Data Analysis_HB_28042016

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Descriptive Statistics

Following are relevant aggregated statistics and statistics by each of the 11 treatments for each of three relevant dependent variables. These relevant dependent variables are 1. Donation, which is the amount the subject donated in order to retire emission rights 2. Donated, which is equal to 1 if the subject donated a positive amount, and 0 otherwise 3. Belief, which is the amount the subject thinks other participants in this experiment donated on average (not incentivized)

1. Variable: Donation to retire carbon licenses

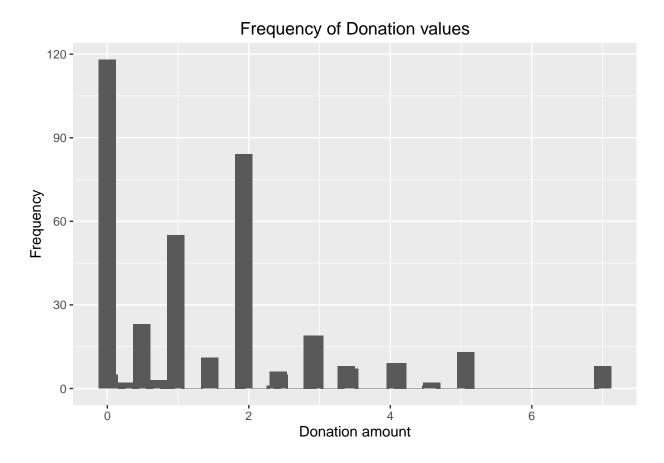
Aggregated descriptive statistics

```
summary(df$Donation)

## Min. 1st Qu. Median Mean 3rd Qu. Max.
## 0.00 0.00 1.00 1.44 2.00 7.00

sd(df$Donation)
```

Distribution of aggregated donations

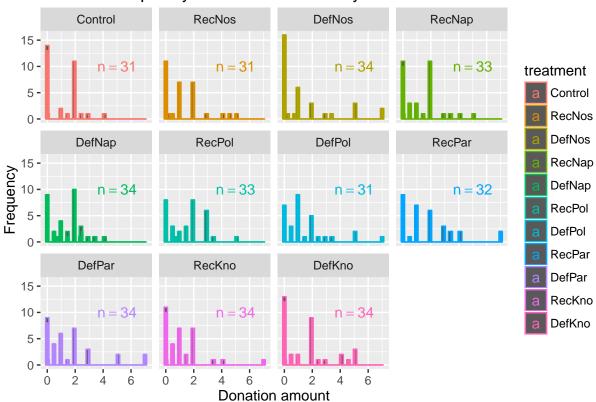


Distribution of donations by treatment

```
## group: Control
## vars n mean sd median trimmed mad min max range skew kurtosis
## -----
## group: RecNos
 vars n mean sd median trimmed mad min max range skew kurtosis
## -----
## group: DefNos
## vars n mean sd median trimmed mad min max range skew kurtosis
   1 34 1.42 2.08 0.5 1.05 0.74 0 7 7 1.46 0.93 0.36
## -----
## group: RecNap
## vars n mean sd median trimmed mad min max range skew kurtosis
## 1 1 33 1.32 1.31 1 1.15 1.48 0 5 5 0.83
## -----
## group: DefNap
## vars n mean sd median trimmed mad min max range skew kurtosis se
  1 34 1.36 1.11 1.45 1.28 1.19 0 4 4 0.3 -0.81 0.19
## group: RecPol
```

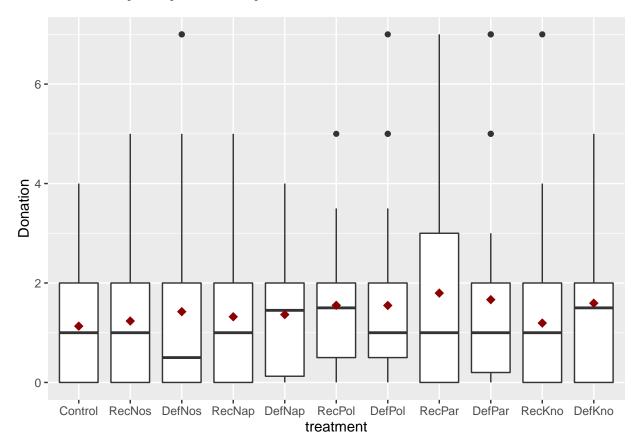
```
## vars n mean sd median trimmed mad min max range skew kurtosis se
## -----
## group: DefPol
## vars n mean sd median trimmed mad min max range skew kurtosis se
## 1 1 31 1.55 1.68 1 1.24 1.48 0 7 7 1.55
## -----
## group: RecPar
## vars n mean sd median trimmed mad min max range skew kurtosis se
## 1 1 32 1.8 1.88 1 1.52 1.48 0 7 7 1.23
                                    1.1 0.33
## group: DefPar
## vars n mean sd median trimmed mad min max range skew kurtosis se
## -----
## group: RecKno
## vars n mean sd median trimmed mad min max range skew kurtosis se
## group: DefKno
## vars n mean sd median trimmed mad min max range skew kurtosis
## 1 1 34 1.59 1.7 1.5 1.4 2.22 0 5 5 0.73 -0.74 0.29
```

Frequency of Donation values by treatment



Donations by treatment (Boxplot)

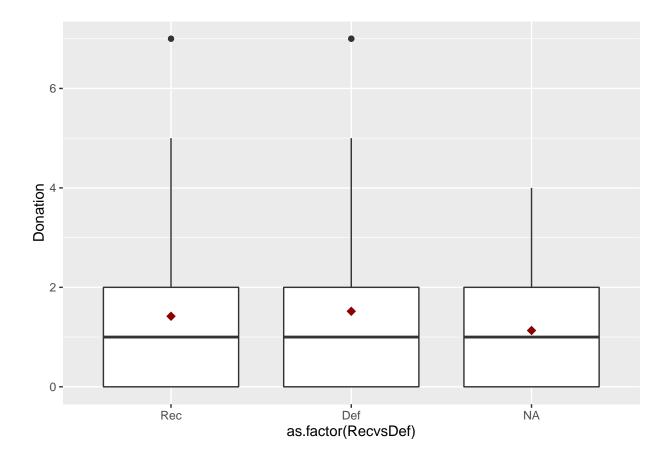
Red diamonds in boxplots represent the respective means



Donations by aggregated treatment (Boxplot), i.e. Def vs. Rec vs. Control

describeBy(df\$Donation, df\$RecvsDef)

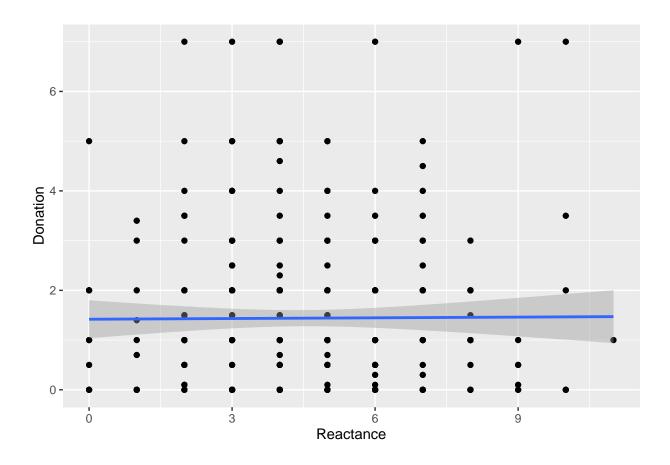
```
## group: Rec
##
    vars
                  sd median trimmed mad min max range skew kurtosis
          n mean
                               1.2 1.48 0
                                                  7 1.36
       1 163 1.42 1.48
                         1
## group: Def
    vars
          n mean sd median trimmed mad min max range skew kurtosis
    1 167 1.52 1.71 1 1.21 1.48
                                         0
                                             7
                                                   7 1.4
```

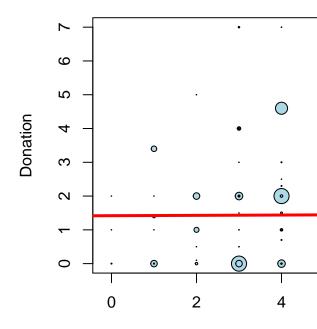


Donations by Reactance score

The reactance score was constructed by changing each of the 11 rectance-items to a dummy variable equal to 1 if the subject chose 3 or 4 on the respective item, and 0 otherwise. Afterwards, all 11 dummies were added to construct an ordinal Reactance score.

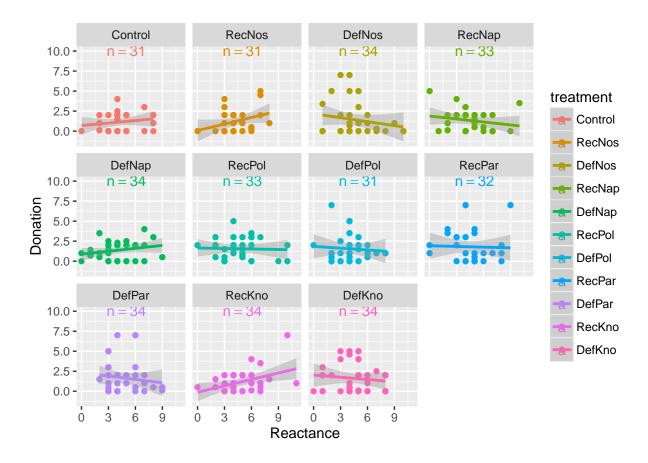
Shows a point plot (not jittered) with Donation amount and the respective Reactance score of each participant. Includes a linear regression line, including the 95% confidence region, of the Reactance score as a predictor for the Donation amount.





To see how often combinations of Reactance and Donation scores appeared $\,$

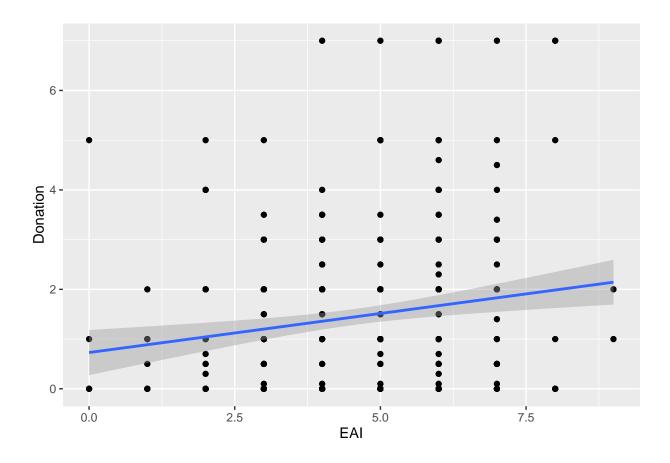
Donations by Reactance score per treatment Shows a point plot (not jittered) with Donation amount and the respective Reactance score of each participant, for each treatment. Includes a linear regression line, including the 95% confidence region, of the Reactance score as a predictor for the Donation amount, for each treatment.



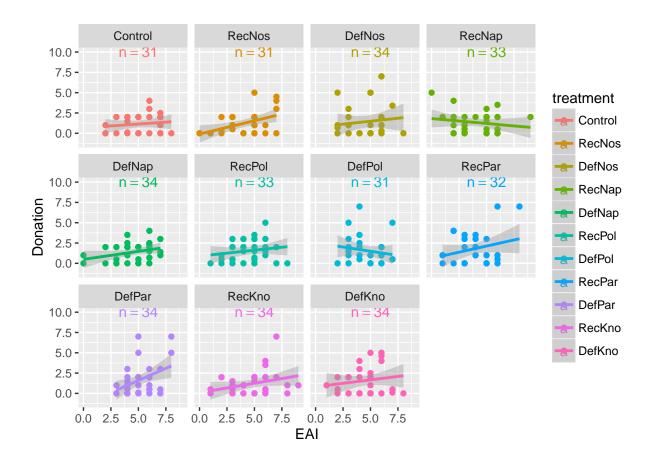
Donations by EAI score

The EAI score was constructed by changing each of the 12 EAI-items to a dummy variable equal to 1 if the subject chose 3 or 4 on the respective item, and 0 otherwise. Afterwards, all 12 dummies were added to construct an ordinal EAI score.

Shows a point plot (not jittered) with Donation amount and the respective EAI score of each participant. Includes a linear regression line, including the 95% confidence region, of the EAI score as a predictor for the Donation amount.



Donations by EAI score per treatment Shows a point plot (not jittered) with Donation amount and the respective EAI score of each participant, for each treatment. Includes a linear regression line, including the 95% confidence region, of the EAI score as a predictor for the Donation amount, for each treatment.



2. Variable: Donation dummy (1 if donated, 0 otherwise)

Aggregated descriptive statistics

```
summary(df$Donated)
```

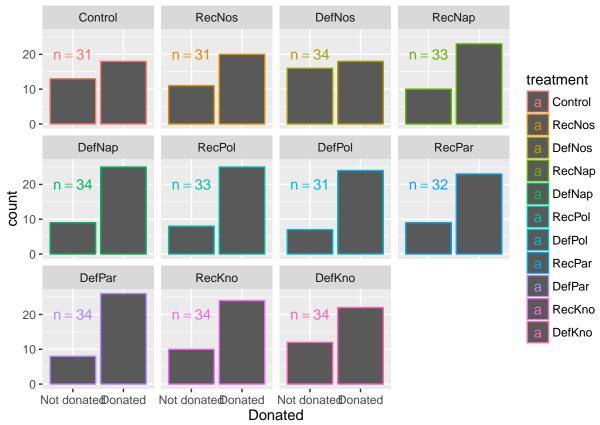
Not donated Donated ## 113 248

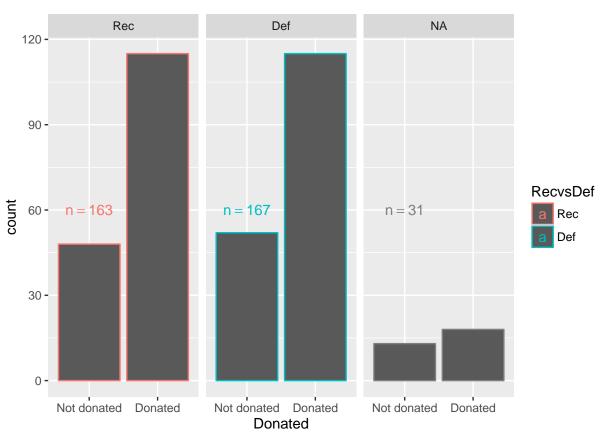
Distribution of donation dummy by treatment

table(df\$Donated, df\$treatment)

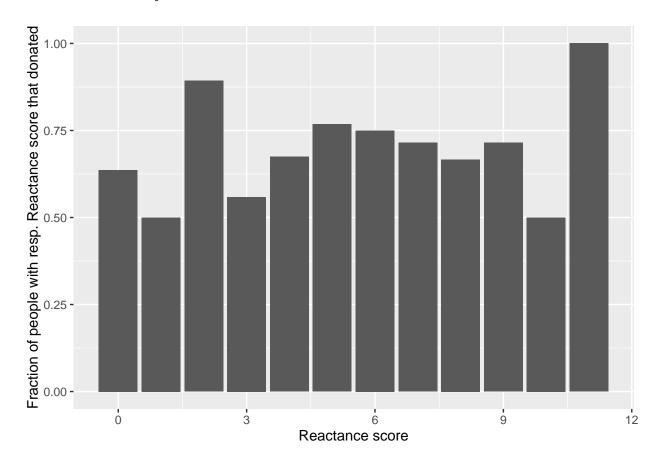
```
##
##
                  Control RecNos DefNos RecNap DefNap RecPol DefPol RecPar
     Not donated
                        13
                               11
                                       16
                                               10
                                                               8
                                                                      7
                                                                              9
##
##
     Donated
                        18
                               20
                                       18
                                               23
                                                      25
                                                              25
                                                                     24
                                                                             23
##
                  DefPar RecKno DefKno
##
##
     Not donated
                       8
                              10
                                      12
##
     Donated
                      26
                              24
                                      22
```

Decision to donate by treatment graph





Decision to donate by Reactance score

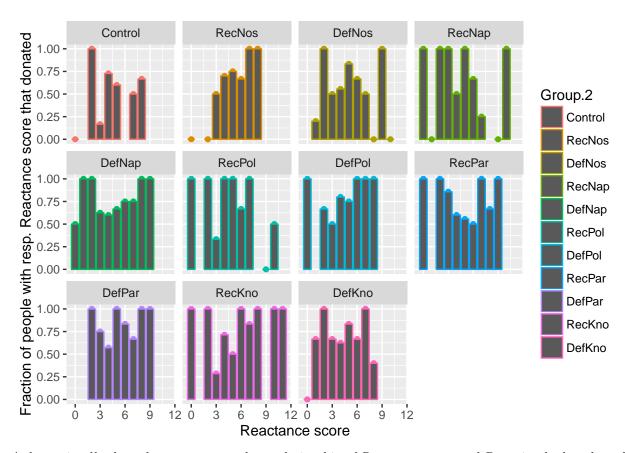


chisq.test(table(df\$Donated, df\$Reactance))

```
## Warning in chisq.test(table(df$Donated, df$Reactance)): Chi-squared
## approximation may be incorrect

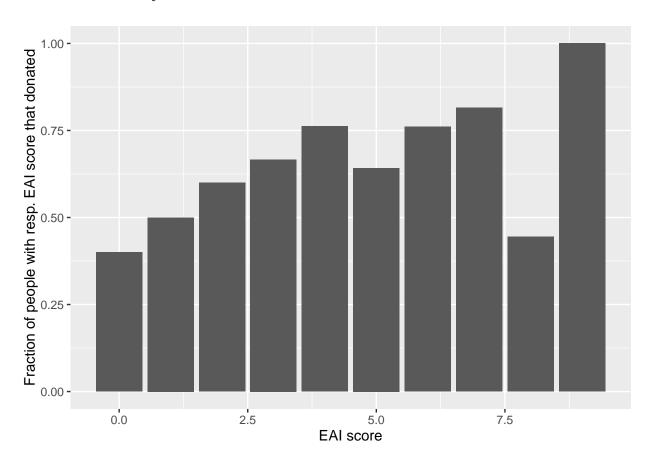
##
## Pearson's Chi-squared test
##
## data: table(df$Donated, df$Reactance)
## X-squared = 16.887, df = 11, p-value = 0.1113
```

Decision to donate by Reactance score and treatment



At least visually there does not seem to be a relationship of Reactance score and Donation broken down by treatment.

Decision to donate by EAI score



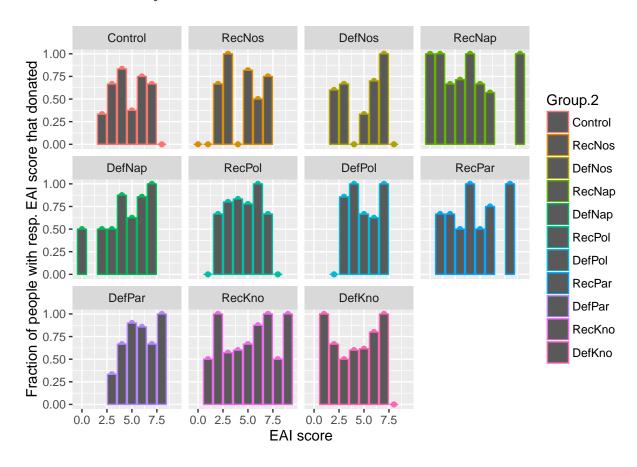
chisq.test(table(df\$Donated, df\$EAI))

```
## Warning in chisq.test(table(df$Donated, df$EAI)): Chi-squared approximation
## may be incorrect

##
## Pearson's Chi-squared test
##
## data: table(df$Donated, df$EAI)
## X-squared = 14.31, df = 9, p-value = 0.1117
```

The Chi^2 -test statistic is not significant, indicating that the decision whether or not to donate anything does not depend on the EAI.

Decision to donate by EAI score and treatment



3. Variable: Beliefs about other participants donations

1.974

Aggregated descriptive statistics

1.000

2.000

##

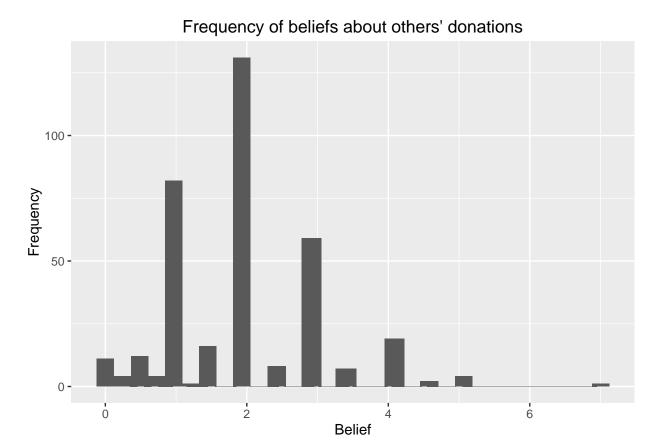
0.000

```
summary(df$belief)
## Min. 1st Qu. Median Mean 3rd Qu. Max.
```

7.000

3.000

Distribution of aggregated beliefs about donations

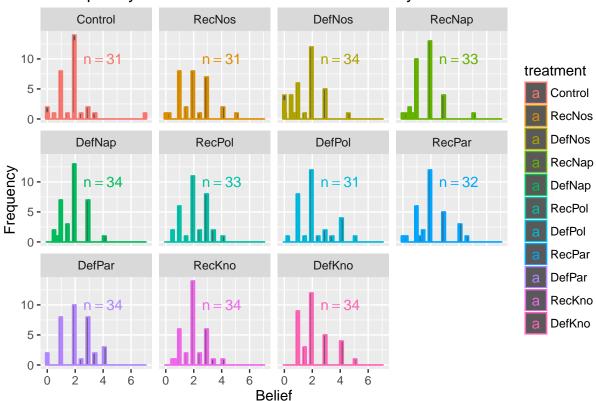


Distribution of beliefs by treatment

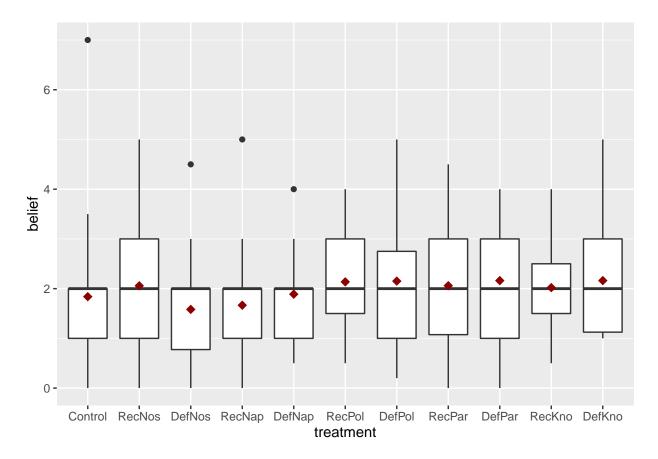
```
## group: Control
## vars n mean sd median trimmed mad min max range skew kurtosis
## 1 1 31 1.84 1.25 2 1.72 0.74 0 7 7 2.1
## -----
## group: RecNos
 vars n mean sd median trimmed mad min max range skew kurtosis
## -----
## group: DefNos
## vars n mean sd median trimmed mad min max range skew kurtosis
   1 34 1.58 1.07 2 1.55 1.48 0 4.5 4.5 0.43 -0.24 0.18
## -----
## group: RecNap
## vars n mean sd median trimmed mad min max range skew kurtosis
## 1 1 33 1.67 1.01 2 1.6 1.48 0 5 5 0.97 1.55 0.18
## group: DefNap
## vars n mean sd median trimmed mad min max range skew kurtosis se
   1 34 1.89 0.86 2 1.88 1.48 0.5 4 3.5 0.32 -0.63 0.15
## group: RecPol
```

```
## vars n mean sd median trimmed mad min max range skew kurtosis se
## -----
## group: DefPol
## vars n mean sd median trimmed mad min max range skew kurtosis se
## 1 1 31 2.15 1.16 2 2.06 1.48 0.2 5 4.8 0.67 -0.39 0.21
## -----
## group: RecPar
## vars n mean sd median trimmed mad min max range skew kurtosis
## 1 1 32 2.06 1.1 2 2 1.48 0 4.5 4.5 0.43 -0.46 0.19
## group: DefPar
## vars n mean sd median trimmed mad min max range skew kurtosis se
## 1 1 34 2.16 1.1 2 2.16 1.48 0 4 4 -0.1 -0.93 0.19
## group: RecKno
## vars n mean sd median trimmed mad min max range skew kurtosis se
## group: DefKno
## vars n mean sd median trimmed mad min max range skew kurtosis se
## 1 1 34 2.16 1.09 2 2.05 1.48 1 5 4 0.82 -0.24 0.19
```

Frequency of beliefs about others donations by treatment



Beliefs by treatment (Boxplot)



Inferential Statistics

Following are relevant inferential statistics for each of three relevant dependent variables. These relevant dependent variables are 1. Donation, which is the amount the subject donated in order to retire emission rights 2. Donated, which is equal to 1 if the subject donated a positive amount, and 0 otherwise 3. Belief, which is the amount the subject thinks other participants in this experiment donated on average (not incentivized)

1. Variable: Donation to retire carbon licenses

Kruskal-Wallis-Test

The following KW-test tests the null-hypothesis that the median donations in each treatment are the same. The test assumes variance homogeneity and equal distributions of donations in each treatment. It basically tests whether the distributions from the different treatments are shifted.

```
kruskal.test(df$Donation ~ df$treatment)
```

```
##
## Kruskal-Wallis rank sum test
##
## data: df$Donation by df$treatment
```

```
## Kruskal-Wallis chi-squared = 6.3484, df = 10, p-value = 0.7852   
We do not reject the null (p=.05).
```

ANOVA (one-way)

```
## Df Sum Sq Mean Sq F value Pr(>F)
## df$treatment 10 14.3 1.433 0.578 0.832
## Residuals 350 867.1 2.477

We do not reject the null (p = .05).
```

2. Variable: Donation dummy (1 if donated, 0 otherwise)

Chi² Test

The following Chi²-test tests the null-hypothesis that whether or not a participant decides to donate anything to retire emission rights (extensive margin) is independent of the treatments.

```
table(df$Donated, df$treatment)
```

```
##
##
                   Control RecNos DefNos RecNap DefNap RecPol DefPol RecPar
##
     Not donated
                        13
                                11
                                        16
                                                10
                                                                         7
##
     Donated
                        18
                                20
                                        18
                                                23
                                                        25
                                                                25
                                                                        24
                                                                                23
##
##
                   DefPar RecKno DefKno
##
     Not donated
                        8
                               10
                                       12
                       26
                                       22
##
     Donated
                               24
```

```
chisq.test(table(df$Donated, df$treatment))
```

```
##
## Pearson's Chi-squared test
##
## data: table(df$Donated, df$treatment)
## X-squared = 9.4673, df = 10, p-value = 0.4884
We fail to reject the null (p = .05)
```

3. Variable: Beliefs about other participants donations

Kruskal-Wallis Test

The following KW-test tests the null-hypothesis that the median beliefs about other participants average donations in each treatment are the same. The test assumes variance homogeneity and equal distributions of donations in each treatment. It basically tests whether the distributions from the different treatments are shifted.

kruskal.test(df\$belief ~ df\$treatment)

```
##
## Kruskal-Wallis rank sum test
##
## data: df$belief by df$treatment
## Kruskal-Wallis chi-squared = 14.092, df = 10, p-value = 0.1688
```

We reject the null (p = .05) in favor of the alternative hypothesis that in at least two treatments do the beliefs about other participants donation amounts differ significantly.

Test of hypotheses from the working paper

H₀a

Mean and median payments to retire carbon licenses in the control condition are close to zero.

H0: Average Donations = 0 HA: Average Donations >< 0

```
t.test(df$Donation, mu = 0)
```

```
##
## One Sample t-test
##
## data: df$Donation
## t = 17.487, df = 360, p-value < 2.2e-16
## alternative hypothesis: true mean is not equal to 0
## 95 percent confidence interval:
## 1.278209 1.602123
## sample estimates:
## mean of x
## 1.440166</pre>
```

```
wilcox.test(df$Donation, mu = 0)
```

```
##
## Wilcoxon signed rank test with continuity correction
##
## data: df$Donation
## V = 30876, p-value < 2.2e-16
## alternative hypothesis: true location is not equal to 0</pre>
```

We reject the null that Donations are equal to 0

H0b

The share of subjects whose payments correspond to the recommended, respectively defaulted payment-value (convergence) is higher than in the control condition. Additionally, we expect that the share of subjects converging to the default is higher than the share converging to the recommendation.

Aggregated donations in recommendation treatments > donations in control group

```
describeBy(df$Donation, df$RecvsC)
## group: Control
## vars n mean sd median trimmed mad min max range skew kurtosis
      1 31 1.13 1.15 1
                           1.02 1.48 0 4
                                             4 0.42
                                                      -0.940.21
## -----
## group: Rec
## vars n mean sd median trimmed mad min max range skew kurtosis se
      ## 1
                                                        2.27 0.12
t.test(df$Donation ~ df$RecvsC)
##
## Welch Two Sample t-test
##
## data: df$Donation by df$RecvsC
## t = -1.2075, df = 50.813, p-value = 0.2328
## alternative hypothesis: true difference in means is not equal to 0
## 95 percent confidence interval:
## -0.7619248 0.1896311
## sample estimates:
## mean in group Control mean in group Rec
                               1.418405
##
             1.132258
wilcox.test(df$Donation ~ df$RecvsC)
##
## Wilcoxon rank sum test with continuity correction
## data: df$Donation by df$RecvsC
## W = 2307.5, p-value = 0.4336
\#\# alternative hypothesis: true location shift is not equal to 0
We cannot reject the null that Donations in recommendation treatments are equal to donations in control
condition.
Aggregated donations in default treatments > donations in control group
describeBy(df$Donation, df$DefvsC)
## group: Control
   vars n mean sd median trimmed mad min max range skew kurtosis
     1 31 1.13 1.15 1
                           1.02 1.48 0 4
                                            4 0.42 -0.94 0.21
## -----
## group: Def
   vars n mean sd median trimmed mad min max range skew kurtosis
      1.63 0.13
```

```
t.test(df$Donation ~ df$DefvsC)
##
## Welch Two Sample t-test
## data: df$Donation by df$DefvsC
## t = -1.5742, df = 57.756, p-value = 0.1209
\#\# alternative hypothesis: true difference in means is not equal to 0
## 95 percent confidence interval:
## -0.8775591 0.1049495
## sample estimates:
## mean in group Control mean in group Def
              1.132258
                                   1.518563
wilcox.test(df$Donation ~ df$DefvsC)
##
## Wilcoxon rank sum test with continuity correction
## data: df$Donation by df$DefvsC
## W = 2354.5, p-value = 0.4134
\#\# alternative hypothesis: true location shift is not equal to 0
We cannot reject the null that Donations in default treatments are equal to donations in control condition.
Aggregated donations in default treatments > donations in recommendation treatments
describeBy(df$Donation, df$RecvsDef)
## group: Rec
## vars n mean sd median trimmed mad min max range skew kurtosis
       2.27 0.12
## group: Def
## vars n mean sd median trimmed mad min max range skew kurtosis
     1 167 1.52 1.71 1
                             1.21 1.48 0 7 7 1.4
t.test(df$Donation ~ df$RecvsDef)
##
## Welch Two Sample t-test
## data: df$Donation by df$RecvsDef
## t = -0.57023, df = 323.38, p-value = 0.5689
## alternative hypothesis: true difference in means is not equal to 0
## 95 percent confidence interval:
## -0.4457065 0.2453905
## sample estimates:
## mean in group Rec mean in group Def
          1.418405
                           1.518563
##
```

```
wilcox.test(df$Donation ~ df$RecvsDef)
```

```
##
## Wilcoxon rank sum test with continuity correction
##
## data: df$Donation by df$RecvsDef
## W = 13584, p-value = 0.9755
## alternative hypothesis: true location shift is not equal to 0
```

We cannot reject the null that Donations in default treatments are equal to donations in recommendation treatments.

H0c

The share of subjects converging to the recommended, respectively defaulted payment-values in the name and picture condition is higher than in the neutral source-condition.

For Recommendations: Donations in Name and Picture treatments > Donations in No-Source treatments

```
describeBy(df$Donation, df$RecNapvsRecNos)
## group: RecNap
   vars n mean sd median trimmed mad min max range skew kurtosis
       1 33 1.32 1.31
                                          0 5
                                                   5 0.83
                         1
                               1.15 1.48
                                                              0.13 0.23
## group: RecNos
    vars n mean sd median trimmed mad min max range skew kurtosis
## 1
       1 31 1.24 1.39
                          1
                               0.99 1.48 0
                                              5
                                                   5 1.14
                                                              0.52 0.25
```

```
##
## Welch Two Sample t-test
##
## data: df$Donation by df$RecNapvsRecNos
## t = 0.25345, df = 61.007, p-value = 0.8008
## alternative hypothesis: true difference in means is not equal to 0
## 95 percent confidence interval:
## -0.5906449 0.7621014
## sample estimates:
## mean in group RecNap mean in group RecNos
## 1.321212 1.235484

wilcox.test(df$Donation ~ df$RecNapvsRecNos)
```

Warning in wilcox.test.default(x = c(2, 0.5, 0, 2, 1, 2, 3.5, 2, 3, 0,

0.5, : cannot compute exact p-value with ties

```
##
## Wilcoxon rank sum test with continuity correction
##
## data: df$Donation by df$RecNapvsRecNos
## W = 546, p-value = 0.6373
## alternative hypothesis: true location shift is not equal to 0
```

We cannot reject the null that Donations in recommendation treatments informing about the name and picture of the source are equal to donations in recommendation treatments providing no information about the source of the recommendation.

For Defaults: Donations in Name and Picture treatments > Donations in No-Source treatments

```
describeBy(df$Donation, df$DefNapvsDefNos)
## group: DefNap
                    sd median trimmed mad min max range skew kurtosis
     vars n mean
                                 1.28 1.19
                                                       4 0.3
        1 34 1.36 1.11
                         1.45
                                             0
                                                                 -0.810.19
## group: DefNos
     vars n mean sd median trimmed mad min max range skew kurtosis
        1 34 1.42 2.08
                          0.5
                                 1.05 0.74
                                             0
                                                       7 1.46
                                                                  0.93 0.36
## 1
t.test(df$Donation ~ df$DefNapvsDefNos)
##
   Welch Two Sample t-test
##
## data: df$Donation by df$DefNapvsDefNos
## t = -0.14519, df = 50.431, p-value = 0.8851
## alternative hypothesis: true difference in means is not equal to 0
## 95 percent confidence interval:
## -0.872389 0.754742
## sample estimates:
## mean in group DefNap mean in group DefNos
##
               1.364706
                                    1.423529
wilcox.test(df$Donation ~ df$DefNapvsDefNos)
## Warning in wilcox.test.default(x = c(2, 1, 0.7, 2, 3.5, 1, 0.5, 2, 0, 3, :
## cannot compute exact p-value with ties
##
##
   Wilcoxon rank sum test with continuity correction
##
## data: df$Donation by df$DefNapvsDefNos
## W = 677, p-value = 0.2127
## alternative hypothesis: true location shift is not equal to 0
```

We cannot reject the null that Donations in default treatments informing about the name and picture of the source are equal to donations in default treatments providing no information about the source of the default.

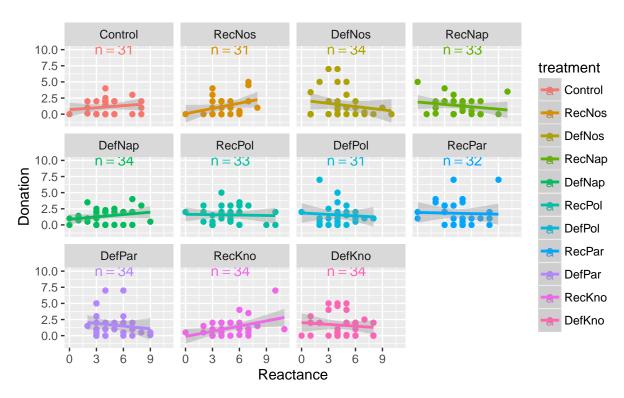
H1

A subject's reaction towards the respective intervention depends on is predicted by trait reactance.

The following are not rigurous tests of the respective hypotheses, but rather approaches to get an idea about relationships and predictions.

Relation between Donation and Reactance score

conditional on treatment

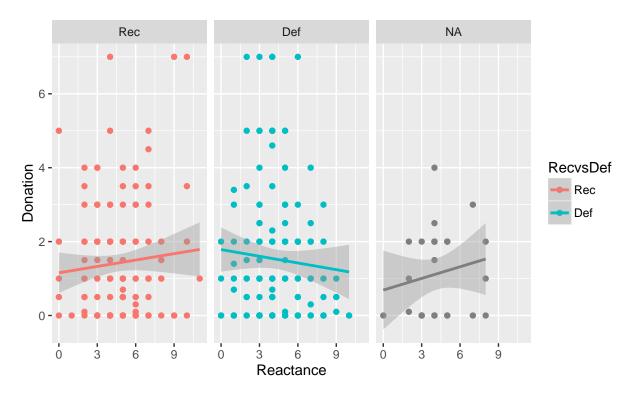


H1a

A subject that scores high on trait reactance is less likely to converge to the recommended and defaulted payment-values, than a subject scoring low on trait reactance. The following treats the Reactance score as metric.

Relation between Donation and Reactance score

resp. for Rec and Def treatment groups

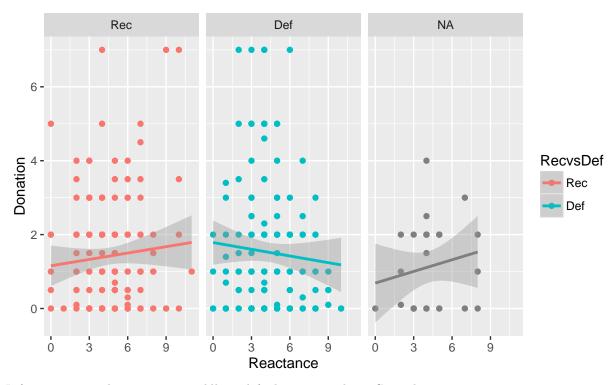


H1b

A subject that scores high on trait reactance is less likely to converge to the defaulted than to the recommended payment-value.

Relation between Donation and Reactance score

resp. for Rec and Def treatment groups



Left is recommendation group, middle is default group, right is Control.

H2

The share of subjects converging to the recommended, respectively defaulted payment-values in the condition informing about the academic degree of the source is higher than in the name and picture condition.

For Recommendations: Donations in Knowledge treatments > Donations in Name and Picture treatments

```
describeBy(df$Donation, df$RecNapvsRecKno)

## group: RecNap

## vars n mean sd median trimmed mad min max range skew kurtosis se
```

```
5 0.83
                                              0
                                                                    0.13 0.23
        1 33 1.32 1.31
                             1
                                  1.15 1.48
  group: RecKno
                    sd median trimmed mad min max range skew kurtosis
          n mean
     vars
        1 34 1.19 1.45
                             1
                                  0.93 1.48
                                              0
                                                         7 2.07
                                                                    5.36 0.25
## 1
```

t.test(df\$Donation ~ df\$RecNapvsRecKno)

##

```
## Welch Two Sample t-test
##
## data: df$Donation by df$RecNapvsRecKno
## t = 0.37725, df = 64.659, p-value = 0.7072
## alternative hypothesis: true difference in means is not equal to 0
## 95 percent confidence interval:
## -0.5458115 0.8000005
## sample estimates:
## mean in group RecNap mean in group RecKno
             1.321212
                               1.194118
wilcox.test(df$Donation ~ df$RecNapvsRecKno)
## Warning in wilcox.test.default(x = c(2, 0.5, 0, 2, 1, 2, 3.5, 2, 3, 0,
## 0.5, : cannot compute exact p-value with ties
##
## Wilcoxon rank sum test with continuity correction
## data: df$Donation by df$RecNapvsRecKno
## W = 608.5, p-value = 0.5453
## alternative hypothesis: true location shift is not equal to 0
For Defaults: Donations in Knowledge treatments > Donations in Name and Picture treat-
ments
describeBy(df$Donation, df$DefNapvsDefKno)
## group: DefNap
## vars n mean sd median trimmed mad min max range skew kurtosis
## group: DefKno
## vars n mean sd median trimmed mad min max range skew kurtosis
t.test(df$Donation ~ df$DefNapvsDefKno)
##
## Welch Two Sample t-test
## data: df$Donation by df$DefNapvsDefKno
## t = -0.65788, df = 56.889, p-value = 0.5133
## alternative hypothesis: true difference in means is not equal to 0
## 95 percent confidence interval:
## -0.9277281 0.4689046
## sample estimates:
## mean in group DefNap mean in group DefKno
##
             1.364706
                                1.594118
```

```
wilcox.test(df$Donation ~ df$DefNapvsDefKno)

## Warning in wilcox.test.default(x = c(2, 1, 0.7, 2, 3.5, 1, 0.5, 2, 0, 3, :
## cannot compute exact p-value with ties

##
## Wilcoxon rank sum test with continuity correction
##
## data: df$Donation by df$DefNapvsDefKno
## W = 573.5, p-value = 0.9598
## alternative hypothesis: true location shift is not equal to 0
```

H3-1

The share of subjects converging to the recommended, respectively defaulted payment-values in the condition informing about the political characteristic of the source is lower than in the name and picture condition.

For Recommendations: Donations in Political treatments < Donations in Name and Picture treatments

```
describeBy(df$Donation, df$RecNapvsRecPol)
## group: RecNap
                    sd median trimmed mad min max range skew kurtosis
    vars n mean
                                                       5 0.83
        1 33 1.32 1.31
                           1
                                 1.15 1.48
                                             0
                                                                  0.13 0.23
## group: RecPol
     vars n mean sd median trimmed mad min max range skew kurtosis
        1 33 1.55 1.3
                         1.5
                                1.47 1.48
                                                5
                                                      5 0.46
## 1
                                            0
                                                                 -0.50.23
t.test(df$Donation ~ df$RecNapvsRecPol)
##
##
   Welch Two Sample t-test
##
## data: df$Donation by df$RecNapvsRecPol
## t = -0.71862, df = 63.996, p-value = 0.475
## alternative hypothesis: true difference in means is not equal to 0
## 95 percent confidence interval:
## -0.8705319 0.4099258
## sample estimates:
## mean in group RecNap mean in group RecPol
               1.321212
                                    1.551515
wilcox.test(df$Donation ~ df$RecNapvsRecPol)
## Warning in wilcox.test.default(x = c(2, 0.5, 0, 2, 1, 2, 3.5, 2, 3, 0,
```

0.5, : cannot compute exact p-value with ties

```
##
## Wilcoxon rank sum test with continuity correction
##
## data: df$Donation by df$RecNapvsRecPol
## W = 486, p-value = 0.4462
## alternative hypothesis: true location shift is not equal to 0
```

We cannot reject the null that Donations in recommendation treatments informing about the political mandate of the source are equal to donations in recommendations treatments providing the name and picture of the source.

For Defaults: Donations in Political treatments < Donations in Name and Picture treatments

```
describeBy(df$Donation, df$DefNapvsDefPol)
## group: DefNap
    vars n mean
                   sd median trimmed mad min max range skew kurtosis
       1 34 1.36 1.11 1.45
                                1.28 1.19 0 4
                                                      4 0.3
## group: DefPol
    vars n mean sd median trimmed mad min max range skew kurtosis se
       1 31 1.55 1.68
                           1
                                1.24 1.48
                                            0
                                              7
                                                      7 1.55
                                                                 2.05 0.3
t.test(df$Donation ~ df$DefNapvsDefPol)
##
##
   Welch Two Sample t-test
## data: df$Donation by df$DefNapvsDefPol
## t = -0.51546, df = 51.451, p-value = 0.6084
## alternative hypothesis: true difference in means is not equal to 0
## 95 percent confidence interval:
## -0.8989215 0.5315591
## sample estimates:
## mean in group DefNap mean in group DefPol
##
              1.364706
                                   1.548387
wilcox.test(df$Donation ~ df$DefNapvsDefPol)
## Warning in wilcox.test.default(x = c(2, 1, 0.7, 2, 3.5, 1, 0.5, 2, 0, 3, :
## cannot compute exact p-value with ties
##
##
   Wilcoxon rank sum test with continuity correction
## data: df$Donation by df$DefNapvsDefPol
## W = 538, p-value = 0.8883
## alternative hypothesis: true location shift is not equal to 0
```

We cannot reject the null that donations in default treatments informing about the political mandate of the source are equal to donations in default treatments providing the name and picture of the source.

H3-2

When the source is political the share of subjects converging to the default is lower than the share of subjects converging to the recommendation.

Donations in default treatments informing about the political characteristics of the source < donations in recommendation treatments informing about the political characteristics of the source

```
describeBy(df$Donation, df$RecPolvsDefPol)
## group: RecPol
   vars n mean sd median trimmed mad min max range skew kurtosis
      1 33 1.55 1.3
                        1.5
                               1.47 1.48
                                           0
                                               5
                                                     5 0.46
## group: DefPol
                   sd median trimmed mad min max range skew kurtosis se
    vars n mean
       1 31 1.55 1.68
                           1
                                1.24 1.48
                                            0
                                              7
                                                      7 1.55
t.test(df$Donation ~ df$RecPolvsDefPol)
##
##
   Welch Two Sample t-test
## data: df$Donation by df$RecPolvsDefPol
## t = 0.0083167, df = 56.497, p-value = 0.9934
## alternative hypothesis: true difference in means is not equal to 0
## 95 percent confidence interval:
## -0.7501750 0.7564311
## sample estimates:
## mean in group RecPol mean in group DefPol
##
              1.551515
                                   1.548387
wilcox.test(df$Donation ~ df$RecPolvsDefPol)
## Warning in wilcox.test.default(x = c(3, 1, 0.5, 3.5, 0, 0, 0.5, 0, 2, 1, :
## cannot compute exact p-value with ties
##
## Wilcoxon rank sum test with continuity correction
## data: df$Donation by df$RecPolvsDefPol
## W = 546.5, p-value = 0.6382
## alternative hypothesis: true location shift is not equal to 0
```

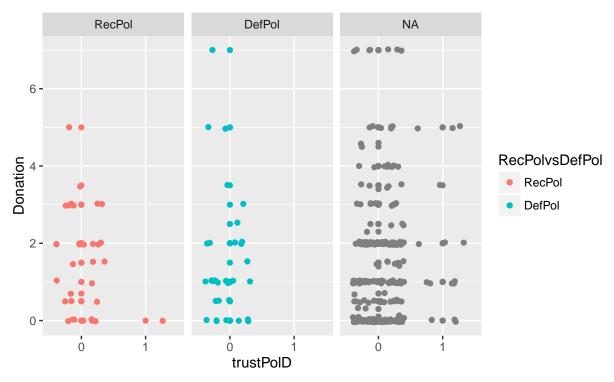
We cannot reject the null that donations in default treatments informing about the political characteristics of the source are equal to donations in recommendation treatments informing about the political characteristics of the source.

H3a (HERE ALSO INCLUDE PARTY TREATMENTS, NOT JUST POLITICAL?)

A subject that scores high on trust in politics is more likely to converge to the recommended and defaulted payment-values, than a subject scoring low on trust in politics. In treatments informing about the political characteristics of the source.

Relationship between trust in politics dummy and Donation

resp. for RecPol and DefPol treatment groups



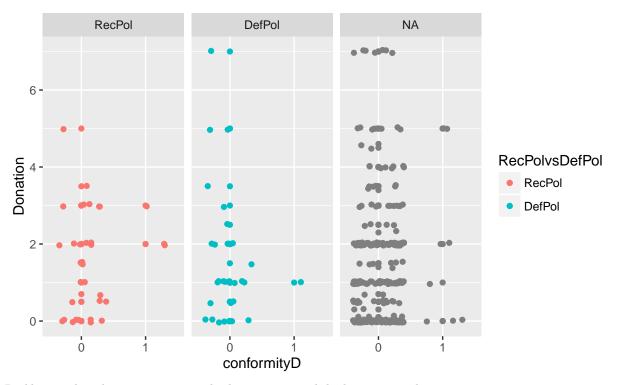
Problem is that there are not enough observations with high trust in politics (no observation in Default x Political treatment.

H3b

A subject that values conformity, i.e. doing what the majority does, is more likely to converge to the recommended and defaulted payment-values, than a subject that does not value conformity.

Relationship between conformity dummy and Donation

resp. for RecPol and DefPol treatment groups



Problem is that there are not enough observations with high trust in politics.

H4

The share of subjects converging to the recommended, respectively defaulted payment-values, relative to the political-characteristic condition, is higher for subjects with same party preferences, and lower for subjects with different party preferences. **Hypothesis is possibly phrased wrongly.**

table(df\$party)

##			
##	AfD	Andere	Bündnis90/Grüne
##	10	21	72
##	CDU/CSU	Die Linke	FDP
##	58	67	11
## Keir	ne (Nichtwähler)	Keine Angabe	SPD
##	45	7	70

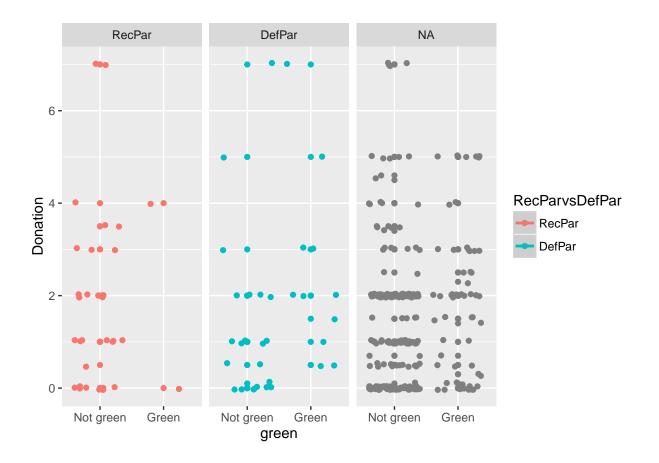
table(df\$party, df\$treatment)

##								
##		Control	RecNos	DefNos	RecNap	DefNap	${\tt RecPol}$	DefPol
##	AfD	2	1	0	2	0	0	2
##	Andere	2	3	1	3	1	1	1
##	Bündnis90/Grüne	6	6	7	8	7	7	6

```
CDU/CSU
##
                                 5
                                         7
                                                 4
                                                        2
                                                                5
                                                                                5
##
     Die Linke
                                  7
                                         4
                                                 6
                                                         5
                                                                8
                                                                                5
                                                                        8
     FDP
                                                                                0
##
                                  0
                                                 2
                                                                1
                                                                        0
                                                         1
##
     Keine (Nichtwähler)
                                  5
                                         5
                                                 6
                                                         3
                                                                3
                                                                        2
                                                                                3
                                                                                2
     Keine Angabe
                                                 3
                                                                0
                                                                        0
##
                                  1
                                         0
                                                         0
                                                                9
##
     SPD
                                  3
                                         4
                                                 5
                                                         9
                                                                       11
                                                                                7
##
##
                           RecPar DefPar RecKno DefKno
     AfD
##
                                 0
                                        0
     Andere
##
                                 2
                                        1
                                                3
                                                        3
                                                5
                                                        7
##
     Bündnis90/Grüne
                                 2
                                       11
##
     CDU/CSU
                                 6
                                        7
                                               10
                                                        3
##
     Die Linke
                                 8
                                        5
                                                7
                                                        4
                                        0
                                                        3
##
     FDP
                                 0
                                                3
##
     Keine (Nichtwähler)
                                 7
                                        5
                                                3
                                                        3
##
     Keine Angabe
                                        0
                                 1
                                                0
                                                        0
##
     SPD
                                        5
```

chisq.test(table(df\$party, df\$treatment))

```
## Warning in chisq.test(table(df$party, df$treatment)): Chi-squared
## approximation may be incorrect
##
## Pearson's Chi-squared test
## data: table(df$party, df$treatment)
## X-squared = 83.27, df = 80, p-value = 0.3792
##
               Control RecNos DefNos RecNap DefNap RecPol DefPol RecPar
##
##
     Not green
                    25
                           25
                                  27
                                          25
                                                 27
                                                        26
                                                               25
                                                                      30
                     6
                            6
                                   7
                                          8
                                                 7
                                                         7
                                                                6
                                                                       2
##
     Green
##
               DefPar RecKno DefKno
##
##
                   23
                          29
     Not green
                           5
                                  7
##
     Green
                   11
```



Further Statistics and Tests

Compare observations that believe we cooperated with Julia Verlinden vs. those who don't

Variable: Donation amount

```
## group: Ja
         n mean sd median trimmed mad min max range skew kurtosis se
       1 156 1.3 1.27
                          1
                                1.14 1.48 0 5
                                                   5 0.9
## group: Nein
          n mean sd median trimmed mad min max range skew kurtosis
                                1.44 1.48 0
                                                    7 1.33
       1 113 1.75 1.84
                         1.5
                                               7
                                                                1.37 0.17
##
##
  Welch Two Sample t-test
##
## data: df$Donation by df$believe2
## t = -2.2729, df = 186.78, p-value = 0.02417
## alternative hypothesis: true difference in means is not equal to 0
## 95 percent confidence interval:
## -0.85041305 -0.06011565
## sample estimates:
```

```
## mean in group Ja mean in group Nein
## 1.298718 1.753982

##
## Wilcoxon rank sum test with continuity correction
##
## data: df$Donation by df$believe2
## W = 7952.5, p-value = 0.1631
## alternative hypothesis: true location shift is not equal to 0
```

Participants who believe we cooperated with Julia Verlinden have a **lower** mean Donation, and also a **lower** variance. The difference is significant (p < .1) judged by the Welch Two Sample t-test, but insignificant judged by the Wilcoxon-Mann-Whitney-U test.

Variable: Decision to donate

```
##
##
                  Ja Nein
##
    Not donated 44
                       31
##
    Donated
                 112
                       82
chisq.test(table(df$Donated, df$believe2))
##
##
   Pearson's Chi-squared test with Yates' continuity correction
##
## data: table(df$Donated, df$believe2)
## X-squared = 2.3598e-06, df = 1, p-value = 0.9988
```

The Chi²-Test is not significant. This implies that the decision whether or not to contribute anything vs. nothing is not dependent on the answer to the question whether the respondent believed that we really cooperated with Julia Verlinden.

Decision to donate for subjects seeing a recommendation vs. subjects seeing a default, irrespective of source

```
table(df$Donated, df$RecvsDef)
##
##
                 Rec Def
    Not donated 48 52
##
##
    Donated
                115 115
chisq.test(table(df$Donated, df$RecvsDef))
##
   Pearson's Chi-squared test with Yates' continuity correction
##
##
## data: table(df$Donated, df$RecvsDef)
## X-squared = 0.04587, df = 1, p-value = 0.8304
```

The Chi²-Test is not significant. This implies that the decision whether or not to contribute anything vs. nothing is not dependent on whether the subjects encountered a recommendation or a default value with our without any specific source or information on the source.

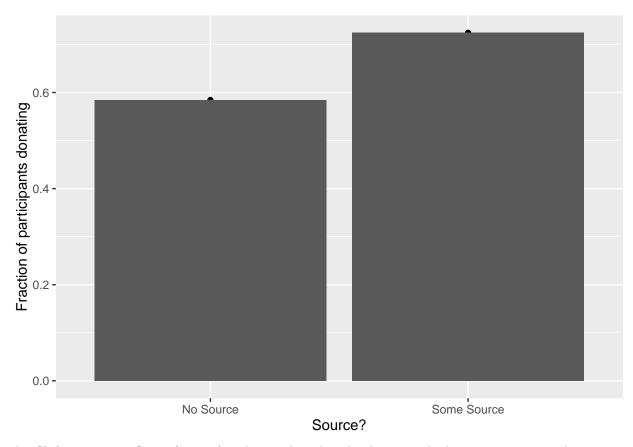
Decision to donate for subjects seeing a recommendation vs. subjects seeing a default, with non-political source-information

```
##
##
                 Non-political/partisan Rec Non-political/partisan Def
##
     Not donated
                                          31
                                          67
                                                                      65
##
     Donated
##
##
   Pearson's Chi-squared test with Yates' continuity correction
##
## data: table(df$Donated, df$RecvsDefNonPolPar)
## X-squared = 0.29534, df = 1, p-value = 0.5868
```

The Chi²-Test is not significant. This implies that the decision whether or not to contribute anything vs. nothing is not dependent on whether the subjects encountered a non-political or non-partisan recommendation or a respective default value.

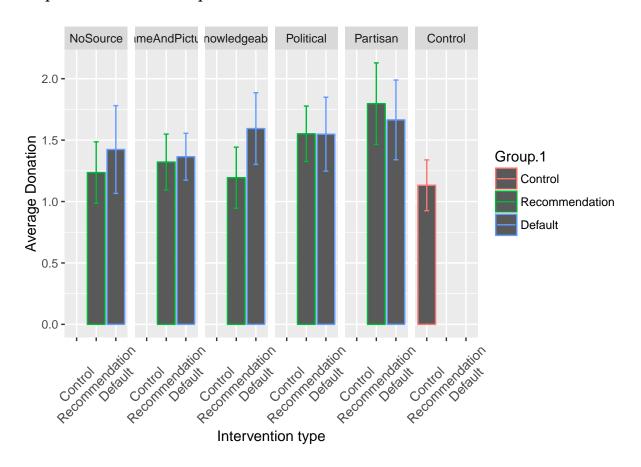
Decision to donate for subjects seeing an intervention without source-information vs. some source-information

```
##
##
                 No Source Some Source
##
     Not donated
                        27
                                    73
##
     Donated
                        38
                                    192
##
##
   Pearson's Chi-squared test with Yates' continuity correction
##
## data: table(df$Donated, df$NosvsSome)
## X-squared = 4.1982, df = 1, p-value = 0.04047
## Warning in Ops.factor(left, right): '/' not meaningful for factors
```



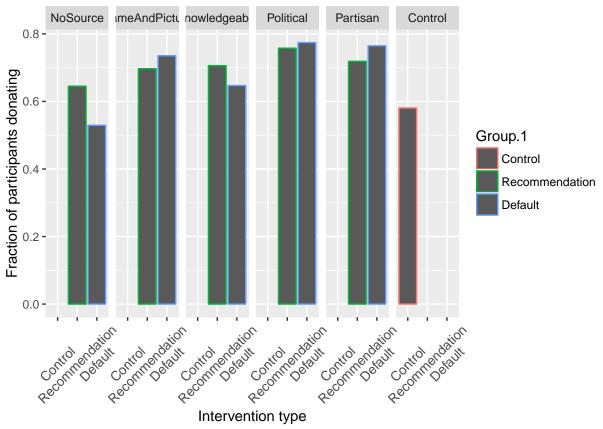
The Chi² test is significant (p < .1). This implies that the decision whether or not to contribute or not depends on whether or not some source-information vs. no source-information is provided.

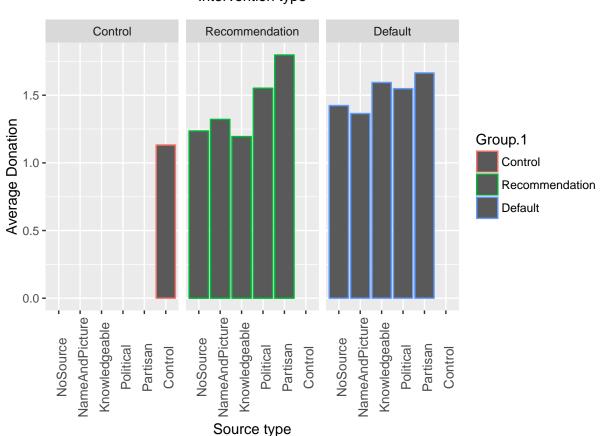
Graphs in order to see potential interactions



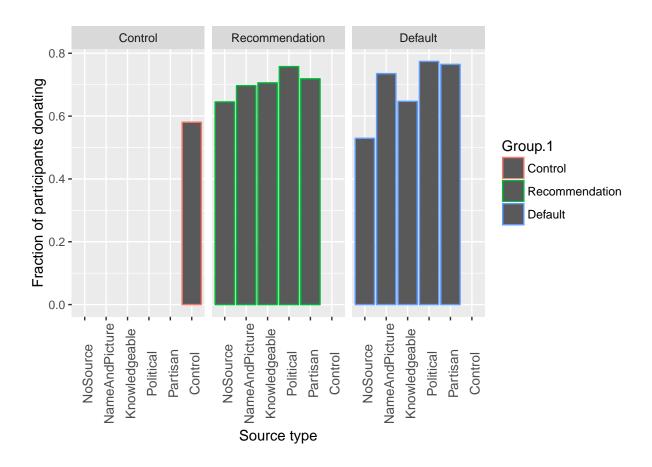
```
## Warning in Ops.factor(left, right): ^{\prime}/^{\prime} not meaningful for factors
```

^{##} Warning in Ops.factor(left, right): $^{\prime}/^{\prime}$ not meaningful for factors

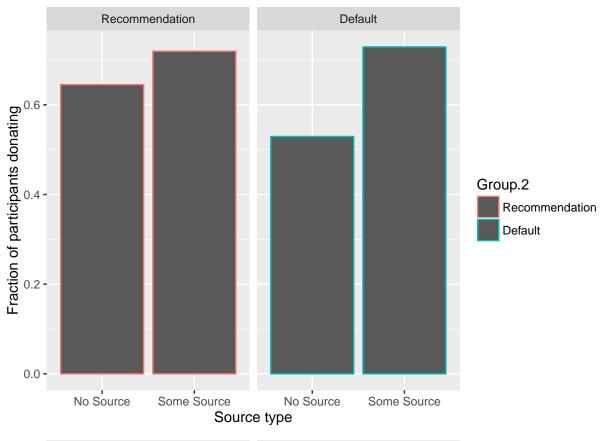


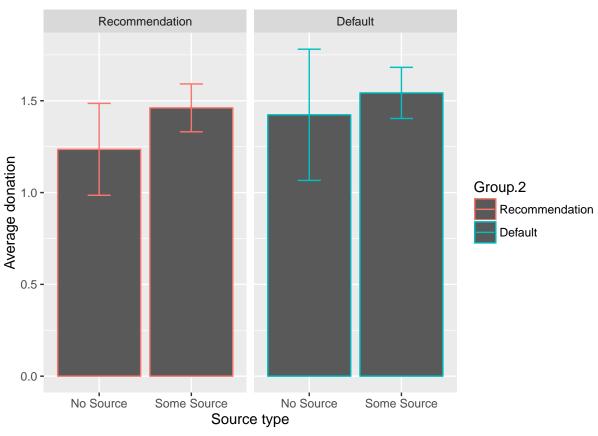


Warning in Ops.factor(left, right): '/' not meaningful for factors
Warning in Ops.factor(left, right): '/' not meaningful for factors



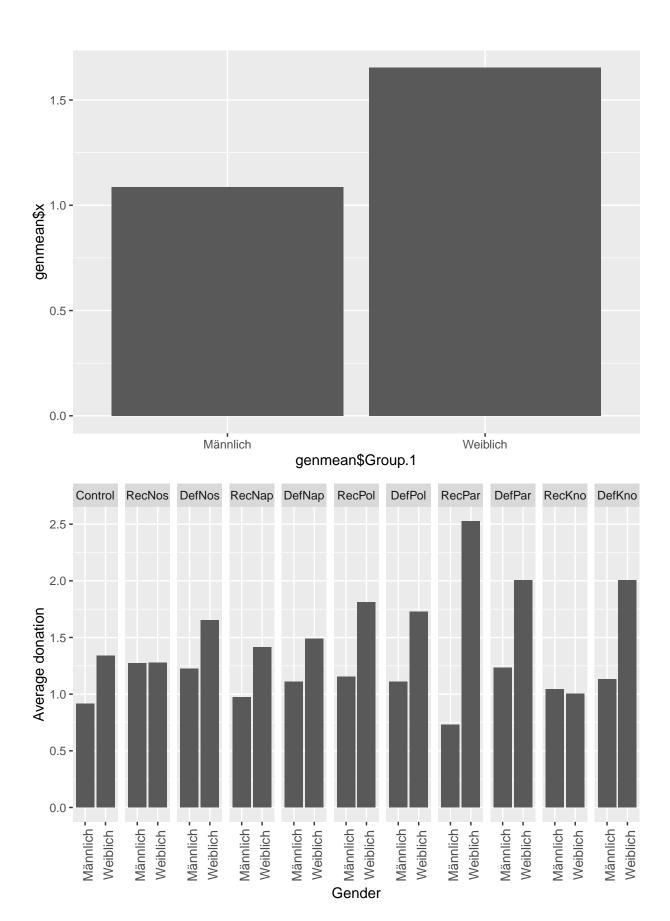
- ## Warning in Ops.factor(left, right): '/' not meaningful for factors
- ## Warning in Ops.factor(left, right): '/' not meaningful for factors





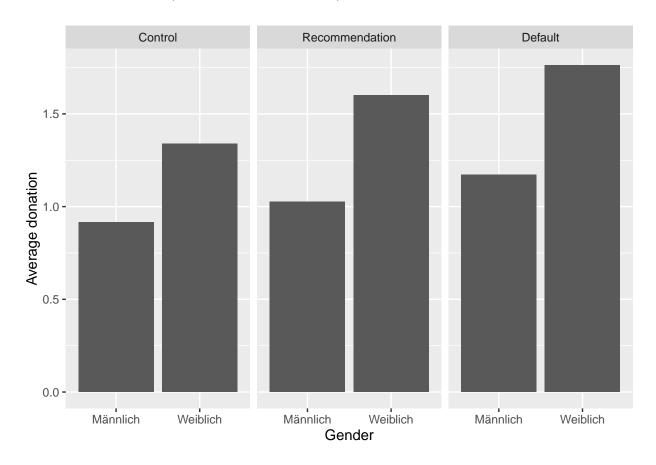
Gender differences (by treatment)

```
## group: Keine Angabe
## vars n mean sd median trimmed mad min max range skew kurtosis se
## group: Männlich
## vars n mean sd median trimmed mad min max range skew kurtosis se
## 1 1 145 1.09 1.37 0.7 0.84 1.04 0 7 7 1.83 4.14 0.11
## -----
## group: Weiblich
## vars n mean sd median trimmed mad min max range skew kurtosis
##
## Welch Two Sample t-test
##
## data: dfsub$Donation by dfsub$gender
## t = -3.594, df = 336.21, p-value = 0.0003743
\#\# alternative hypothesis: true difference in means is not equal to 0
## 95 percent confidence interval:
## -0.8785382 -0.2570311
## sample estimates:
## mean in group Männlich mean in group Weiblich
             1.085517
                               1.653302
##
## Wilcoxon rank sum test with continuity correction
##
## data: dfsub$Donation by dfsub$gender
## W = 11718, p-value = 9.388e-05
\#\# alternative hypothesis: true location shift is not equal to 0
```

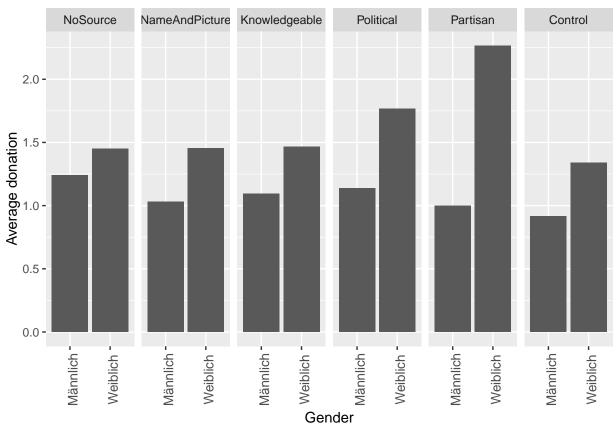


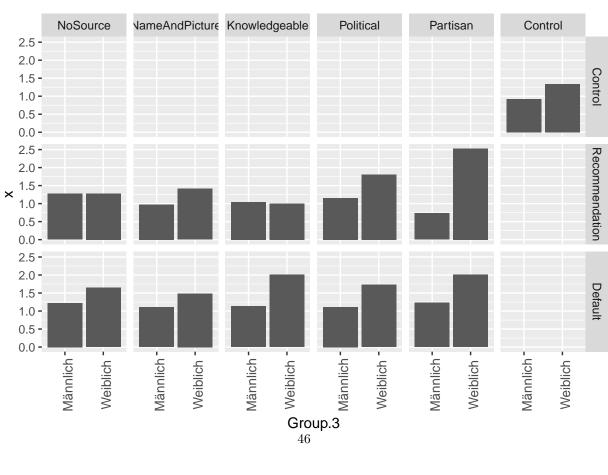
Could be interesting. For example, when looking at Control, RecNos, RecNap, RecPol, RecPar, (And RecKno), the treatment-effect seems to be present for women, but not (or eben negatively) for men. This seems to be also true for Defaults, but slightly different

Gender differences (by Intervention type)



Gender differences (by Source type)





Same as above, the treatment effect seems to be present for females, but only when the source is political or associated with the green party. For males, the treatment effect seems to be zero, or slightly negative.

Regression analysis of interaction effect between treatment and gender

```
summary(lm(Donation ~ treatment*gender, dfsub))
##
## Call:
## lm(formula = Donation ~ treatment * gender, data = dfsub)
## Residuals:
##
       Min
                10 Median
                                3Q
                                        Max
  -2.5263 -1.1100 -0.2789
                            0.7778
##
                                   5.7778
##
## Coefficients:
##
                                   Estimate Std. Error t value Pr(>|t|)
## (Intercept)
                                   0.916667
                                               0.437058
                                                          2.097
                                                                  0.0367 *
## treatmentRecNos
                                   0.356061
                                               0.631985
                                                          0.563
                                                                  0.5735
## treatmentDefNos
                                   0.305556
                                               0.564239
                                                          0.542
                                                                  0.5885
## treatmentRecNap
                                   0.056667
                                               0.586375
                                                          0.097
                                                                  0.9231
## treatmentDefNap
                                   0.192424
                                               0.631985
                                                          0.304
                                                                  0.7610
## treatmentRecPol
                                   0.237179
                                                          0.391
                                                                  0.6958
                                               0.606090
## treatmentDefPol
                                   0.194444
                                               0.667617
                                                          0.291
                                                                  0.7710
## treatmentRecPar
                                  -0.185897
                                               0.606090 -0.307
                                                                  0.7593
## treatmentDefPar
                                   0.316667
                                               0.586375
                                                          0.540
                                                                  0.5895
## treatmentRecKno
                                   0.125000
                                               0.618093
                                                          0.202
                                                                  0.8399
## treatmentDefKno
                                   0.214583
                                               0.578173
                                                          0.371
                                                                  0.7108
## genderWeiblich
                                   0.422222
                                               0.564239
                                                          0.748
                                                                  0.4548
## treatmentRecNos:genderWeiblich -0.416002
                                               0.804609
                                                         -0.517
                                                                  0.6055
## treatmentDefNos:genderWeiblich
                                                          0.007
                                                                  0.9942
                                   0.005556
                                               0.767448
## treatmentRecNap:genderWeiblich
                                   0.016209
                                               0.778472
                                                          0.021
                                                                  0.9834
## treatmentDefNap:genderWeiblich -0.044357
                                               0.791463 -0.056
                                                                  0.9553
## treatmentRecPol:genderWeiblich
                                               0.780579
                                                          0.300
                                                                  0.7646
                                   0.233932
## treatmentDefPol:genderWeiblich
                                   0.193939
                                               0.822953
                                                          0.236
                                                                  0.8138
## treatmentRecPar:genderWeiblich
                                                                  0.0809
                                   1.373324
                                               0.784433
                                                          1.751
## treatmentDefPar:genderWeiblich
                                  0.349708
                                               0.769302
                                                          0.455
                                                                  0.6497
## treatmentRecKno:genderWeiblich -0.459127
                                                         -0.584
                                               0.786473
                                                                  0.5598
## treatmentDefKno:genderWeiblich 0.452083
                                               0.767448
                                                          0.589
                                                                  0.5562
## ---
## Signif. codes: 0 '***' 0.001 '**' 0.05 '.' 0.1 ' ' 1
## Residual standard error: 1.514 on 335 degrees of freedom
## Multiple R-squared: 0.0788, Adjusted R-squared: 0.02105
## F-statistic: 1.365 on 21 and 335 DF, p-value: 0.1328
```

Still need to verify whether this is equivalent to using dummy variables for the treatments. However, I do not think that treatment variable is treated as metric.

Party preference differences

Including only "believers" in Julia Verlinden

Treatment interactions with...

- ... personal Interest and Knowledge w.r.t climate protection
- ... warm Glow feeling/ feeling of guilt when (not) protecting the climate
- ... judging carbon offsetting as a efficient way to protect the climate
- ... moral importance of climate protection
- ... income
- ... assessment of sources knowledge and interest w.r.t. climate protection