
run1 = run1[, run := 0]
run1[, condition := treatment]

# One small run was done in evening 7/24 treatment only
run2_small <- parse_survey_data("data/run2.small.2020.07.24.csv", TRUE)
```

```
## data/run2.small.2020.07.24.csv
## number of responses 33
## number of responses after dropping na 30
## number in treatment 30
## number in control 0
```

```
run2_small[, run := 1]
run2_small[, condition := treatment]
run2_small[, control_viz_is_accurate := ""]

# A large run of 270 run on 7/25 treatment and control
run2_large <- parse_survey_data("data/run2.2020.07.25.csv")
```

```
## data/run2.2020.07.25.csv
## number of responses 270
## number of responses after dropping na 262
## number in treatment 130
## number in control 132
```

```
run2_large = run2_large[, run := 1]
run2_large[, condition := treatment]
run2 = rbind(run2_small, run2_large)
```

```
# Combine the runs on 7/21 and 7/25
combined = rbind(run1, run2)
```

2. EDA

2.1 Check Duration of Survey

```
show_duration = function(d) {
  cat('duration for control ', mean(d[condition == 0, duration_of_survey]), '\n')
  cat('duration for treatment', mean(d[condition == 1, duration_of_survey]), '\n\n')
}
show_duration(run1)
```

```
## duration for control 153.1875
## duration for treatment 172.2061
```

```
show_duration(run2)
```

```
## duration for control    136.7955  
## duration for treatment 205.5813
```

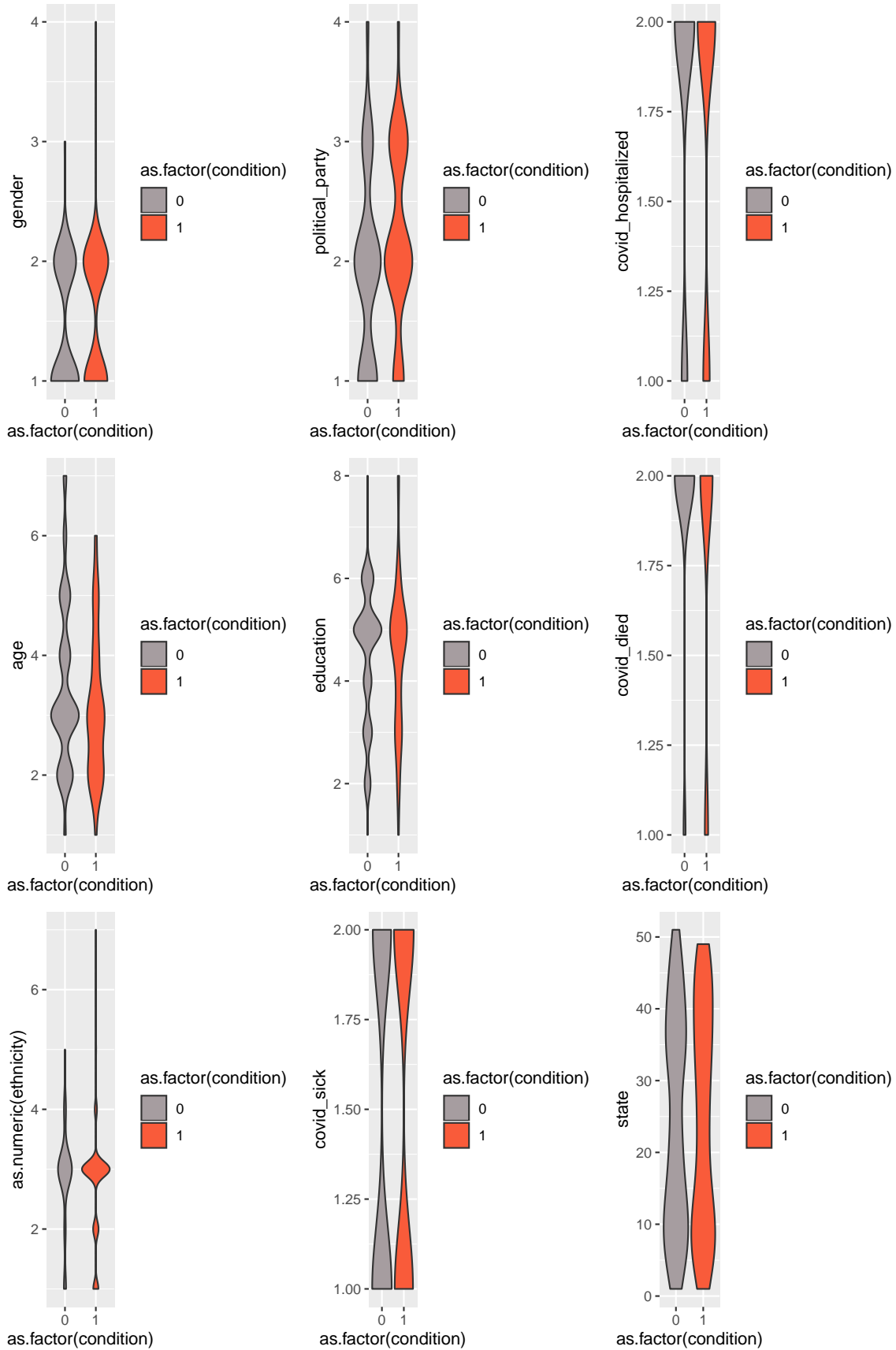
```
show_duration(combined)
```

```
## duration for control    144.8654  
## duration for treatment 190.5567
```

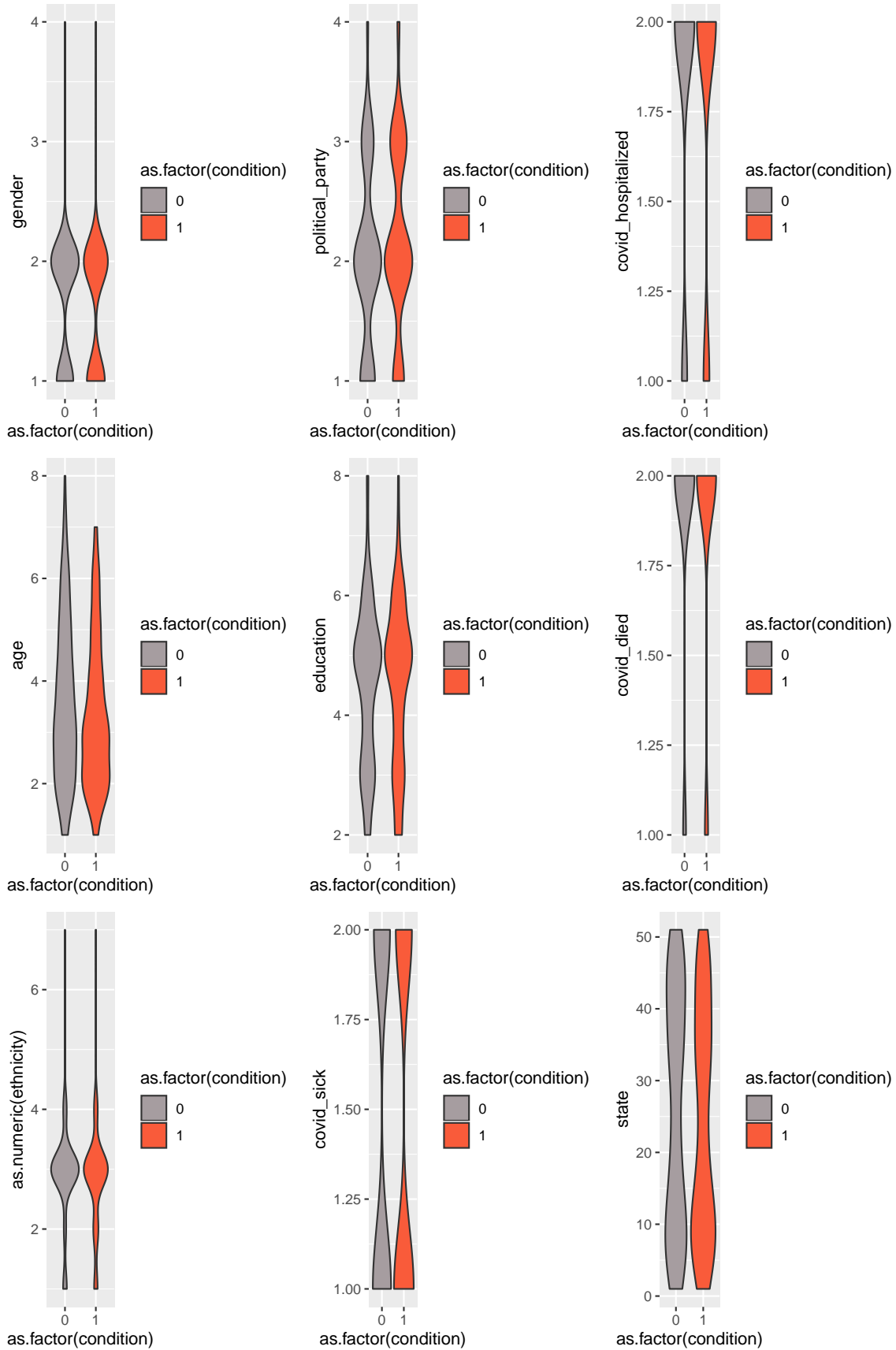
3. Covariate balance

3.1 Compare distributions using violin plote

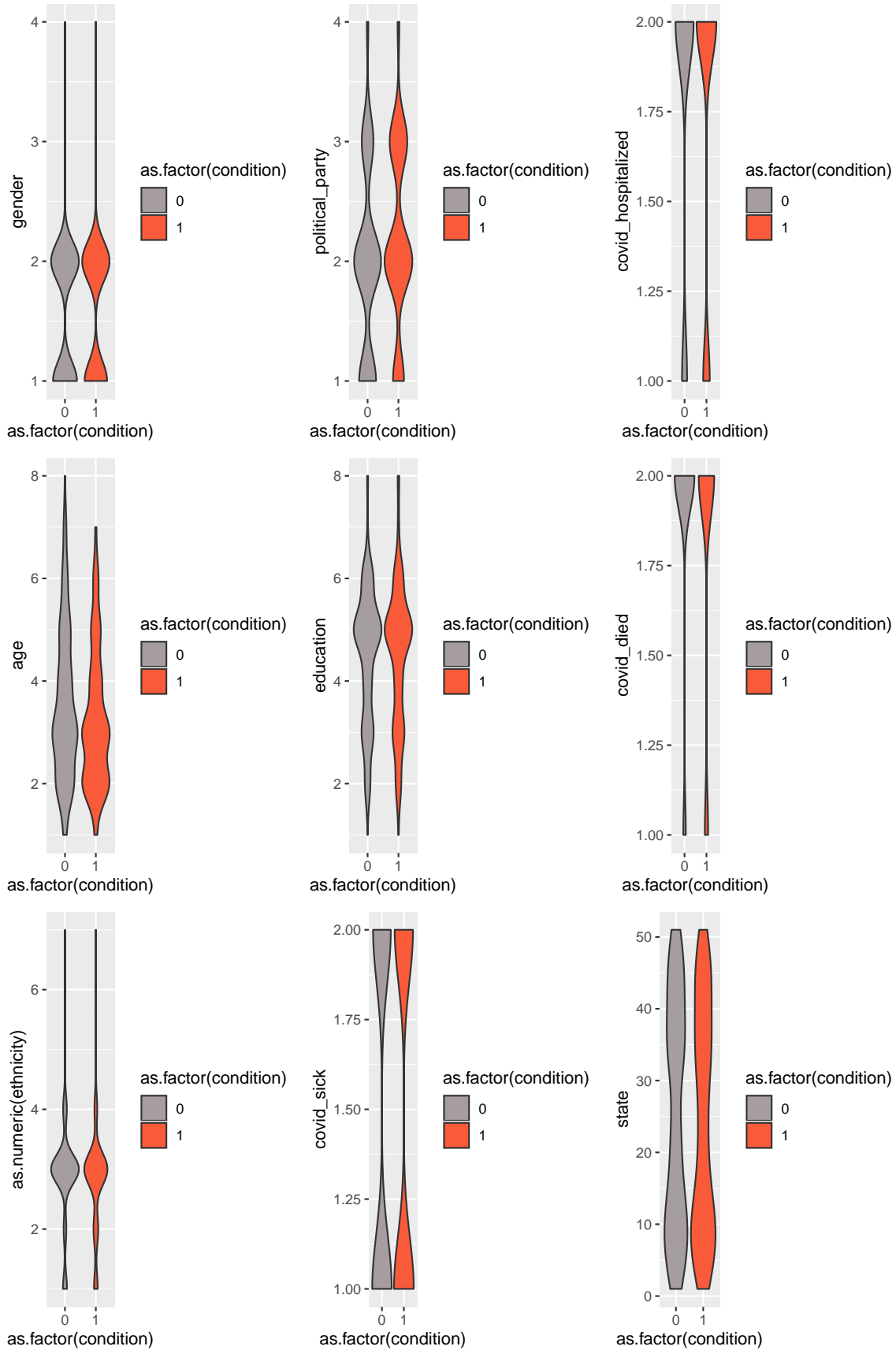
```
check_covariate_balance = function(d) {  
  options(repr.plot.width = 14, repr.plot.height = 8)  
  
  p1 = ggplot(d, aes(x=as.factor(condition), y=gender, fill=as.factor(condition))) +  
    geom_violin() + scale_fill_manual(values=c("#a69da0", "#f95b3a"))  
  
  p2 = ggplot(d, aes(x=as.factor(condition), y=age, fill=as.factor(condition))) +  
    geom_violin() + scale_fill_manual(values=c("#a69da0", "#f95b3a"))  
  
  p3 = ggplot(d, aes(x=as.factor(condition), y=as.numeric(ethnicity), fill=as.factor(condition))) +  
    geom_violin() + scale_fill_manual(values=c("#a69da0", "#f95b3a"))  
  
  p4 = ggplot(d, aes(x=as.factor(condition), y=political_party, fill=as.factor(condition))) +  
    geom_violin() + scale_fill_manual(values=c("#a69da0", "#f95b3a"))  
  
  p5 = ggplot(d, aes(x=as.factor(condition), y=education, fill=as.factor(condition))) +  
    geom_violin() + scale_fill_manual(values=c("#a69da0", "#f95b3a"))  
  
  p6 = ggplot(d, aes(x=as.factor(condition), y=covid_sick, fill=as.factor(condition))) +  
    geom_violin() + scale_fill_manual(values=c("#a69da0", "#f95b3a"))  
  
  p7 = ggplot(d, aes(x=as.factor(condition), y=covid_hospitalized, fill=as.factor(condition))) +  
    geom_violin() + scale_fill_manual(values=c("#a69da0", "#f95b3a"))  
  
  p8 = ggplot(d, aes(x=as.factor(condition), y=covid_died, fill=as.factor(condition))) +  
    geom_violin() + scale_fill_manual(values=c("#a69da0", "#f95b3a"))  
  
  p9 = ggplot(d, aes(x=as.factor(condition), y=state, fill=as.factor(condition))) +  
    geom_violin() + scale_fill_manual(values=c("#a69da0", "#f95b3a"))  
  
  multiplot(p1, p2, p3, p4, p5, p6, p7, p8, p9, cols=3)  
}  
check_covariate_balance(run1)
```



```
check_covariate_balance(run2)
```

```
check_covariate_balance(combined)
```



4. Estimate ATE

4.1 Estimate the ATE for both outcomes

```
estimate_ate = function(d, outcome_field) {
  g <- d[, .(group_mean = mean(get(outcome_field))), keyby = .(condition)]

  ate <- g[, diff(group_mean)]

  res <- NA
  for (i in 1:10000) {
    res[i] <- d[, .(group_mean = mean(get(outcome_field))), keyby = .(sample(condition))][, diff(gr
  ]
  dist_sharp_null <- res
  #hist(dist_sharp_null)
  #abline(v=ate, lwd=3, col='blue')
  #abline(v=abs(ate), lwd=3, col='blue')
  p_value_one_tailed <- mean(dist_sharp_null >= ate)
  p_value_two_tailed <- mean(abs(dist_sharp_null) >= abs(ate))

  cat(outcome_field, '\n')
  cat(' mean control      ', g[condition == '0', group_mean], '\n')
  cat(' mean traitement  ', g[condition == '1', group_mean], '\n')
  cat(' ATE                ', ate, '\n')
  cat(' p_value 1-tailed', p_value_one_tailed, '\n')
  cat(' p_value 2-tailed', p_value_two_tailed, '\n\n')
}

cat('run1', '\n')

## run1

cat('*****', '\n')

## *****

estimate_ate(run1, 'outcome_spread')

## outcome_spread
## mean control      3.132812
## mean traitement  3.473282
## ATE                0.3404699
## p_value 1-tailed 0.0013
## p_value 2-tailed 0.0022
```

```
estimate_ate(run1, 'outcome_death')
```

```
## outcome_death
## mean control      2.398438
## mean traitement   2.656489
## ATE                0.258051
## p_value 1-tailed  0.0013
## p_value 2-tailed  0.0021
```

```
cat('run2', '\n')
```

```
## run2
```

```
cat('*****', '\n')
```

```
## *****
```

```
estimate_ate(run2, 'outcome_spread')
```

```
## outcome_spread
## mean control      3.151515
## mean traitement   3.30625
## ATE                0.1547348
## p_value 1-tailed  0.0876
## p_value 2-tailed  0.1659
```

```
estimate_ate(run2, 'outcome_death')
```

```
## outcome_death
## mean control      2.5
## mean traitement   2.60625
## ATE                0.10625
## p_value 1-tailed  0.1059
## p_value 2-tailed  0.1922
```

```
cat('combined', '\n')
```

```
## combined
```

```
cat('*****', '\n')
```

```
## *****
```

```
estimate_ate(combined, 'outcome_spread')
```

```
## outcome_spread
## mean control      3.142308
## mean traitement   3.381443
## ATE                0.2391356
## p_value 1-tailed  0.0022
## p_value 2-tailed  0.0034
```

```
estimate_ate(combined, 'outcome_death')
```

```
## outcome_death
## mean control      2.45
## mean treatment    2.628866
## ATE               0.178866
## p_value 1-tailed  0.0016
## p_value 2-tailed  0.0024
```

```
meanFunction <- function(x){
return(data.frame(y=round(mean(x),2),label=round(mean(x,na.rm=T),2)))
}
```

```
show_box_plot_spread = function(df, title) {
  df$treatment_group <- as.factor(df$condition)
  p = ggplot(df, aes(x=treatment_group, y=outcome_spread, fill=treatment_group)) + geom_boxplot() +
    scale_fill_manual(values=c("#a69da0", "#f95b3a")) + labs(title=title, x="Control vs. Treatment", y =
      width = 1, linetype = "dashed") + stat_summary(fun.data = meanFunction, geom="text", s

  p + geom_jitter(shape=16, alpha=.3, position=position_jitter(0.2))
}
```

```
show_box_plot_death= function(df, title) {
  df$treatment_group <- as.factor(df$condition)
  p = ggplot(df, aes(x=treatment_group, y=outcome_death, fill=treatment_group)) + geom_boxplot() +
    scale_fill_manual(values=c("#a69da0", "#f95b3a")) + labs(title=title, x="Control vs. Treatment", y =
      width = 1, linetype = "dashed") + stat_summary(fun.data = meanFunction, geom="text", s

  #stat_summary(fun.y=mean, geom="line", shape=10, size=14, color="black", fill="red")

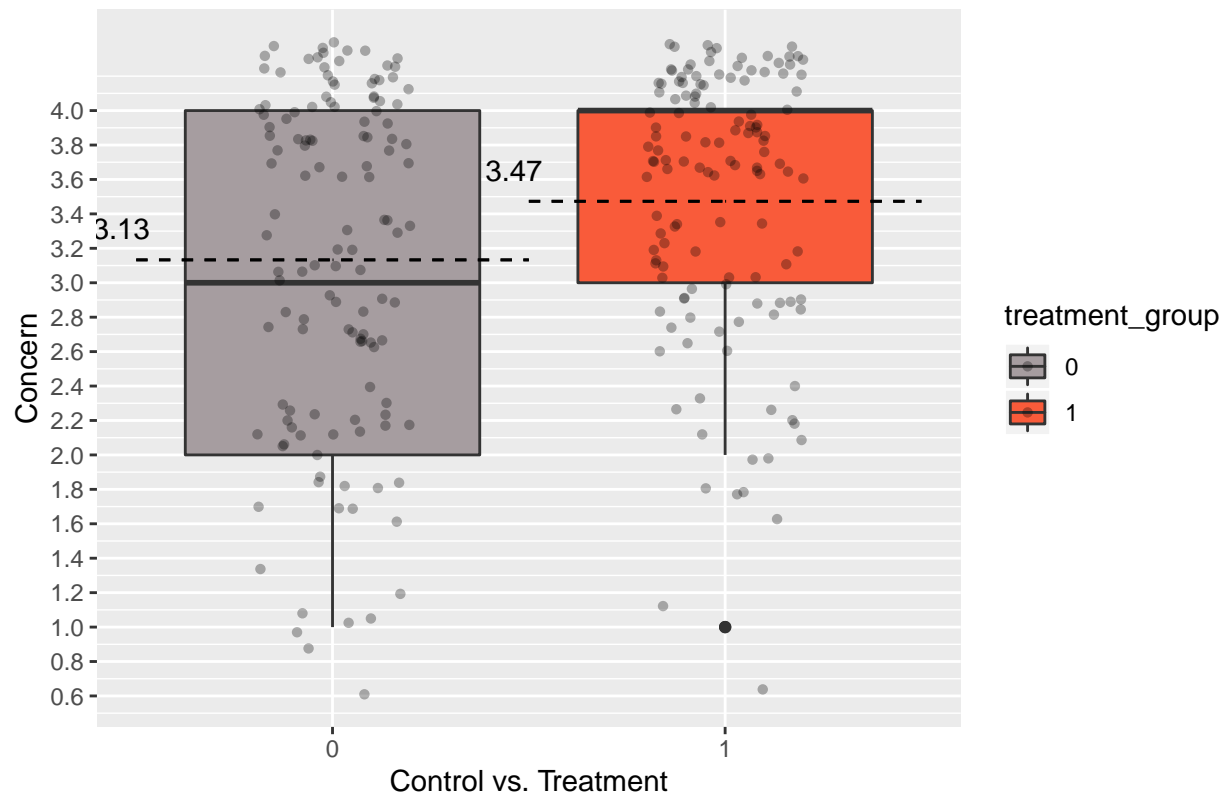
  p + geom_jitter(shape=16, size=2, alpha=.3, position=position_jitter(0.2) )
}
```

```
meanFunction1 <- function(x){
return(data.frame(y=round(mean(x),0),label=round(mean(x,na.rm=T),0)))
}
show_box_plot_duration= function(df, title) {
  df$treatment_group <- as.factor(df$condition)
  p = ggplot(df, aes(x=treatment_group, y=duration_of_survey, fill=treatment_group)) + geom_boxplot() +
    scale_fill_manual(values=c("#a69da0", "#f95b3a")) + labs(title=title, x="Control vs. Treatment", y =
      width = 1, linetype = "dashed") + stat_summary(fun.data = meanFunction1, geom="text", s

  p + geom_jitter(shape=16, size=2, alpha=.3, position=position_jitter(0.2) )
}
```

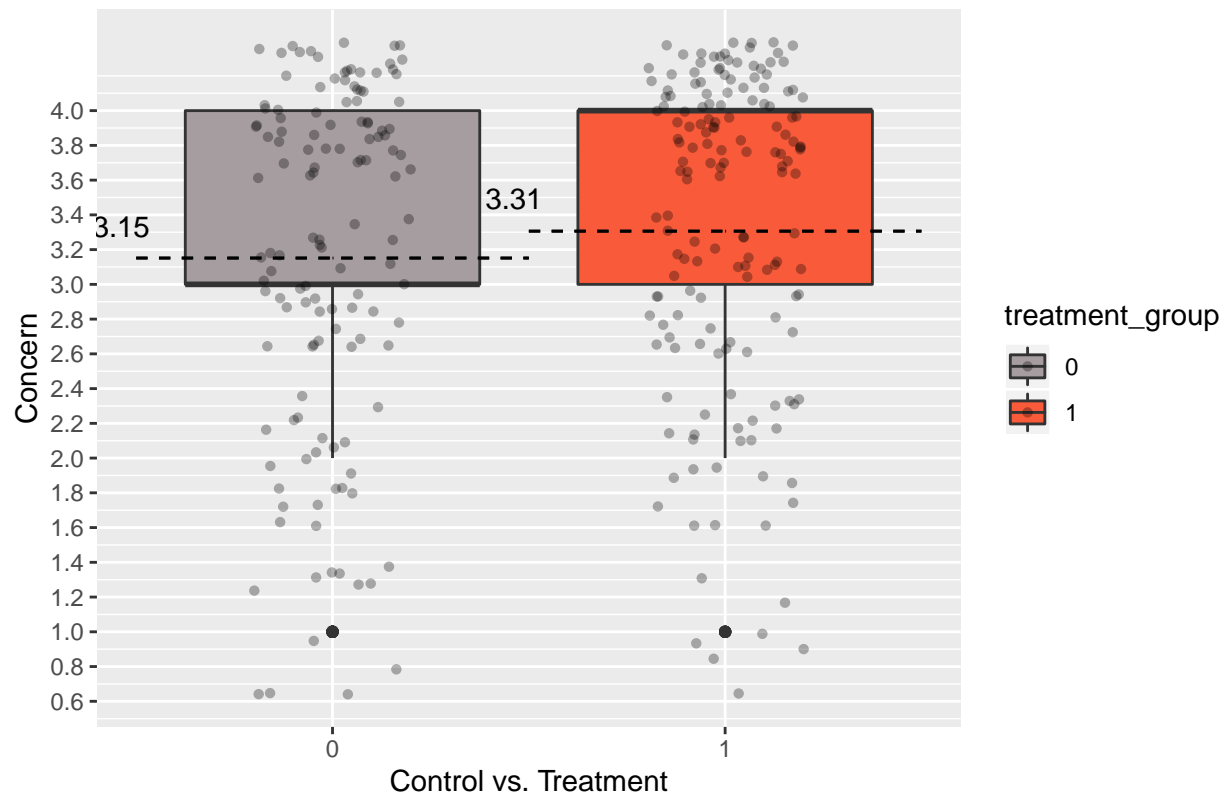
```
show_box_plot_spread(run1, "Concern over spread of COVID-19 in U.S. - Run 1")
```

Concern over spread of COVID-19 in U.S. – Run 1



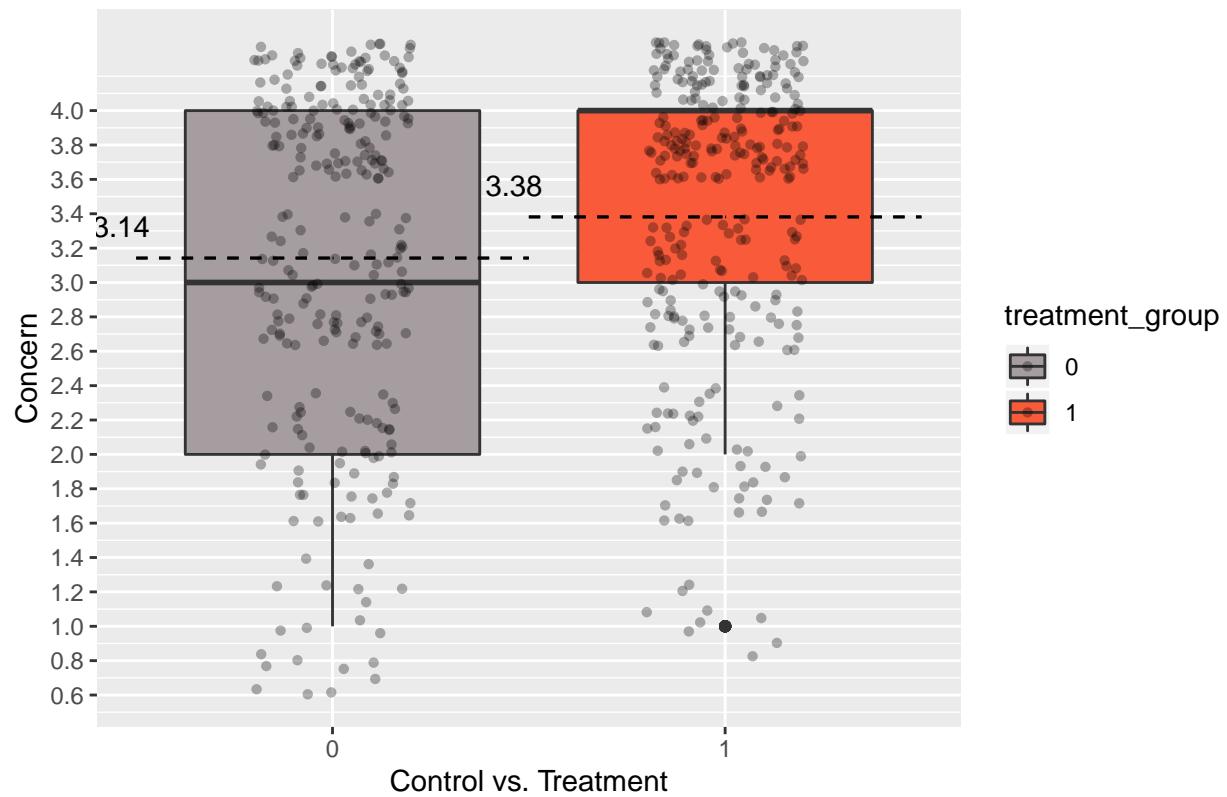
```
show_box_plot_spread(run2, "Concern over spread of COVID-19 in U.S. – Run 2")
```

Concern over spread of COVID-19 in U.S. – Run 2



```
show_box_plot_spread(combined, "Concern over spread of COVID-19 in U.S. - Both Runs")
```

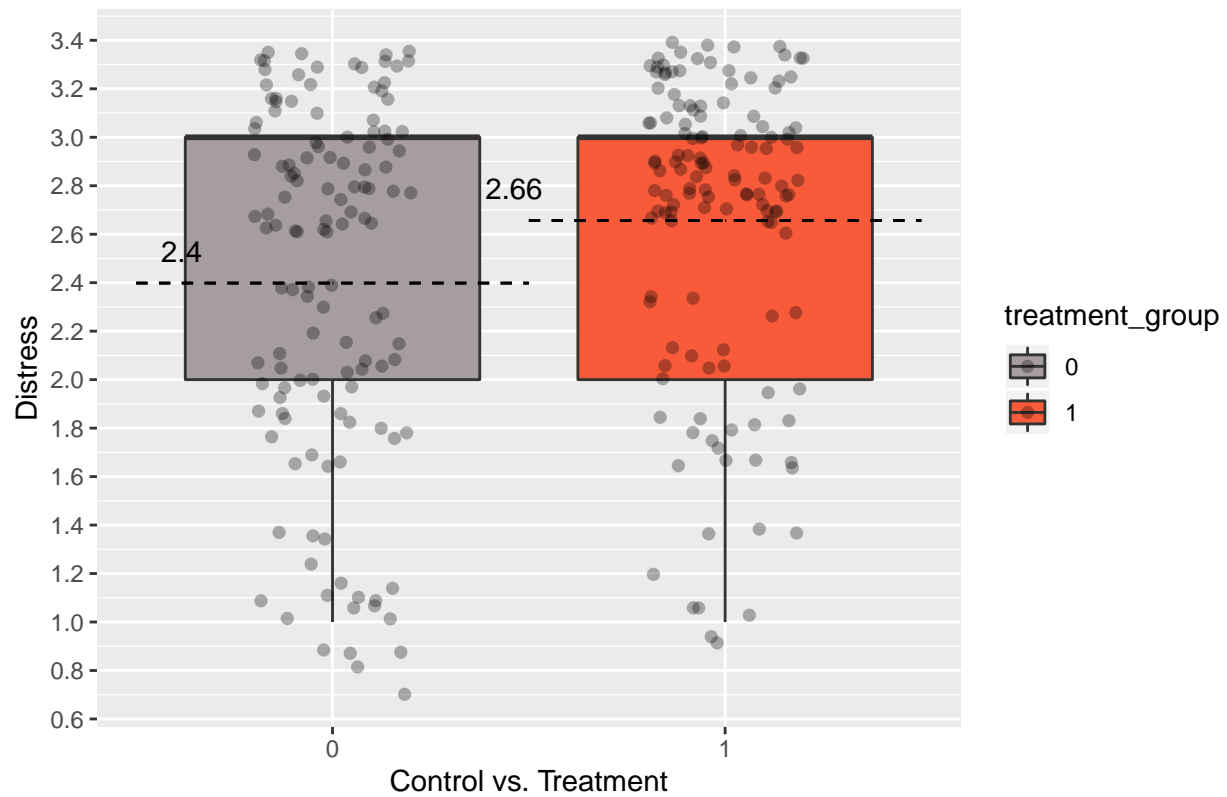

Concern over spread of COVID-19 in U.S. – Both Runs



```
show_box_plot_death(run1, "Distress over COVID-19 deaths in U.S. - Run 1")
```

```
## No summary function supplied, defaulting to 'mean_se()'
```

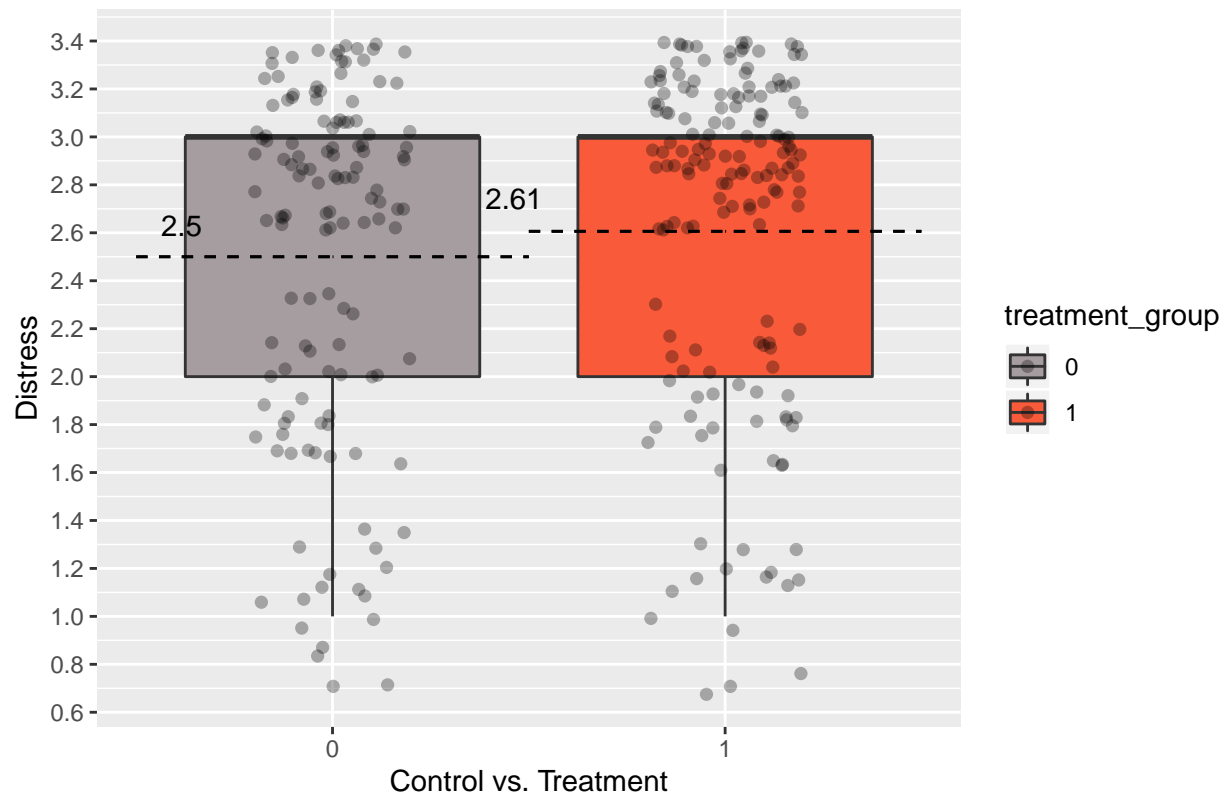
Distress over COVID-19 deaths in U.S. – Run 1



```
show_box_plot_death(run2, "Distress over COVID-19 deaths in U.S. - Run 2")
```

```
## No summary function supplied, defaulting to 'mean_se()
```

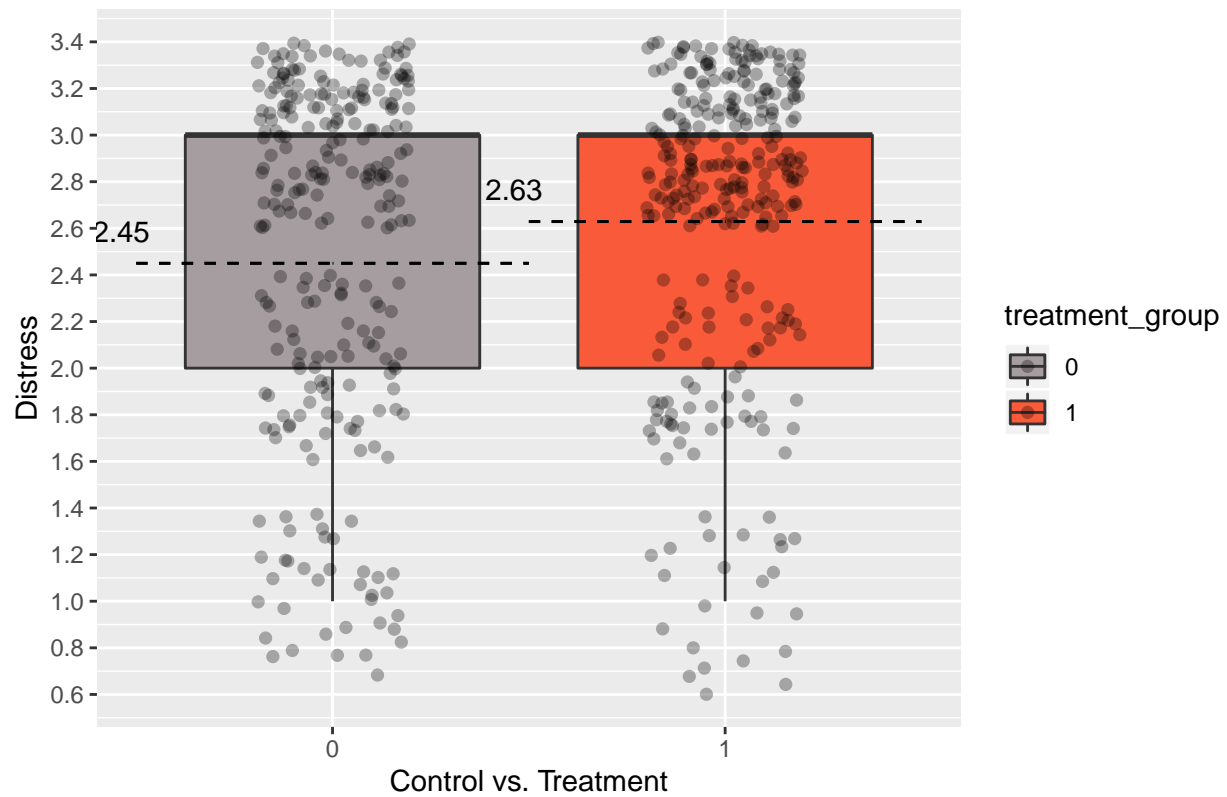
Distress over COVID-19 deaths in U.S. – Run 2



```
show_box_plot_death(combined, "Distress over COVID-19 deaths in U.S. - Both runs")
```

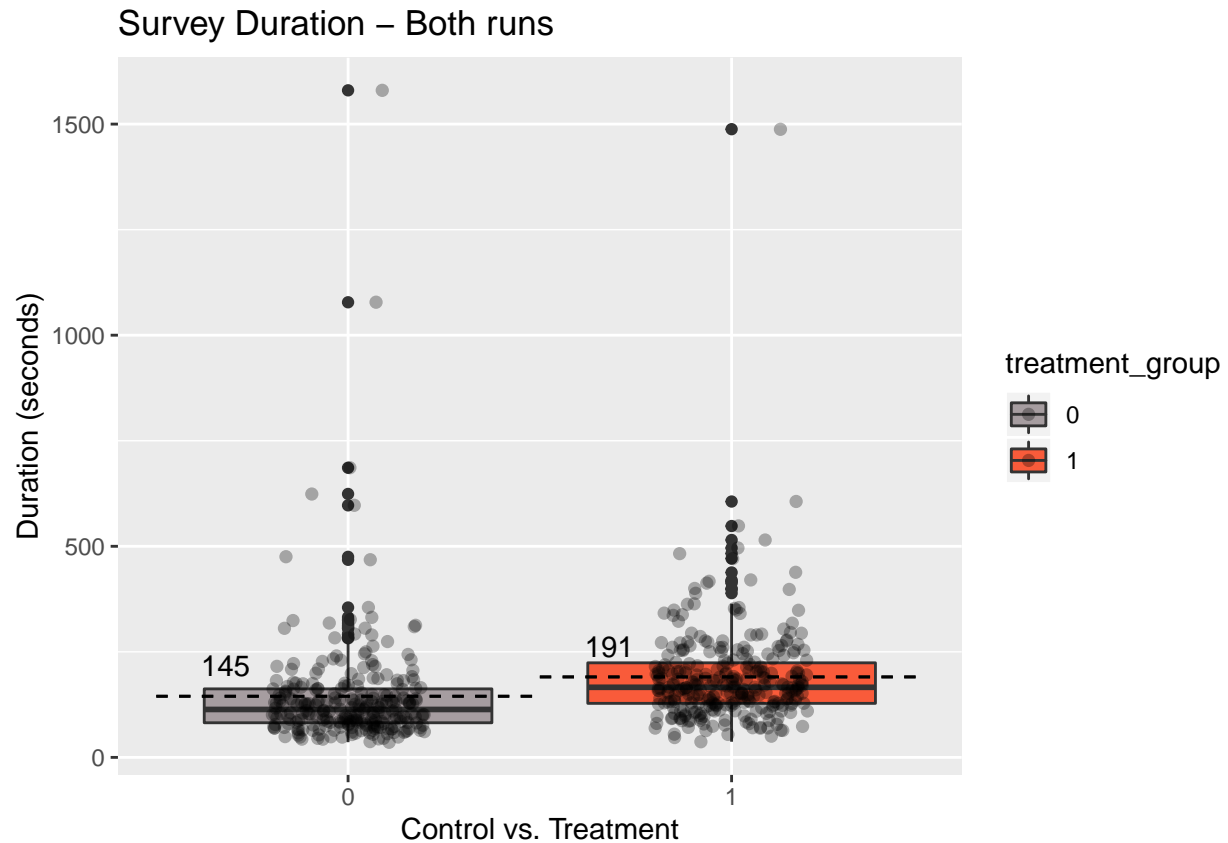
```
## No summary function supplied, defaulting to 'mean_se()'
```

Distress over COVID-19 deaths in U.S. – Both runs



```
show_box_plot_duration(combined, "Survey Duration - Both runs")
```

```
## No summary function supplied, defaulting to 'mean_se()'
```



```
cat(combined[treatment == 0, mean(duration_of_survey)], '\n')
```

```
## 144.8654
```

```
cat(combined[treatment == 1, mean(duration_of_survey)], '\n')
```

```
## 190.5567
```

5. Linear Regression

5.1 Perform linear regression the two outcomes (concern about COVID-19 spread, concern about COVID-19 deaths)

```
run_regression_outcome1 = function(d, control_for_run=FALSE) {
  model_spread = lm(outcome_spread ~ condition, d)
  if (control_for_run) {
    model_spread_adv = lm(outcome_spread ~ condition
      + run
      + as.factor(gender)
      + as.factor(age)
      + as.factor(ethnicity)
      + as.factor(political_party)
```

```

        + as.factor(education)
        + as.factor(covid_sick)
        + as.factor(covid_hospitalized)
        + as.factor(covid_died), d)

} else {
  model_spread_adv = lm(outcome_spread ~ condition
    + as.factor(gender)
    + as.factor(age)
    + as.factor(ethnicity)
    + as.factor(political_party)
    + as.factor(education)
    + as.factor(covid_sick)
    + as.factor(covid_hospitalized)
    + as.factor(covid_died), d)

}
est_spread      = get_regression_results_robust_se(model_spread, d, c('condition'), FALSE)
est_spread_adv = get_regression_results_robust_se(model_spread_adv, d, c('condition'), FALSE)
return ( list('model'= model_spread,
              'model_adv'=model_spread_adv,
              'est'=est_spread,
              'est_adv'=est_spread_adv))
}

run_regression_outcome2 = function(d, control_for_run) {
  model_spread = lm(outcome_spread ~ condition, d)
  if (control_for_run) {
    model_spread_adv = lm(outcome_death ~ condition
      + run
      + as.factor(gender)
      + as.factor(age)
      + as.factor(ethnicity)
      + as.factor(political_party)
      + as.factor(education)
      + as.factor(covid_sick)
      + as.factor(covid_hospitalized)
      + as.factor(covid_died), d)

  } else {
    model_spread_adv = lm(outcome_death ~ condition
      + as.factor(gender)
      + as.factor(age)
      + as.factor(ethnicity)
      + as.factor(political_party)
      + as.factor(education)
      + as.factor(covid_sick)
      + as.factor(covid_hospitalized)
      + as.factor(covid_died), d)

  }
}

```

```

est_spread      = get_regression_results_robust_se(model_spread, d, c('condition'), FALSE)
est_spread_adv  = get_regression_results_robust_se(model_spread_adv, d, c('condition'), FALSE)
return ( list('model'= model_spread,
              'model_adv'=model_spread_adv,
              'est'=est_spread,
              'est_adv'=est_spread_adv))
}

aggregate_covariates = function(df) {
  df[, college_educated := ifelse(education > 4, 1, 0)]
  df[, caucasian := ifelse(ethnicity == 3, 1, 0)]
  df[, age_under_40 := ifelse(age < 4, 1, 0)]
  df[, age_40_60 := ifelse(age > 4 & age <= 6, 1, 0)]
  df[, age_over_60 := ifelse(age >= 7, 1, 0)]
}

```

```

regression_labels_run = c('Treatment',
  'Run',
  'Female', 'Non-binary', 'Gender not answered',
  '20-29', '30-39', '40-49', '50-59', '60-69', '70-79', 'Over 80',
  'Black/African American', 'Caucasian', 'Hispanic/Latinx',
  'Native American', 'Pacific Islander', 'Ethnicity not answered',
  'Democrat', 'Independent', 'Party other',
  'High school', 'Some college', 'Associates', 'Bachelors', 'Masters', 'Doctoral', 'JD/MD',
  'Sick from COVID-19',
  'Hospitalized from COVID-19',
  'Died COVID-19')

```

```

mi_spread_run1 = run_regression_outcome1(run1, TRUE)
mi_spread_run2 = run_regression_outcome1(run2, TRUE)
mi_spread_combined_run = run_regression_outcome1(combined, TRUE)

```

```

stargazer(mi_spread_run1$model, mi_spread_run1$model_adv,
  mi_spread_run2$model_adv,
  mi_spread_combined_run$model_adv,
  type="text",
  se = list(mi_spread_run1$est$robust_se_all, mi_spread_run1$est_adv$robust_se_all,
            mi_spread_run2$est_adv$robust_se_all,
            mi_spread_combined_run$est_adv$robust_se_all),
  title=paste('Response to COVID-19 Spread'),
  dep.var.caption = "Response to COVID-19 Spread",
  dep.var.labels.include = FALSE, model.numbers=FALSE,
  column.labels = c("July 21, 2020", "July 21, 2020", "July 25, 2020", "Both dates"),
  align=TRUE,
  covariate.labels = regression_labels_run)

```

```
##
```

```
## Response to COVID-19 Spread
```

```
## =====
```

	Response to COVID-19 Spread		
	July 21, 2020	July 21, 2020	July 25, 2020
Treatment	0.340*** (0.107)	0.339*** (0.106)	0.172* (0.095)
Run			
Female		-0.063 (0.105)	0.187* (0.107)
Non-binary		0.886** (0.420)	1.742*** (0.409)
Gender not answered		0.646*** (0.192)	0.629** (0.292)
20-29		0.323 (0.380)	-0.454* (0.234)
30-39		0.295 (0.381)	-0.209 (0.264)
40-49		0.181 (0.402)	-0.460* (0.272)
50-59		0.370 (0.412)	-0.246 (0.255)
60-69		0.393 (0.441)	0.096 (0.267)
70-79		1.156*** (0.437)	-0.458 (0.336)
Over 80			1.121** (0.463)
Black/African American		0.195 (0.201)	-0.100 (0.217)
Caucasian		-0.087 (0.165)	0.060 (0.187)
Hispanic/Latinx		0.230 (0.205)	0.184 (0.252)
Native American		-0.471 (0.513)	0.482 (0.316)
Pacific Islander		-2.519*** (0.232)	

##			
## Ethnicity not answered		0.715**	-0.121
##		(0.304)	(0.585)
##			
## Democrat		0.675***	1.118***
##		(0.141)	(0.134)
##			
## Independent		0.357**	0.602***
##		(0.171)	(0.162)
##			
## Party other		-0.222	-0.318
##		(0.370)	(0.376)
##			
## High school		-0.050	
##		(0.800)	
##			
## Some college		0.194	0.135
##		(0.791)	(0.191)
##			
## Associates		0.017	-0.020
##		(0.795)	(0.207)
##			
## Bachelors		0.188	0.125
##		(0.781)	(0.174)
##			
## Masters		0.337	0.223
##		(0.786)	(0.195)
##			
## Doctoral		-0.166	0.029
##		(0.780)	(0.337)
##			
## JD/MD		0.502	0.842**
##		(0.778)	(0.339)
##			
## Sick from COVID-19		-0.200*	-0.162
##		(0.119)	(0.117)
##			
## Hospitalized from COVID-19		-0.166	-0.056
##		(0.165)	(0.161)
##			
## Died COVID-19		0.092	-0.110
##		(0.158)	(0.157)
##			
## Constant	3.133***	2.450***	2.660***
##	(0.085)	(0.896)	(0.321)
##			
## -----			
## Observations	259	259	292
## R2	0.038	0.314	0.351
## Adjusted R2	0.034	0.227	0.282
## Residual Std. Error	0.858 (df = 257)	0.768 (df = 229)	0.796 (df = 263)
## F Statistic	10.183*** (df = 1; 257)	3.614*** (df = 29; 229)	5.084*** (df = 28; 263)
## =====			
## Note:			*p<0.

```

mi_death_run1 = run_regression_outcome2(run1, TRUE)
mi_death_run2 = run_regression_outcome2(run2, TRUE)
mi_death_combined_run = run_regression_outcome2(combined, TRUE)

stargazer(mi_death_run1$model, mi_death_run1$model_adv,
  mi_death_run2$model_adv,
  mi_death_combined_run$model_adv,
  type="text",
  se = list(mi_death_run1$est$robust_se_all, mi_death_run1$est_adv$robust_se_all,
    mi_death_run2$est_adv$robust_se_all,
    mi_death_combined_run$est_adv$robust_se_all),
  title=paste('Response to COVID-19 Death'),
  dep.var.caption = "Response to COVID-19 Death",
  dep.var.labels.include = FALSE, model.numbers=FALSE,
  column.labels = c("July 21, 2020", "July 21, 2020", "July 25, 2020", "Both dates"),
  align=TRUE,
  covariate.labels = regression_labels_run)

```

```

##
## Response to COVID-19 Death
## =====
##                                     Response to COVID-19 Death
##                                     -----
##                                     July 21, 2020      July 21, 2020      July 25, 2020
## -----
## Treatment                0.340***                0.250***                0.109
##                           (0.107)                  (0.090)                  (0.072)
##
## Run
##
##
## Female                    0.050                    0.082
##                           (0.085)                  (0.079)
##
## Non-binary                0.573*                    0.899***
##                           (0.308)                  (0.308)
##
## Gender not answered       0.437***                0.454**
##                           (0.159)                  (0.207)
##
## 20-29                     -0.371*                  -0.083
##                           (0.191)                  (0.175)
##
## 30-39                     -0.323*                    0.071
##                           (0.186)                  (0.190)
##
## 40-49                     -0.345*                    0.142
##                           (0.209)                  (0.195)
##
## 50-59                     -0.258                    0.153
##                           (0.211)                  (0.195)
##

```

## 60-69	-0.279	0.370*
##	(0.260)	(0.213)
##		
## 70-79	0.170	-0.288
##	(0.254)	(0.295)
##		
## Over 80		0.873***
##		(0.256)
##		
## Black/African American	0.041	-0.022
##	(0.168)	(0.147)
##		
## Caucasian	-0.103	-0.016
##	(0.134)	(0.118)
##		
## Hispanic/Latinx	-0.019	0.193
##	(0.212)	(0.160)
##		
## Native American	-0.261	-0.020
##	(0.289)	(0.218)
##		
## Pacific Islander	-1.667***	
##	(0.176)	
##		
## Ethnicity not answered	0.374	-0.169
##	(0.239)	(0.291)
##		
## Democrat	0.543***	0.811***
##	(0.117)	(0.108)
##		
## Independent	0.340**	0.414***
##	(0.134)	(0.132)
##		
## Party other	-0.107	0.141
##	(0.302)	(0.291)
##		
## High school	0.233	
##	(0.530)	
##		
## Some college	0.216	0.040
##	(0.516)	(0.137)
##		
## Associates	0.135	-0.095
##	(0.524)	(0.166)
##		
## Bachelors	0.088	-0.071
##	(0.512)	(0.133)
##		
## Masters	0.263	-0.036
##	(0.517)	(0.142)
##		
## Doctoral	-0.475	0.041
##	(0.512)	(0.141)
##		

```
## JD/MD                                0.290                0.469*
##                                (0.513)                (0.239)
##
## Sick from COVID-19                  -0.102                -0.097
##                                (0.098)                (0.084)
##
## Hospitalized from COVID-19          -0.084                0.003
##                                (0.125)                (0.116)
##
## Died COVID-19                       0.067                -0.090
##                                (0.128)                (0.118)
##
## Constant                           3.133***                2.327***                2.017***
##                                (0.085)                (0.570)                (0.239)
## -----
## Observations                        259                259                292
## R2                                0.038                0.252                0.325
## Adjusted R2                       0.034                0.157                0.254
## Residual Std. Error                0.858 (df = 257)        0.628 (df = 229)        0.591 (df = 263)
## F Statistic                       10.183*** (df = 1; 257)  2.657*** (df = 29; 229)  4.533*** (df = 28; 263) 5
## =====
## Note:                                                                    *p<0.
```

```
state_ordinals <- fread('data/state_ordinals.csv')
states_northeast = c('Connecticut','Maine','Massachusetts','New Hampshire','Rhode Island','Vermont','New
states_midwest  = c('Illinois','Indiana','Michigan','Ohio','Wisconsin', 'Iowa', 'Kansas', 'Minnesota', 'Ne
states_south    = c('Delaware', 'Florida', 'Georgia', 'Maryland', 'North Carolina', 'South Carolina', 'Virg
states_west     = c('Arizona', 'Colorado', 'Idaho', 'Montana', 'Nevada', 'New Mexico', 'Utah', 'Wyoming', '
states_other    = c('Puerto Rico')

get_region_for_state = function(ordinal) {
  state_name = state_ordinals[ordinal, V1]
  if (is.element(state_name, states_northeast)) {
    return("northeast")
  } else if (is.element(state_name, states_midwest)) {
    return("midwest")
  } else if (is.element(state_name, states_south)) {
    return("south")
  } else if (is.element(state_name, states_west)) {
    return("west")
  } else if (is.element(state_name, states_other)) {
    return ("other")
  } else {
    return("other")
  }
}
get_region_for_state(5)
```

```
## [1] "west"
```

```

regions = c(rep('',53))
i = 1
for (ordinal in state_ordinals$V2) {
  the_region = get_region_for_state(ordinal)
  cat(the_region, i, '\n')
  regions[i] = the_region
  i = i + 1
}

```

```

## south 1
## west 2
## west 3
## south 4
## west 5
## west 6
## northeast 7
## south 8
## south 9
## south 10
## south 11
## west 12
## west 13
## midwest 14
## midwest 15
## midwest 16
## midwest 17
## south 18
## south 19
## northeast 20
## south 21
## northeast 22
## midwest 23
## midwest 24
## south 25
## midwest 26
## west 27
## midwest 28
## west 29
## northeast 30
## northeast 31
## west 32
## northeast 33
## south 34
## midwest 35
## midwest 36
## south 37
## west 38
## northeast 39
## other 40
## northeast 41
## south 42
## midwest 43
## south 44

```

```
## south 45
## west 46
## northeast 47
## south 48
## west 49
## south 50
## midwest 51
## west 52
## other 53
```

```
state_ordinals[, region:= regions]
```

```
state_ordinals
```

```
##           V1 V2   region
## 1:      Alabama 1   south
## 2:      Alaska 2   west
## 3:      Arizona 3   west
## 4:      Arkansas 4  south
## 5:      California 5  west
## 6:      Colorado 6  west
## 7:      Connecticut 7 northeast
## 8: District of Columbia 8  south
## 9:      Delaware 9  south
## 10:     Florida 10  south
## 11:     Georgia 11  south
## 12:     Hawaii 12  west
## 13:     Idaho 13  west
## 14:     Illinois 14  midwest
## 15:     Indiana 15  midwest
## 16:     Iowa 16  midwest
## 17:     Kansas 17  midwest
## 18:     Kentucky 18  south
## 19:     Louisiana 19  south
## 20:     Maine 20  northeast
## 21:     Maryland 21  south
## 22:     Massachusetts 22 northeast
## 23:     Michigan 23  midwest
## 24:     Minnesota 24  midwest
## 25:     Mississippi 25  south
## 26:     Missouri 26  midwest
## 27:     Montana 27  west
## 28:     Nebraska 28  midwest
## 29:     Nevada 29  west
## 30:     New Hampshire 30 northeast
## 31:     New Jersey 31 northeast
## 32:     New Mexico 32  west
## 33:     New York 33  northeast
## 34:     North Carolina 34  south
## 35:     North Dakota 35  midwest
## 36:     Ohio 36  midwest
## 37:     Oklahoma 37  south
## 38:     Oregon 38  west
## 39:     Pennsylvania 39 northeast
```

```
## 40:      Puerto Rico 40      other
## 41:      Rhode Island 41 northeast
## 42:      South Carolina 42      south
## 43:      South Dakota 43      midwest
## 44:      Tennessee 44      south
## 45:      Texas 45      south
## 46:      Utah 46      west
## 47:      Vermont 47 northeast
## 48:      Virginia 48      south
## 49:      Washington 49      west
## 50:      West Virginia 50      south
## 51:      Wisconsin 51      midwest
## 52:      Wyoming 52      west
## 53:      Not in US 53      other
##              V1 V2      region
```

```
aggregate_covariates = function(df) {
  df[, gender_male := ifelse(gender == 1, 1, 0)]
  df[, gender_female := ifelse(gender == 2, 1, 0)]
  df[, gender_other := ifelse(gender == 3 | gender == 4, 1, 0)]
  df[, college_educated := ifelse(education > 4, 1, 0)]
  df[, not_caucasian := ifelse(ethnicity != 3, 1, 0)]
  df[, age_under_40 := ifelse(age < 4, 1, 0)]
  df[, age_40_60 := ifelse(age > 4 & age <= 6, 1, 0)]
  df[, age_over_60 := ifelse(age >= 7, 1, 0)]
  df[, party_republican := ifelse(political_party == 1, 1, 0)]
  df[, party_democrat := ifelse(political_party == 2, 1, 0)]
  df[, party_other := ifelse(political_party == 3 | political_party == 4, 1, 0)]
  df[, party_non_republican := ifelse( political_party == 1, 0, 1)]
  df[, treatment_assignment := condition]
  df[, state_name := state_ordinals[state, V1]]
  df[, region:= state_ordinals[state, region]]
  df[, region_northeast := ifelse(region == 'northeast', 1,0)]
  df[, region_midwest := ifelse(region == 'midwest', 1,0)]
  df[, region_south := ifelse(region == 'south', 1,0)]
  df[, region_west := ifelse(region == 'west', 1,0)]
  df[, region_other := ifelse(region == 'other', 1,0)]
}

aggregate_covariates(run1)
aggregate_covariates(run2)
aggregate_covariates(combined)

head(combined[, c('state', 'state_name', 'region', 'region_northeast', 'region_midwest')])
```

```
##      state  state_name  region region_northeast region_midwest
## 1:     36      Ohio  midwest              0              1
## 2:     39 Pennsylvania northeast              1              0
## 3:     38      Oregon   west              0              0
## 4:     45      Texas   south              0              0
## 5:     13      Idaho   west              0              0
## 6:     36      Ohio  midwest              0              1
```

```
nrow(combined[covid_died == 1,])
```

```
## [1] 80
```

```
run_regression_outcome1_simplified = function(d, control_for_run=FALSE) {
  model_spread = lm(outcome_spread ~ treatment_assignment, d)
  if (control_for_run) {
    model_spread_adv = lm(outcome_spread ~ treatment_assignment
      + run
      + gender_female
      + gender_other
      + age_40_60
      + age_over_60
      + party_democrat
      + party_other
      + not_caucasian
      + college_educated
      + region_south
      + region_midwest
      + region_west
      + covid_sick
      + covid_hospitalized
      + covid_died, d)
  } else {
    model_spread_adv = lm(outcome_spread ~ treatment_assignment
      + gender_female
      + gender_other
      + age_under_40
      + age_40_60
      + age_over_60
      + party_democrat
      + party_other
      + not_caucasian
      + college_educated
      + region_south
      + region_midwest
      + region_west
      + covid_sick
      + covid_hospitalized
      + covid_died, d)
  }
  est_spread      = get_regression_results_robust_se(model_spread, d, c('treatment_assignment'), FALSE)
  est_spread_adv = get_regression_results_robust_se(model_spread_adv, d, c('treatment_assignment'), FALSE)
  return ( list('model'= model_spread,
    'model_adv'=model_spread_adv,
    'est'=est_spread,
    'est_adv'=est_spread_adv))
}

run_regression_outcome2_simplified = function(d, control_for_run=FALSE) {
  model_spread = lm(outcome_spread ~ treatment_assignment, d)
  if (control_for_run) {
```



```

model_spread_adv = lm(outcome_death ~ treatment_assignment
+ run
+ gender_female
+ gender_other
+ age_40_60
+ age_over_60
+ party_democrat
+ party_other
+ not_caucasian
+ college_educated
+ region_south
+ region_midwest
+ region_west
+ covid_sick
+ covid_hospitalized
+ covid_died, d)

} else {
  model_spread_adv = lm(outcome_death ~ treatment_assignment
+ gender_female
+ gender_other
+ age_40_60
+ age_over_60
+ party_democrat
+ party_other
+ not_caucasian
+ college_educated
+ region_south
+ region_midwest
+ region_west
+ covid_sick
+ covid_hospitalized
+ covid_died, d)
}
est_spread      = get_regression_results_robust_se(model_spread, d, c('treatment_assignment'), FALSE)
est_spread_adv = get_regression_results_robust_se(model_spread_adv, d, c('treatment_assignment'), FALSE)
return ( list('model'= model_spread,
              'model_adv'=model_spread_adv,
              'est'=est_spread,
              'est_adv'=est_spread_adv))
}

```

```

mi_spread_run1 = run_regression_outcome1_simplified(run1, TRUE)
mi_spread_run2 = run_regression_outcome1_simplified(run2, TRUE)
mi_spread_combined_run = run_regression_outcome1_simplified(combined, TRUE)

stargazer(mi_spread_run1$model, mi_spread_run1$model_adv,
mi_spread_run2$model_adv,
mi_spread_combined_run$model_adv,
type="text", out="output/model1_regression.html", report="vcsp*",
se = list(mi_spread_run1$est$robust_se_all, mi_spread_run1$est_adv$robust_se_all,
mi_spread_run2$est_adv$robust_se_all,

```

```

mi_spread_combined_run$est_adv$robust_se_all),
title=paste('Concerned over COVID-19 Spread in U.S. '),
dep.var.caption = "Concerned over COVID-19 Spread in U.S.",
dep.var.labels.include = FALSE, model.numbers=FALSE,
column.labels = c("July 21, 2020", "July 21, 2020", "July 25, 2020", "Both dates"),
align=TRUE)

```

```

##
## Concerned over COVID-19 Spread in U.S.
## =====
##                               Concerned over COVID-19 Spread in U.S.
##                               -----
##                               July 21, 2020      July 21, 2020      July 25, 2020
## -----
## treatment_assignment          0.340            0.334            0.146
##                               (0.107)          (0.104)          (0.096)
##                               p = 0.002***      p = 0.002***      p = 0.128
##
## run
##
##
## gender_female                 -0.091           0.166
##                               (0.105)          (0.106)
##                               p = 0.388          p = 0.117
##
## gender_other                  0.645           0.930
##                               (0.214)          (0.248)
##                               p = 0.003***      p = 0.0002***
##
## age_40_60                    0.143           0.232
##                               (0.125)          (0.112)
##                               p = 0.254          p = 0.040**
##
## age_over_60                  0.940           0.017
##                               (0.221)          (0.259)
##                               p = 0.00003***     p = 0.947
##
## party_democrat               0.681           1.053
##                               (0.133)          (0.132)
##                               p = 0.00000***     p = 0.000***
##
## party_other                   0.261           0.438
##                               (0.167)          (0.156)
##                               p = 0.118          p = 0.006***
##
## not_caucasian                0.171           -0.052
##                               (0.104)          (0.109)
##                               p = 0.099*         p = 0.635
##
## college_educated             0.153           0.071
##                               (0.102)          (0.102)
##                               p = 0.134          p = 0.491

```

```
##
## region_south                0.054                0.044
##                            (0.137)              (0.141)
##                            p = 0.695            p = 0.755
##
## region_midwest              0.098                -0.206
##                            (0.147)              (0.175)
##                            p = 0.509            p = 0.239
##
## region_west                 0.280                0.077
##                            (0.139)              (0.166)
##                            p = 0.045**          p = 0.643
##
## covid_sick                  -0.225               -0.176
##                            (0.120)              (0.118)
##                            p = 0.062*          p = 0.137
##
## covid_hospitalized          -0.186               -0.070
##                            (0.155)              (0.156)
##                            p = 0.231            p = 0.657
##
## covid_died                  0.051                -0.117
##                            (0.153)              (0.160)
##                            p = 0.741            p = 0.465
##
## Constant                    3.133                3.057                2.922
##                            (0.085)              (0.283)              (0.306)
##                            p = 0.000***          p = 0.000***          p = 0.000***
##
## -----
## Observations                259                259                292
## R2                          0.038                0.263                0.291
## Adjusted R2                 0.034                0.218                0.252
## Residual Std. Error        0.858 (df = 257)        0.773 (df = 243)        0.812 (df = 276)
## F Statistic                 10.183*** (df = 1; 257)  5.782*** (df = 15; 243)  7.541*** (df = 15; 276)
## =====
## Note:                                                                *p<0.1; **]
```

```
mi_spread_run1 = run_regression_outcome2_simplified(run1, TRUE)
mi_spread_run2 = run_regression_outcome2_simplified(run2, TRUE)
mi_spread_combined_run = run_regression_outcome2_simplified(combined, TRUE)

stargazer(mi_spread_run1$model, mi_spread_run1$model_adv,
  mi_spread_run2$model_adv,
  mi_spread_combined_run$model_adv,
  type="html", out="output/model2_regression.html", report="vcsp",
  se = list(mi_spread_run1$est$robust_se_all, mi_spread_run1$est_adv$robust_se_all,
    mi_spread_run2$est_adv$robust_se_all,
    mi_spread_combined_run$est_adv$robust_se_all),
  title=paste('Distressed by COVID-19 Deaths in U.S. '),
  dep.var.caption = "Distressed by COVID-19 Deaths in U.S.",
  dep.var.labels.include = FALSE, model.numbers=FALSE,
  column.labels = c("July 21, 2020", "July 21, 2020", "July 25, 2020", "Both dates"),
```



```

## <tr><td style="text-align:left"></td><td></td><td>p = 0.644</td><td>p = 0.204</td><td>p = 0.437</td>
## <tr><td style="text-align:left"></td><td></td><td></td><td></td><td></td></tr>
## <tr><td style="text-align:left">region_west</td><td></td><td>0.030</td><td>0.076</td><td>0.049</td><
## <tr><td style="text-align:left"></td><td></td><td>(0.120)</td><td>(0.123)</td><td>(0.086)</td></tr>
## <tr><td style="text-align:left"></td><td></td><td>p = 0.806</td><td>p = 0.536</td><td>p = 0.568</td>
## <tr><td style="text-align:left"></td><td></td><td></td><td></td><td></td></tr>
## <tr><td style="text-align:left">covid_sick</td><td></td><td>-0.093</td><td>-0.110</td><td>-0.111</td>
## <tr><td style="text-align:left"></td><td></td><td>(0.096)</td><td>(0.082)</td><td>(0.062)</td></tr>
## <tr><td style="text-align:left"></td><td></td><td>p = 0.330</td><td>p = 0.181</td><td>p = 0.075<sup>
## <tr><td style="text-align:left"></td><td></td><td></td><td></td><td></td></tr>
## <tr><td style="text-align:left">covid_hospitalized</td><td></td><td>-0.111</td><td>0.031</td><td>-0.
## <tr><td style="text-align:left"></td><td></td><td>(0.114)</td><td>(0.112)</td><td>(0.077)</td></tr>
## <tr><td style="text-align:left"></td><td></td><td>p = 0.332</td><td>p = 0.783</td><td>p = 0.752</td>
## <tr><td style="text-align:left"></td><td></td><td></td><td></td><td></td></tr>
## <tr><td style="text-align:left">covid_died</td><td></td><td>0.054</td><td>-0.106</td><td>-0.056</td>
## <tr><td style="text-align:left"></td><td></td><td>(0.123)</td><td>(0.119)</td><td>(0.082)</td></tr>
## <tr><td style="text-align:left"></td><td></td><td>p = 0.664</td><td>p = 0.371</td><td>p = 0.497</td>
## <tr><td style="text-align:left"></td><td></td><td></td><td></td><td></td></tr>
## <tr><td style="text-align:left">Constant</td><td></td><td>3.133</td><td>2.255</td><td>2.192</td><td>2.310</td>
## <tr><td style="text-align:left"></td><td></td><td>(0.085)</td><td>(0.242)</td><td>(0.225)</td><td>(0.165)</td>
## <tr><td style="text-align:left"></td><td></td><td>p = 0.000<sup>***</sup></td><td>p = 0.000<sup>***</sup></td>
## <tr><td style="text-align:left"></td><td></td><td></td><td></td><td></td></tr>
## <tr><td colspan="5" style="border-bottom: 1px solid black"></td></tr><tr><td style="text-align:left".
## <tr><td style="text-align:left">R<sup>2</sup></td><td></td><td>0.038</td><td>0.198</td><td>0.295</td><td>0.22
## <tr><td style="text-align:left">Adjusted R<sup>2</sup></td><td></td><td>0.034</td><td>0.149</td><td>0.257</td>
## <tr><td style="text-align:left">Residual Std. Error</td><td></td><td>0.858 (df = 257)</td><td>0.631 (df = 243
## <tr><td style="text-align:left">F Statistic</td><td></td><td>10.183<sup>***</sup> (df = 1; 257)</td><td>4.004
## <tr><td colspan="5" style="border-bottom: 1px solid black"></td></tr><tr><td style="text-align:left".
## </table>

```

```

run_regression_outcome1_hte = function(d) {

  model_spread_adv = lm(outcome_spread ~ treatment_assignment
                        + gender_female
                        + gender_other
                        + age_40_60
                        + age_over_60
                        + party_democrat
                        + party_other
                        + party_other:treatment_assignment
                        + not_caucasian
                        + college_educated
                        + region_south
                        + region_midwest
                        + region_west
                        + covid_sick
                        + covid_hospitalized
                        + covid_died, d)

  est_spread_adv = get_regression_results_robust_se(model_spread_adv, d, c('treatment_assignment'), FALSE)
  return ( list('model_adv'=model_spread_adv,
                'est_adv'=est_spread_adv))
}

```

```

run_regression_outcome2_hte = function(d) {

  model_spread_adv = lm(outcome_death ~ treatment_assignment
                        + gender_female
                        + gender_other
                        + age_40_60
                        + age_over_60
                        + party_democrat
                        + party_other
                        + party_other:treatment_assignment
                        + not_caucasian
                        + college_educated
                        + region_south
                        + region_midwest
                        + region_west
                        + covid_sick
                        + covid_hospitalized
                        + covid_died, d)

  est_spread_adv = get_regression_results_robust_se(model_spread_adv, d, c('treatment_assignment'), FALSE)
  return ( list('model_adv'=model_spread_adv,
                'est_adv'=est_spread_adv))
}

```

```

mi_spread_combined_run = run_regression_outcome1_hte(combined)
mi_death_combined_run  = run_regression_outcome2_hte(combined)

stargazer(mi_spread_combined_run$model_adv,
           #type = "text",
           type="html", out="output/model_combined_spread_hte_regression.html", report="vcsp*",
           se = list(mi_spread_combined_run$est_adv$robust_se_all),
           dep.var.caption = "Concerned over COVID-19 Spread in U.S. - HTE",
           dep.var.labels.include = FALSE, model.numbers=FALSE,
           column.labels = c("Both dates"),
           align=TRUE)

```

```

##
## <table style="text-align:center"><tr><td colspan="2" style="border-bottom: 1px solid black"></td></tr>
## <tr><td></td><td colspan="1" style="border-bottom: 1px solid black"></td></tr>
## <tr><td style="text-align:left"></td><td>Both dates</td></tr>
## <tr><td colspan="2" style="border-bottom: 1px solid black"></td></tr><tr><td style="text-align:left">
## <tr><td style="text-align:left"></td><td>(0.078)</td></tr>
## <tr><td style="text-align:left"></td><td>p = 0.053<sup>*</sup></td></tr>
## <tr><td style="text-align:left"></td><td></td></tr>
## <tr><td style="text-align:left">gender_female</td><td>0.043</td></tr>
## <tr><td style="text-align:left"></td><td>(0.072)</td></tr>
## <tr><td style="text-align:left"></td><td>p = 0.555</td></tr>
## <tr><td style="text-align:left"></td><td></td></tr>
## <tr><td style="text-align:left">gender_other</td><td>0.802</td></tr>
## <tr><td style="text-align:left"></td><td>(0.149)</td></tr>
## <tr><td style="text-align:left"></td><td>p = 0.00000<sup>***</sup></td></tr>
## <tr><td style="text-align:left"></td><td></td></tr>

```

[illegible]

```
## <tr><td style="text-align:left"></td><td>p = 0.000<sup>***</sup></td></tr>
## <tr><td style="text-align:left"></td><td></td></tr>
## <tr><td colspan="2" style="border-bottom: 1px solid black"></td></tr><tr><td style="text-align:left">
## <tr><td style="text-align:left">R<sup>2</sup></td><td>0.251</td></tr>
## <tr><td style="text-align:left">Adjusted R<sup>2</sup></td><td>0.229</td></tr>
## <tr><td style="text-align:left">Residual Std. Error</td><td>0.798 (df = 534)</td></tr>
## <tr><td style="text-align:left">F Statistic</td><td>11.194<sup>***</sup> (df = 16; 534)</td></tr>
## <tr><td colspan="2" style="border-bottom: 1px solid black"></td></tr><tr><td style="text-align:left">
## </table>
```

```
stargazer(mi_death_combined_run$model_adv,
  #type = "text",
  type="html", out="output/model_combined_death_hte_regression.html", report="vcsp",
  se = list(mi_death_combined_run$est_adv$robust_se_all),
  dep.var.caption = "Distressed by COVID-19 Spread in U.S. - HTE",
  dep.var.labels.include = FALSE, model.numbers=FALSE,
  column.labels = c("Both dates"),
  align=TRUE)
```

```
##
## <table style="text-align:center"><tr><td colspan="2" style="border-bottom: 1px solid black"></td></tr>
## <tr><td></td><td colspan="1" style="border-bottom: 1px solid black"></td></tr>
## <tr><td style="text-align:left"></td><td>Both dates</td></tr>
## <tr><td colspan="2" style="border-bottom: 1px solid black"></td></tr><tr><td style="text-align:left">
## <tr><td style="text-align:left"></td><td>(0.061)</td></tr>
## <tr><td style="text-align:left"></td><td>p = 0.227</td></tr>
## <tr><td style="text-align:left"></td><td></td></tr>
## <tr><td style="text-align:left">gender_female</td><td>0.064</td></tr>
## <tr><td style="text-align:left"></td><td>(0.055)</td></tr>
## <tr><td style="text-align:left"></td><td>p = 0.242</td></tr>
## <tr><td style="text-align:left"></td><td></td></tr>
## <tr><td style="text-align:left">gender_other</td><td>0.553</td></tr>
## <tr><td style="text-align:left"></td><td>(0.106)</td></tr>
## <tr><td style="text-align:left"></td><td>p = 0.00000<sup>***</sup></td></tr>
## <tr><td style="text-align:left"></td><td></td></tr>
## <tr><td style="text-align:left">age_40_60</td><td>0.154</td></tr>
## <tr><td style="text-align:left"></td><td>(0.063)</td></tr>
## <tr><td style="text-align:left"></td><td>p = 0.015<sup>***</sup></td></tr>
## <tr><td style="text-align:left"></td><td></td></tr>
## <tr><td style="text-align:left">age_over_60</td><td>0.011</td></tr>
## <tr><td style="text-align:left"></td><td>(0.188)</td></tr>
## <tr><td style="text-align:left"></td><td>p = 0.954</td></tr>
## <tr><td style="text-align:left"></td><td></td></tr>
## <tr><td style="text-align:left">party_democrat</td><td>0.671</td></tr>
## <tr><td style="text-align:left"></td><td>(0.076)</td></tr>
## <tr><td style="text-align:left"></td><td>p = 0.000<sup>***</sup></td></tr>
## <tr><td style="text-align:left"></td><td></td></tr>
## <tr><td style="text-align:left">party_other</td><td>0.153</td></tr>
## <tr><td style="text-align:left"></td><td>(0.123)</td></tr>
## <tr><td style="text-align:left"></td><td>p = 0.216</td></tr>
## <tr><td style="text-align:left"></td><td></td></tr>
## <tr><td style="text-align:left">not_caucasian</td><td>0.057</td></tr>
## <tr><td style="text-align:left"></td><td>(0.058)</td></tr>
## <tr><td style="text-align:left"></td><td>p = 0.326</td></tr>
```



```

## <tr><td style="text-align:left"></td><td></td></tr>
## <tr><td style="text-align:left">college_educated</td><td>-0.062</td></tr>
## <tr><td style="text-align:left"></td><td>(0.054)</td></tr>
## <tr><td style="text-align:left"></td><td>p = 0.253</td></tr>
## <tr><td style="text-align:left"></td><td></td></tr>
## <tr><td style="text-align:left">region_south</td><td>0.052</td></tr>
## <tr><td style="text-align:left"></td><td>(0.072)</td></tr>
## <tr><td style="text-align:left"></td><td>p = 0.472</td></tr>
## <tr><td style="text-align:left"></td><td></td></tr>
## <tr><td style="text-align:left">region_midwest</td><td>-0.065</td></tr>
## <tr><td style="text-align:left"></td><td>(0.087)</td></tr>
## <tr><td style="text-align:left"></td><td>p = 0.454</td></tr>
## <tr><td style="text-align:left"></td><td></td></tr>
## <tr><td style="text-align:left">region_west</td><td>0.046</td></tr>
## <tr><td style="text-align:left"></td><td>(0.086)</td></tr>
## <tr><td style="text-align:left"></td><td>p = 0.593</td></tr>
## <tr><td style="text-align:left"></td><td></td></tr>
## <tr><td style="text-align:left">covid_sick</td><td>-0.108</td></tr>
## <tr><td style="text-align:left"></td><td>(0.061)</td></tr>
## <tr><td style="text-align:left"></td><td>p = 0.080<sup>*</sup></td></tr>
## <tr><td style="text-align:left"></td><td></td></tr>
## <tr><td style="text-align:left">covid_hospitalized</td><td>-0.025</td></tr>
## <tr><td style="text-align:left"></td><td>(0.076)</td></tr>
## <tr><td style="text-align:left"></td><td>p = 0.748</td></tr>
## <tr><td style="text-align:left"></td><td></td></tr>
## <tr><td style="text-align:left">covid_died</td><td>-0.048</td></tr>
## <tr><td style="text-align:left"></td><td>(0.081)</td></tr>
## <tr><td style="text-align:left"></td><td>p = 0.554</td></tr>
## <tr><td style="text-align:left"></td><td></td></tr>
## <tr><td style="text-align:left">treatment_assignment:party_other</td><td>0.247</td></tr>
## <tr><td style="text-align:left"></td><td>(0.130)</td></tr>
## <tr><td style="text-align:left"></td><td>p = 0.059<sup>*</sup></td></tr>
## <tr><td style="text-align:left"></td><td></td></tr>
## <tr><td style="text-align:left">Constant</td><td>2.318</td></tr>
## <tr><td style="text-align:left"></td><td>(0.164)</td></tr>
## <tr><td style="text-align:left"></td><td>p = 0.000<sup>***</sup></td></tr>
## <tr><td style="text-align:left"></td><td></td></tr>
## <tr><td colspan="2" style="border-bottom: 1px solid black"></td></tr><tr><td style="text-align:left">
## <tr><td style="text-align:left">R<sup>2</sup></td><td>0.234</td></tr>
## <tr><td style="text-align:left">Adjusted R<sup>2</sup></td><td>0.211</td></tr>
## <tr><td style="text-align:left">Residual Std. Error</td><td>0.607 (df = 534)</td></tr>
## <tr><td style="text-align:left">F Statistic</td><td>10.180<sup>***</sup> (df = 16; 534)</td></tr>
## <tr><td colspan="2" style="border-bottom: 1px solid black"></td></tr><tr><td style="text-align:left">
## </table>

```

```
t.test(gender ~ treatment_assignment, data = run2)
```

```

##
## Welch Two Sample t-test
##
## data: gender by treatment_assignment
## t = 0.48861, df = 286.52, p-value = 0.6255
## alternative hypothesis: true difference in means is not equal to 0
## 95 percent confidence interval:

```

```
## -0.09520922  0.15808801
## sample estimates:
## mean in group 0 mean in group 1
##      1.643939      1.612500
```

```
t.test(age ~ treatment_assignment, data = run2)
```

```
##
## Welch Two Sample t-test
##
## data: age by treatment_assignment
## t = 1.3895, df = 270.08, p-value = 0.1658
## alternative hypothesis: true difference in means is not equal to 0
## 95 percent confidence interval:
## -0.1019390  0.5909542
## sample estimates:
## mean in group 0 mean in group 1
##      3.575758      3.331250
```

```
t.test(age_under_40 ~ treatment_assignment, data = run2)
```

```
##
## Welch Two Sample t-test
##
## data: age_under_40 by treatment_assignment
## t = -1.5912, df = 275.56, p-value = 0.1127
## alternative hypothesis: true difference in means is not equal to 0
## 95 percent confidence interval:
## -0.20591972  0.02182881
## sample estimates:
## mean in group 0 mean in group 1
##      0.5454545      0.6375000
```

```
t.test(political_party ~ treatment_assignment, data = run2)
```

```
##
## Welch Two Sample t-test
##
## data: political_party by treatment_assignment
## t = -1.8651, df = 281.75, p-value = 0.06321
## alternative hypothesis: true difference in means is not equal to 0
## 95 percent confidence interval:
## -0.346847087  0.009347087
## sample estimates:
## mean in group 0 mean in group 1
##      2.00000      2.16875
```

```
t.test(ethnicity ~ treatment_assignment, data = run2)
```

```
##
## Welch Two Sample t-test
```

```
##
## data:  ethnicity by treatment_assignment
## t = 0.53453, df = 285.48, p-value = 0.5934
## alternative hypothesis: true difference in means is not equal to 0
## 95 percent confidence interval:
##  -0.1442760  0.2518517
## sample estimates:
## mean in group 0 mean in group 1
##      2.878788      2.825000
```

```
t.test(state ~ treatment_assignment, data = run2)
```

```
##
## Welch Two Sample t-test
##
## data:  state by treatment_assignment
## t = 0.91773, df = 274.56, p-value = 0.3596
## alternative hypothesis: true difference in means is not equal to 0
## 95 percent confidence interval:
##  -1.992233  5.471778
## sample estimates:
## mean in group 0 mean in group 1
##      25.47727      23.73750
```

```
t.test(college_educated ~ treatment_assignment, data = run2)
```

```
##
## Welch Two Sample t-test
##
## data:  college_educated by treatment_assignment
## t = -0.24298, df = 278.83, p-value = 0.8082
## alternative hypothesis: true difference in means is not equal to 0
## 95 percent confidence interval:
##  -0.12756046  0.09953016
## sample estimates:
## mean in group 0 mean in group 1
##      0.5984848      0.6125000
```

```
t.test(covid_sick ~ treatment_assignment, data = run2)
```

```
##
## Welch Two Sample t-test
##
## data:  covid_sick by treatment_assignment
## t = 0.33496, df = 279.15, p-value = 0.7379
## alternative hypothesis: true difference in means is not equal to 0
## 95 percent confidence interval:
##  -0.09605997  0.13545391
## sample estimates:
## mean in group 0 mean in group 1
##      1.469697      1.450000
```

```
t.test(covid_hospitalized ~ treatment_assignment, data = run2)
```

```
##
## Welch Two Sample t-test
##
## data: covid_hospitalized by treatment_assignment
## t = 0.51646, df = 283.29, p-value = 0.6059
## alternative hypothesis: true difference in means is not equal to 0
## 95 percent confidence interval:
## -0.07134676 0.12210434
## sample estimates:
## mean in group 0 mean in group 1
## 1.787879 1.762500
```

```
t.test(covid_died ~ treatment_assignment, data = run2)
```

```
##
## Welch Two Sample t-test
##
## data: covid_died by treatment_assignment
## t = 0.37052, df = 283.8, p-value = 0.7113
## alternative hypothesis: true difference in means is not equal to 0
## 95 percent confidence interval:
## -0.06452216 0.09444640
## sample estimates:
## mean in group 0 mean in group 1
## 1.871212 1.856250
```

```
t.test(party_republican ~ treatment_assignment, data = run2)
```

```
##
## Welch Two Sample t-test
##
## data: party_republican by treatment_assignment
## t = 1.4367, df = 265.7, p-value = 0.152
## alternative hypothesis: true difference in means is not equal to 0
## 95 percent confidence interval:
## -0.02644936 0.16925239
## sample estimates:
## mean in group 0 mean in group 1
## 0.2651515 0.1937500
```

```
t.test(party_democrat ~ treatment_assignment, data = run2)
```

```
##
## Welch Two Sample t-test
##
## data: party_democrat by treatment_assignment
## t = 0.18947, df = 279.42, p-value = 0.8499
## alternative hypothesis: true difference in means is not equal to 0
## 95 percent confidence interval:
```

```
## -0.1049179 0.1272664
## sample estimates:
## mean in group 0 mean in group 1
##      0.4924242      0.4812500
```

```
t.test(party_other ~ treatment_assignment, data = run2)
```

```
##
## Welch Two Sample t-test
##
## data: party_other by treatment_assignment
## t = -1.5657, df = 286.84, p-value = 0.1185
## alternative hypothesis: true difference in means is not equal to 0
## 95 percent confidence interval:
## -0.18638513 0.02123361
## sample estimates:
## mean in group 0 mean in group 1
##      0.2424242      0.3250000
```