

Experiments for Trust

Group

```
# Load the experiment's data
```

```
dmt <- read.csv("./cleaneddata/mturk.csv")
```

```
head(dmt)
```

```
##           Q6                               Q7    FL_3_D0
## 1 Ketogenic                               MControl
## 2 Pritikin I do not trust what the speaker was saying. FTreatment
## 3 Pritikin          I trust what the speaker was saying. FControl
## 4 Pritikin I do not trust what the speaker was saying. FTreatment
## 5 Pritikin          I trust what the speaker was saying. FControl
## 6   Paleo                               MTreatment
```

```
summary(dmt)
```

```
##           Q6                               Q7
##           : 4                               :276
## Ketogenic:144   I do not trust what the speaker was saying.: 31
## Paleo      :127   I trust what the speaker was saying.      :123
## Pritikin :155
##           FL_3_D0
## FControl  :109
## FTreatment:106
## MControl  :110
## MTreatment:105
```

```
# Adding columns for control, treatment and clusters (M, F)
```

```
t = c("FTreatment", "MTreatment")
```

```
s = c("FTreatment", "FControl")
```

```
dmt$treat = ifelse(dmt$FL_3_D0 %in% t, 1, 0 )
```

```
dmt$female = ifelse(dmt$FL_3_D0 %in% s, 1, 0)
```

```
dmt$trust = ifelse(dmt$Q7 == "I trust what the speaker was saying.", 1,
                   ifelse (dmt$Q6 != "Pritikin" | dmt$Q7 == "", NA, 0))
```

```
head(dmt, 100)
```

```
##           Q6                               Q7    FL_3_D0 treat
## 1 Ketogenic                               MControl      0
## 2 Pritikin I do not trust what the speaker was saying. FTreatment  1
## 3 Pritikin          I trust what the speaker was saying. FControl  0
## 4 Pritikin I do not trust what the speaker was saying. FTreatment  1
## 5 Pritikin          I trust what the speaker was saying. FControl  0
## 6   Paleo                               MTreatment  1
## 7 Ketogenic                               MControl      0
## 8 Pritikin I do not trust what the speaker was saying. FTreatment  1
## 9 Pritikin          I trust what the speaker was saying. MControl  0
## 10 Pritikin          I trust what the speaker was saying. FControl  0
## 11 Pritikin I do not trust what the speaker was saying. MTreatment  1
## 12 Pritikin I do not trust what the speaker was saying. MTreatment  1
## 13 Ketogenic                               FControl      0
## 14 Ketogenic                               MControl      0
## 15   Paleo                               FTreatment  1
```

## 16	Paleo	MTreatment	1
## 17	Ketogenic	FTreatment	1
## 18	Ketogenic	FControl	0
## 19	Paleo	FControl	0
## 20	Pritikin	I trust what the speaker was saying. FTreatment	1
## 21	Paleo	MTreatment	1
## 22	Pritikin	I do not trust what the speaker was saying. MControl	0
## 23	Ketogenic	MTreatment	1
## 24	Paleo	FTreatment	1
## 25	Paleo	FControl	0
## 26	Pritikin	I trust what the speaker was saying. FTreatment	1
## 27	Pritikin	I do not trust what the speaker was saying. MTreatment	1
## 28	Ketogenic	MControl	0
## 29	Pritikin	I trust what the speaker was saying. MTreatment	1
## 30	Pritikin	I trust what the speaker was saying. FTreatment	1
## 31	Pritikin	I trust what the speaker was saying. MControl	0
## 32	Paleo	FControl	0
## 33	Pritikin	I trust what the speaker was saying. MControl	0
## 34	Pritikin	I trust what the speaker was saying. FControl	0
## 35	Paleo	FTreatment	1
## 36		MTreatment	1
## 37		MControl	0
## 38		MControl	0
## 39		FControl	0
## 40	Ketogenic	MTreatment	1
## 41	Paleo	MControl	0
## 42	Pritikin	I trust what the speaker was saying. FTreatment	1
## 43	Paleo	MControl	0
## 44	Paleo	FTreatment	1
## 45	Ketogenic	FControl	0
## 46	Paleo	FControl	0
## 47	Ketogenic	MTreatment	1
## 48	Ketogenic	MTreatment	1
## 49	Paleo	FTreatment	1
## 50	Paleo	FTreatment	1
## 51	Paleo	MControl	0
## 52	Ketogenic	FControl	0
## 53	Paleo	MControl	0
## 54	Pritikin	I trust what the speaker was saying. MTreatment	1
## 55	Paleo	MControl	0
## 56	Pritikin	FTreatment	1
## 57	Ketogenic	FTreatment	1
## 58	Ketogenic	MTreatment	1
## 59	Pritikin	I trust what the speaker was saying. FControl	0
## 60	Ketogenic	FControl	0
## 61	Paleo	MTreatment	1
## 62	Paleo	FTreatment	1
## 63	Paleo	FControl	0
## 64	Ketogenic	MControl	0
## 65	Ketogenic	MControl	0
## 66	Ketogenic	FTreatment	1
## 67	Paleo	MTreatment	1
## 68	Ketogenic	MTreatment	1
## 69	Paleo	FControl	0

## 70	Paleo	FControl	0
## 71	Paleo	FTreatment	1
## 72	Paleo	MControl	0
## 73	Pritikin I do not trust what the speaker was saying.	FTreatment	1
## 74	Pritikin I trust what the speaker was saying.	FControl	0
## 75	Paleo	MTreatment	1
## 76	Pritikin I trust what the speaker was saying.	MTreatment	1
## 77	Ketogenic	FTreatment	1
## 78	Pritikin I trust what the speaker was saying.	MControl	0
## 79	Paleo	FControl	0
## 80	Ketogenic	FControl	0
## 81	Ketogenic	FTreatment	1
## 82	Paleo	MControl	0
## 83	Paleo	MTreatment	1
## 84	Paleo	FTreatment	1
## 85	Pritikin I do not trust what the speaker was saying.	MControl	0
## 86	Ketogenic	FControl	0
## 87	Pritikin I trust what the speaker was saying.	MTreatment	1
## 88	Paleo	FControl	0
## 89	Ketogenic	MControl	0
## 90	Pritikin I trust what the speaker was saying.	FControl	0
## 91	Ketogenic	FTreatment	1
## 92	Paleo	FTreatment	1
## 93	Pritikin I trust what the speaker was saying.	FControl	0
## 94	Ketogenic	MControl	0
## 95	Paleo	FTreatment	1
## 96	Ketogenic	MTreatment	1
## 97	Paleo	MControl	0
## 98	Pritikin I trust what the speaker was saying.	FTreatment	1
## 99	Pritikin I trust what the speaker was saying.	FControl	0
## 100	Paleo	MTreatment	1
##	female trust		
## 1	0 NA		
## 2	1 0		
## 3	1 1		
## 4	1 0		
## 5	1 1		
## 6	0 NA		
## 7	0 NA		
## 8	1 0		
## 9	0 1		
## 10	1 1		
## 11	0 0		
## 12	0 0		
## 13	1 NA		
## 14	0 NA		
## 15	1 NA		
## 16	0 NA		
## 17	1 NA		
## 18	1 NA		
## 19	1 NA		
## 20	1 1		
## 21	0 NA		
## 22	0 0		

## 23	0	NA
## 24	1	NA
## 25	1	NA
## 26	1	1
## 27	0	0
## 28	0	NA
## 29	0	1
## 30	1	1
## 31	0	1
## 32	1	NA
## 33	0	1
## 34	1	1
## 35	1	NA
## 36	0	NA
## 37	0	NA
## 38	0	NA
## 39	1	NA
## 40	0	NA
## 41	0	NA
## 42	1	1
## 43	0	NA
## 44	1	NA
## 45	1	NA
## 46	1	NA
## 47	0	NA
## 48	0	NA
## 49	1	NA
## 50	1	NA
## 51	0	NA
## 52	1	NA
## 53	0	NA
## 54	0	1
## 55	0	NA
## 56	1	NA
## 57	1	NA
## 58	0	NA
## 59	1	1
## 60	1	NA
## 61	0	NA
## 62	1	NA
## 63	1	NA
## 64	0	NA
## 65	0	NA
## 66	1	NA
## 67	0	NA
## 68	0	NA
## 69	1	NA
## 70	1	NA
## 71	1	NA
## 72	0	NA
## 73	1	0
## 74	1	1
## 75	0	NA
## 76	0	1

```
## 77      1    NA
## 78      0     1
## 79      1    NA
## 80      1    NA
## 81      1    NA
## 82      0    NA
## 83      0    NA
## 84      1    NA
## 85      0     0
## 86      1    NA
## 87      0     1
## 88      1    NA
## 89      0    NA
## 90      1     1
## 91      1    NA
## 92      1    NA
## 93      1     1
## 94      0    NA
## 95      1    NA
## 96      0    NA
## 97      0    NA
## 98      1     1
## 99      1     1
## 100     0    NA
```

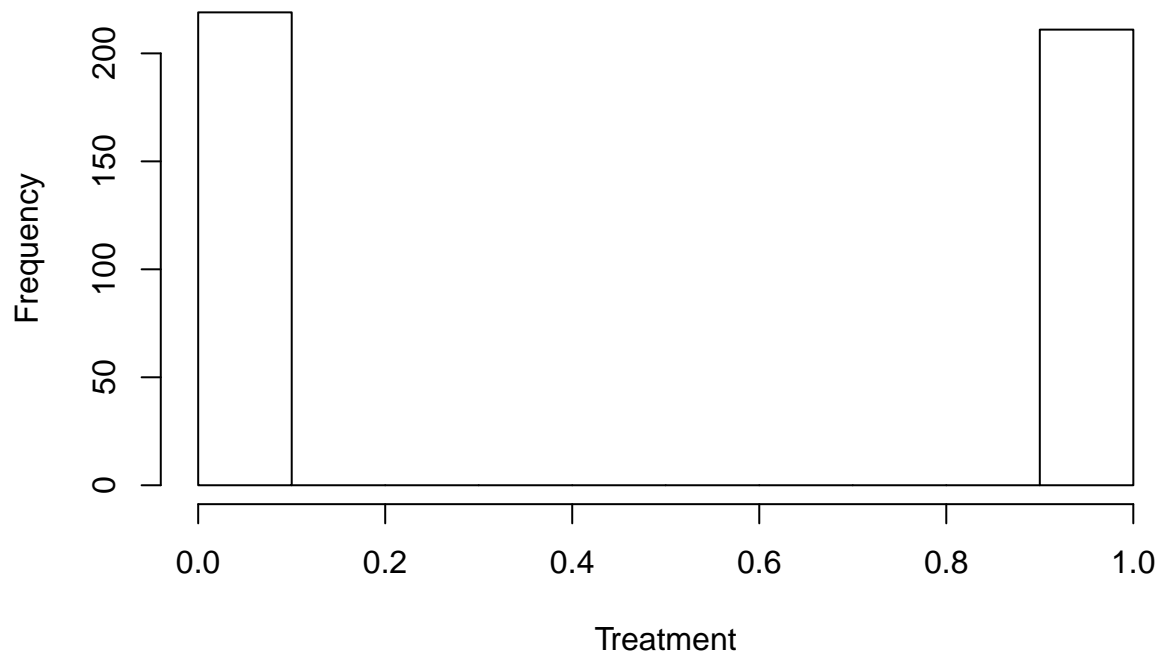
```
summary(dmt)
```

```
##           Q6                      Q7
##           : 4                      :276
## Ketogenic:144 I do not trust what the speaker was saying.: 31
## Paleo      :127 I trust what the speaker was saying.       :123
## Pritikin   :155
##
##
##
##      FL_3_D0      treat      female      trust
## FControl :109  Min.   :0.0000  Min.   :0.0   Min.   :0.0000
## FTreatment:106 1st Qu.:0.0000  1st Qu.:0.0   1st Qu.:1.0000
## MControl  :110 Median :0.0000  Median :0.5   Median :1.0000
## MTreatment:105 Mean   :0.4907  Mean   :0.5   Mean   :0.7987
##           3rd Qu.:1.0000  3rd Qu.:1.0   3rd Qu.:1.0000
##           Max.   :1.0000  Max.   :1.0   Max.   :1.0000
##
##                      NA's      :276
```

```
# Distribution of treatment vs control
```

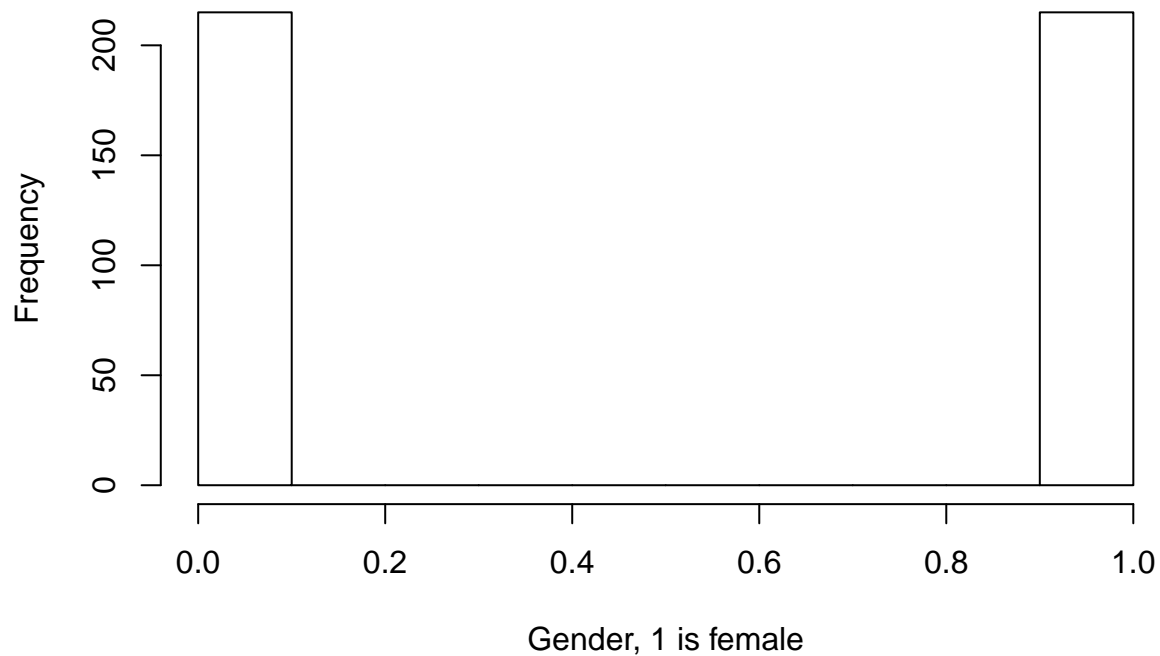
```
hist(dmt$treat, main='Distribution of treatment, 1 is treatment', xlab='Treatment')
```

Distribution of treatment, 1 is treatment



```
# Distribution of male vs female  
hist(dmt$female, main='Gender Distribution', xlab='Gender, 1 is female')
```

Gender Distribution



distributions are fine

These

```
# Regressing trust on treatment  
lrmt = lm(trust ~ treat, data = dmt)
```

```
summary(lrmt)
```

```
##
## Call:
## lm(formula = trust ~ treat, data = dmt)
##
## Residuals:
##      Min       1Q   Median       3Q      Max
## -0.97368  0.02632  0.02632  0.37179  0.37179
##
## Coefficients:
##              Estimate Std. Error t value Pr(>|t|)
## (Intercept)  0.97368    0.04178  23.305 < 2e-16 ***
## treat       -0.34548    0.05871  -5.885 2.45e-08 ***
## ---
## Signif. codes:  0 '***' 0.001 '**' 0.01 '*' 0.05 '.' 0.1 ' ' 1
##
## Residual standard error: 0.3642 on 152 degrees of freedom
## (276 observations deleted due to missingness)
## Multiple R-squared:  0.1856, Adjusted R-squared:  0.1802
## F-statistic: 34.63 on 1 and 152 DF,  p-value: 2.453e-08
```

```
ate = lrmt$coefficients['treat']
```

```
paste0('The ATE of the dissonance video is: ', ate)
```

```
## [1] "The ATE of the dissonance video is: -0.345479082321188"
```

```
# Adding Female as the covariate
```

```
lrmt1 = lm(trust ~ treat + female * treat, data = dmt)
```

```
summary(lrmt1)
```

```
##
## Call:
## lm(formula = trust ~ treat + female * treat, data = dmt)
##
## Residuals:
##      Min       1Q   Median       3Q      Max
## -0.93939  0.00000  0.06061  0.32500  0.42105
##
## Coefficients:
##              Estimate Std. Error t value Pr(>|t|)
## (Intercept)  0.93939    0.06343  14.809 < 2e-16 ***
## treat       -0.36045    0.08671  -4.157 5.4e-05 ***
## female       0.06061    0.08433   0.719  0.473
## treat:female  0.03545    0.11801   0.300  0.764
## ---
## Signif. codes:  0 '***' 0.001 '**' 0.01 '*' 0.05 '.' 0.1 ' ' 1
##
## Residual standard error: 0.3644 on 150 degrees of freedom
## (276 observations deleted due to missingness)
## Multiple R-squared:  0.1956, Adjusted R-squared:  0.1795
## F-statistic: 12.16 on 3 and 150 DF,  p-value: 3.63e-07
```

```
# Randomization inference
```

```
# Creating new variables for randomizing run and run_lag
```

```
dmt$treat_rnd = sample(dmt$treat)
```

```
head(dmt)
```

```
##           Q6                               Q7    FL_3_D0 treat
## 1 Ketogenic                               MControl    0
## 2 Pritikin I do not trust what the speaker was saying. FTreatment    1
## 3 Pritikin          I trust what the speaker was saying.  FControl    0
## 4 Pritikin I do not trust what the speaker was saying. FTreatment    1
## 5 Pritikin          I trust what the speaker was saying.  FControl    0
## 6 Paleo                               MTreatment    1
## female trust treat_rnd
## 1      0    NA      1
## 2      1      0      0
## 3      1      1      0
## 4      1      0      1
## 5      1      1      0
## 6      0    NA      0
```

```
summary(dmt)
```

```
##           Q6                               Q7
##           : 4                               :276
## Ketogenic:144 I do not trust what the speaker was saying.: 31
## Paleo      :127 I trust what the speaker was saying.      :123
## Pritikin   :155
##
##
##           FL_3_D0      treat      female      trust
## FControl :109   Min.   :0.0000   Min.   :0.0   Min.   :0.0000
## FTreatment:106   1st Qu.:0.0000   1st Qu.:0.0   1st Qu.:1.0000
## MControl  :110   Median :0.0000   Median :0.5   Median :1.0000
## MTreatment:105   Mean    :0.4907   Mean    :0.5   Mean    :0.7987
##           3rd Qu.:1.0000   3rd Qu.:1.0   3rd Qu.:1.0000
##           Max.    :1.0000   Max.    :1.0   Max.    :1.0000
##
##                               NA's      :276
## treat_rnd
## Min.      :0.0000
## 1st Qu.:0.0000
## Median :0.0000
## Mean    :0.4907
## 3rd Qu.:1.0000
## Max.    :1.0000
##
```

```
# Repeating the regression with the randomized treatment variable
# Regressing trust on treatment
```

```
lrmt2 = lm(trust ~ treat_rnd, data = dmt, na.action = na.omit)
summary(lrmt2)
```

```
##
## Call:
## lm(formula = trust ~ treat_rnd, data = dmt, na.action = na.omit)
##
## Residuals:
##      Min       1Q   Median       3Q      Max
```



```
## -0.8313  0.1687  0.1687  0.2394  0.2394
##
## Coefficients:
##           Estimate Std. Error t value Pr(>|t|)
## (Intercept)  0.83133    0.04413  18.838  <2e-16 ***
## treat_rnd   -0.07076    0.06499  -1.089    0.278
## ---
## Signif. codes:  0 '***' 0.001 '**' 0.01 '*' 0.05 '.' 0.1 ' ' 1
##
## Residual standard error: 0.402 on 152 degrees of freedom
## (276 observations deleted due to missingness)
## Multiple R-squared:  0.007739, Adjusted R-squared:  0.001211
## F-statistic: 1.185 on 1 and 152 DF, p-value: 0.278

names(summary(lrmt2))

## [1] "call"          "terms"          "residuals"      "coefficients"
## [5] "aliased"        "sigma"          "df"             "r.squared"
## [9] "adj.r.squared" "fstatistic"     "cov.unscaled"   "na.action"

summary(lrmt2)$coefficients

##           Estimate Std. Error  t value    Pr(>|t|)
## (Intercept)  0.83132530 0.04412912 18.838476 1.359259e-41
## treat_rnd   -0.07076192 0.06499148 -1.088788 2.779708e-01

summary(lrmt2)$coefficients[,2]['treat_rnd']

## treat_rnd
## 0.06499148

# Generic function to randomly(pseudo) pick the treatment assignments
coeff_treat_rnd <- function(d) {
  d$treat_rnd = sample(d$treat)
  lmr = lm(trust ~ treat_rnd, data=dmt, na.action = na.omit)
  return (lmr$coefficients['treat_rnd'])
}

# Trying another randomization
coeff_treat_rnd(dmt)

## treat_rnd
## -0.07076192

# 1K Randomizations
h.distribution.under.sharp.null <- replicate(10000, coeff_treat_rnd(dmt))
h.ate.mean <- mean(h.distribution.under.sharp.null)
paste0("Mean from 1k randomizations: ", h.ate.mean)

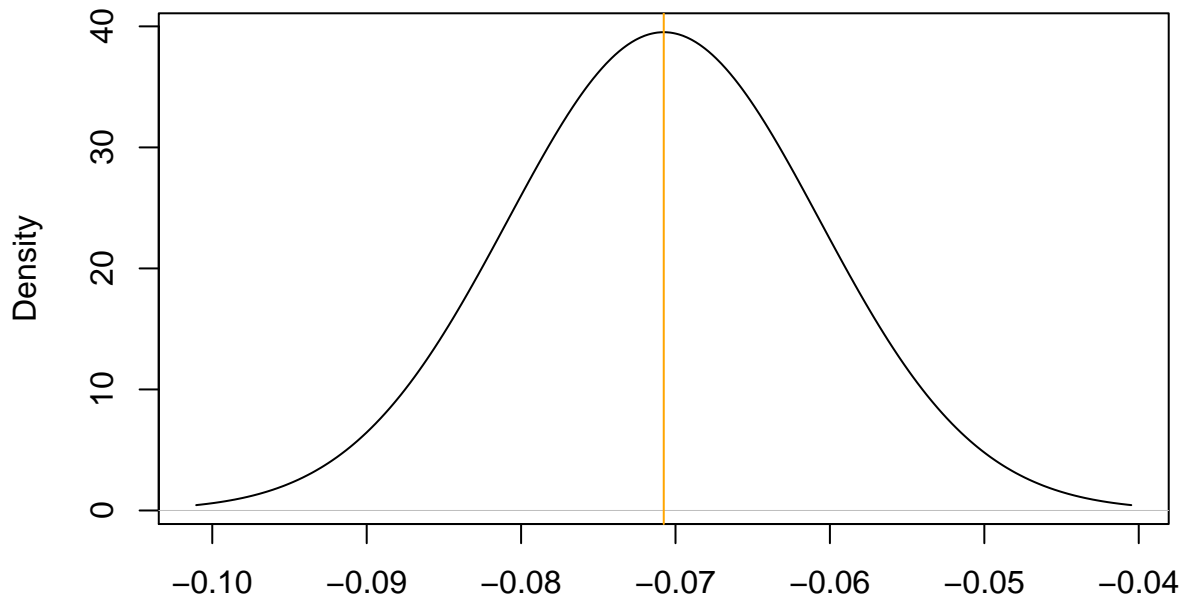
## [1] "Mean from 1k randomizations: -0.0707619209231295"

# Graph for the estimates
plot(density(h.distribution.under.sharp.null),
     main = "Density under Sharp Null")

# Adding our original ATE to the plot
abline(v = ate, col = "blue")
```

```
# Adding the mean to the plot
abline(v = mean(h.distribution.under.sharp.null), col = 'orange')
```

Density under Sharp Null



N = 10000 Bandwidth = 0.01009

```
# p value
pv = mean(h.distribution.under.sharp.null >= ate)
paste0('p value: ', pv)

## [1] "p value: 1"

# num of assignments that generate an estimated ATE at least as large as the actual
n <- sum(h.distribution.under.sharp.null >= ate)
paste0("Number of assignments that generate an ATE at least as large as what we got from the experiment")

## [1] "Number of assignments that generate an ATE at least as large as what we got from the experiment"

# Standard error
se = sd(h.distribution.under.sharp.null)/sqrt(length(h.distribution.under.sharp.null))
paste0('Standard error: ', se)

## [1] "Standard error: 0"
```