#### Data Science for Causal Inference

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The Lab @ DC

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### About Me

- Associate Prof of Government (American University)
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- Senior Social Scientist (The Lab @ DC)
- ➤ Fellow in Methodology (US Office of Evaluation Sciences: "OES")

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- Research agenda: political methodology, causal inference, experimental design, experiments in public policy

Name?

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- ▶ Olympic sport you look forward to?

▶ Data Science in Causal Inference

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  - Calloway-Sant'Anna approach

▶ Introduction to several approaches

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- Materials here: https://github.com/ryantmoore/new-directions-berlin

## Data Science in Causal Inference

## Causal Inference Approaches

The "potential outcomes" framework:  $% \left( 1\right) =\left( 1\right) \left( 1\right) \left($ 

# Causal Inference Approaches

The "potential outcomes" framework:

		Would Enroll if	Would Enroll if	
Citize	n Canvass?	Canvass?	No Canvass?	Enroll
1	Yes			Yes
2	Yes			Yes
3	No			No
4	No			No

# Causal Inference Approaches

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2	Yes	Yes	(No)	Yes
3	No	(Yes)	No	No
4	No	(No)	No	No

The "potential outcomes" framework, more abstractly:

					True $\tau$
Unit $i$	Treatment $T$	Y(1)	Y(0)	$Y^{ m obs}$	Y(1) - Y(0)
1	1	10		10	
2	1	20		20	
3	0		15	15	
4	0		5	5	

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2	1	20	(10)	20	10
3	0	(40)	15	15	25
4	0	(20)	5	5	15

The "potential outcomes" framework, more abstractly:  $\frac{1}{2}$ 

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				$\widehat{ATE} = \hat{\bar{ au}} =$	15 - 10 = 5

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- True treatment effect  $\tau_i = Y_i(1) Y_i(0)$
- True average treatment effect  $\sum_{n=1}^{n} \langle Y_{n}(1) \rangle \langle Y_{n}(2) \rangle$

$$\bar{\tau} = \frac{1}{n} \sum_{i=1}^{n} (Y_i(1) - Y_i(0))$$

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(and we'll draw some DAG's, too)

Three tasks of data science:

Description

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- ▶ Prediction

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Models/algorithms central to all three.

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Hernán, Hsu, and Healy (2019)

Description

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### Description

- ▶ Identifying patterns, etc.
- ► E.g., clustering to discover groups

Prediction

Components

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  - ► Inputs/outputs (predictors/outcomes, features/responses, ...)

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- E.g., regression, random forests, neural networks, ...

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- ► E.g., experiments, observational causal designs, ...



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# I finally found it in real life: the consultant who runs OLS in Excel and calls it machine learning

9:17 AM · Jan 31, 2019 · Twitter for iPhone

<b>54</b> Retweets	7 Quote Tweets	<b>511</b> Likes	
$\Diamond$	$\uparrow \downarrow$	$\bigcirc$	ightharpoons



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Don't do this.



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Don't do this.

(Not "machine learning", probably, but models at least ...)

Consider two loaded datasets:

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str(df2)

```
str(df1)

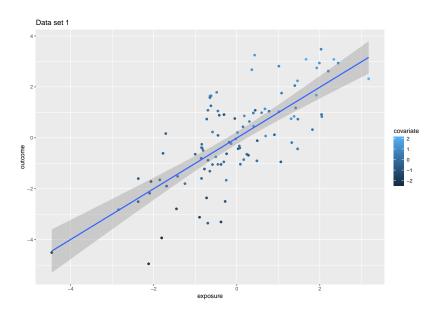
tibble [100 x 3] (S3: tbl_df/tbl/data.frame)
$ covariate: num [1:100] -0.622 1.137 -0.238 1.529 -0.154
$ exposure : num [1:100] 0.0332 0.3627 0.2422 1.4633 0.779
$ outcome : num [1:100] -0.429 2.675 -0.647 2.238 1.044
```

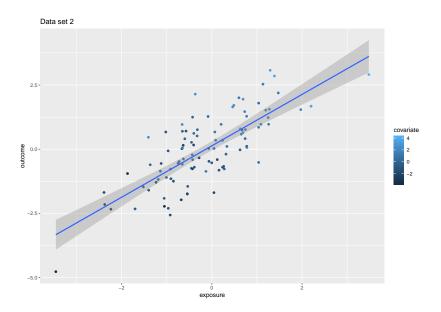
```
tibble [100 x 3] (S3: tbl_df/tbl/data.frame)

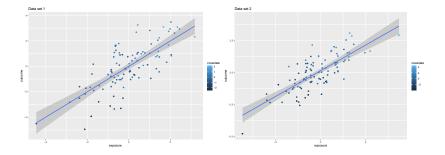
$ exposure : num [1:100] 0.4862 0.0653 -1.4021 -0.546 -0.4

$ outcome : num [1:100] 1.706 0.669 -1.597 -1.733 0.617
```

 $\$  covariate: num [1:100] 2.24 0.924 -0.999 -2.343 0.207 .







#### Model each

```
lm_df1 <- lm(outcome ~ exposure, data = df1)
lm_df2 <- lm(outcome ~ exposure, data = df2)</pre>
```

```
# A tibble: 4 x 4
data term estimate std.error
<chr> <chr> <chr> <chr> 0.00671 0.120
df1 (Intercept) -0.00671 0.120
df1 exposure 0.996 0.0927
df2 (Intercept) 0.133 0.0890
df2 exposure 1.00 0.0841
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▶ Both cases: effect of exposure  $\approx 1$ .

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- ▶ Is this good? Is it correct?

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```

- ▶ Both cases: effect of exposure  $\approx 1$ .
- ▶ Is this good? Is it correct?
- ▶ What if we adjust for covariate?

```
lm_df1_adj <- lm(outcome ~ exposure + covariate, data = df:
lm_df2_adj <- lm(outcome ~ exposure + covariate, data = df:</pre>
```

▶ Both cases: effect of exposure  $\approx 0.5$ .

```
lm_df1_adj <- lm(outcome ~ exposure + covariate, data = df:
lm_df2_adj <- lm(outcome ~ exposure + covariate, data = df:</pre>
```

```
# A tibble: 4 x 4
data term estimate std.error
<chr> <chr> <chr> <chr> 0.501 0.108
df1 exposure 0.501 0.108
df1 covariate 0.970 0.147
df2 exposure 0.554 0.0990
df2 covariate 0.385 0.0598
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- ▶ Both cases: effect of exposure  $\approx 0.5$ .
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- ▶ Both cases: effect of exposure  $\approx 0.5$ .
- ▶ Is this good? Is it correct?
- Which is correct?  $\beta = 1$ ?  $\beta = 0.5$ ?

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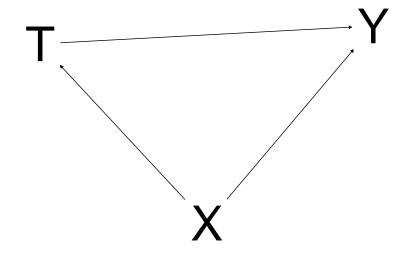
- ▶ Both cases: effect of exposure  $\approx 0.5$ .
- ▶ Is this good? Is it correct?
- ▶ Which is correct?  $\beta = 1$ ?  $\beta = 0.5$ ?
- ► Should we adjust for covariate?

There is nothing in the data that tells us.

There is nothing in the data that tells us. ©

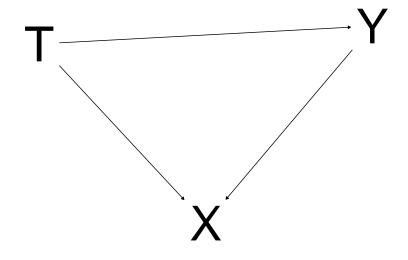
There is nothing in the data that tells us. ©

Here are the true structures: First



There is nothing in the data that tells us. ©

Here are the true structures: Second



When know structures, adjustment sets for unbiasedness differ:

▶ df1: confounding  $\Rightarrow$  adjust for X

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- ▶ df2: collider  $\Rightarrow$  do not adjust for X

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g_conf <- dagitty("dag{ x -> y ; x <- c -> y }")
g_coll <- dagitty("dag{ x -> y ; x -> c <- y }")</pre>
```

- ▶ df1: confounding  $\Rightarrow$  adjust for X
- ▶ df2: collider  $\Rightarrow$  do not adjust for X

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g_conf <- dagitty("dag{ x -> y ; x <- c -> y }")
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```
adjustmentSets(g_conf, "x", "y")
```

```
{ c }
```

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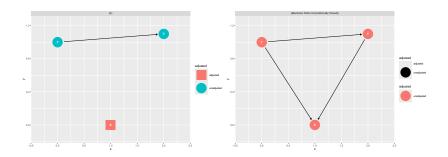
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adjustmentSets(g_coll, "x", "y")
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df2, do not adjust for X,  $\beta = 1$ :

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df1, adjust for X,  $\beta = 0.5$ :

df2, do not adjust for X,  $\beta = 1$ :

(Data from D'Agostino McGowan (2023))

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- ► Importance of identifying "pre-treatment covariates", "proper covariates"; doing "design before analysis"
- ► Importance of experiments: strong knowledge about (part of) causal structure assgn mechanism
- ➤ Causal inference is critical to scientific questions, and separate from prediction
- ➤ Though, methods from prediction can aid causal inference
- ▶ "Causal euphimisms" don't help (Hernán 2018)

# Approaches of Prediction and Causal Inference

Two Cultures, (Breiman 2001)

▶ Data Models: our "social science modeling"

# Approaches of Prediction and Causal Inference

Two Cultures, (Breiman 2001)

- ▶ Data Models: our "social science modeling"
- ▶ Algorithmic Models: our "data science algorithms"

## Methods for Prediction and Causal Inference

- Cross-validation
- ▶ Regression/Decision trees
- ▶ Random forests

James et al. (2021)

k-fold cross-validation to select method

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ightharpoonup Select model that minimises  $CV_{(k)}$ 

```
## Make data
mk_{data} \leftarrow function(n = 90, n_{folds} = 10){
  df <- tibble(
    x1 = rnorm(n),
    x2 = rnorm(n),
    x3 = rnorm(n).
    y = 0.1 * x1 + 0.2 * x2 + 0.5 * x3 + rnorm(n),
    cv_fold = sample(rep(1:n_folds, (n / n_folds)))
df <- mk data()</pre>
```

#### head(df)

```
# A tibble: 6 x 5
     x1
            x2
                  xЗ
                          y cv_fold
  <dbl> <dbl> <dbl> <dbl> <
                              <int>
1 1.35 0.631 -0.448 -1.85
2 -0.805 1.32 -0.981 -1.59
                                 6
3 0.940 -1.09 -0.751 -0.540
 0.610 0.523 -0.363 -0.985
4
5 0.567 0.447 -0.106 0.225
6 0.503 0.349 1.61 0.418
                                  5
```

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6
  0.503 0.349 1.61 0.418
                                  5
```

#### table(df\$cv\_fold)

```
1 2 3 4 5 6 7 8 9 10
9 9 9 9 9 9 9 9 9 9
```

```
cv lm <- function(data, fmla){</pre>
 n folds <- max(data$cv fold)</pre>
  store_mses <- vector("numeric", length = n_folds)</pre>
  for(idx in 1:n folds){
    df_train <- data |> filter(cv_fold != idx)
    df_test <- data |> filter(cv_fold == idx)
    lm_out <- lm(fmla, data = df train)</pre>
    predictions <- predict(lm_out, newdata = df_test)</pre>
    store mses[idx] <- mean((df test$y - predictions)^2)}
  test_error_cv_k <- mean(store_mses)</pre>
  return(test error cv k)
```

```
cv_{lm}(data = df, fmla = y \sim x1 + x2)
```

[1] 1.446812

[1] 1.375958

```
cv_lm(data = df, fmla = y ~ x1 + x2)
[1] 1.446812

df <- mk_data()
cv_lm(df, y ~ x1 + x2)</pre>
```

[1] 0.9928568

```
cv lm(data = df, fmla = y \sim x1 + x2)
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df <- mk data()</pre>
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df <- mk data()</pre>
cv lm(df, y \sim x1 + x2 + x3)
```

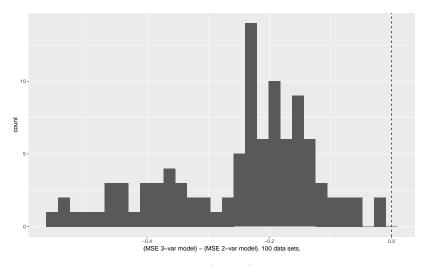


Figure 1: MSE always less (better) for 3-variable model.

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- ► Goal: minimise residual sum of squares (RSS), just like LS regression:

$$\sum_{j=1}^{J} \sum_{i \in R_{j}} \left( y_{i} - \hat{y}_{R_{j}} \right)$$

How to define regions  $R_j$ ?

How to define regions  $R_i$ ?

▶ Top-down, greedy recursive binary split

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$$\sum_{i:x \in R_1(j,s)} \left(y_i - \hat{y}_{R_1(j,s)}\right)^2 + \sum_{i:x \in R_2(j,s)} \left(y_i - \hat{y}_{R_2(j,s)}\right)^2$$

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Sum squared pred. error (plus penalty that grows with tree size) across units in region, then regions.

But, how to choose  $\alpha$ ? (Use cross-validation.)

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- 4. Using that  $\alpha$ , select best subtree from Step 2

Effect of office-holding on wealth (Eggers and Hainmueller 2009):

```
library(qss)
library(rsample)
library(tree)
data("MPs")
mps <- MPs |> mutate(age = yod - yob,
                     is_labour = if_else(party == "labour"
                     is_london = if_else(region == "Greater
                     is_winner = if_else(margin > 0, 1, 0))
  select(ln.net, age, is_labour, is_london, is_winner) |>
  na.omit()
```

```
set.seed(765076184)

mp_split <- initial_split(mps, prop = 0.7)

mp_train <- training(mp_split)

mp_test <- testing(mp_split)</pre>
```

```
tree_mp <- tree(ln.net ~ ., data = mp_train)
plot(tree_mp)
text(tree_mp)</pre>
```

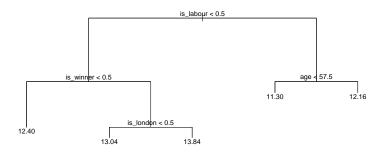


Figure 2: The regression tree (for training data)

Would pruning help?

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```
cv_mps <- cv.tree(tree_mp, K = 10)
plot(cv_mps$size, cv_mps$dev, type = "b")</pre>
```

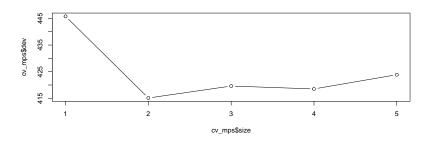


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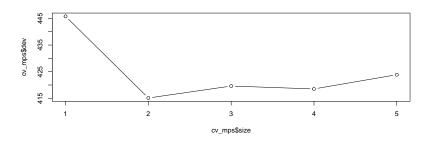


Figure 3: Subtree size 2 minimises SSR

```
prune_mps <- prune.tree(tree_mp, best = 2)

plot(prune_mps)
text(prune_mps)</pre>
```



Figure 4: The pruned tree

#### Predict for test set:

► MSE for pruned: 1.922

▶ MSE for full: 1.945

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(Pretty good for 1 split!?)

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Next: random forest algorithm

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Bagging: bootstrap aggregation

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- ► (Linear regression: lower variance)

Random forests: decorrelated, bagged trees

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- Build deep tree. At each split, randomly sample m of p predictors, build split from only those m.

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- ightharpoonup (Often choose  $m \approx \sqrt{p}$ )
- So, different splits consider different predictors
- So, trees will look very different to each other

```
library(randomForest)
# Full bag:
bag mps <- randomForest(ln.net ~ ., data = mp train,</pre>
                         ntree = 500, mtry = 4,
                         importance = TRUE)
# Decorrelate:
rf mps <- randomForest(ln.net ~ ., data = mp train,
                        ntree = 500, mtry = 2,
                        importance = TRUE)
```

Predict:

```
preds_bag <- predict(bag_mps, newdata = mp_test)
preds_rf <- predict(rf_mps, newdata = mp_test)</pre>
```

- MSE for RF: 1.995
- ▶ MSE for full bag: 2.536

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## Heterogeneous Treatment Effects

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- Notationally,  $\exists i : \tau_i \neq \tau$

## Homogeneous and Heterogeneous Effects: Estimation

Homogeneous effects:

Outcome = 
$$\beta_0 + \beta_1$$
Treatment +  $\epsilon$ 

# Homogeneous and Heterogeneous Effects: Estimation

Homogeneous effects:

$$Outcome = \beta_0 + \beta_1 Treatment + \epsilon$$

```
lm_out <- lm(ln.net ~ is_winner, data = mps)
lm_out</pre>
```

```
Call:
lm(formula = ln.net ~ is_winner, data = mps)
Coefficients:
(Intercept) is_winner
    12.2464    0.5176
```

# Homogeneous and Heterogeneous Effects: Estimation

Homogeneous effects:

```
t.test(ln.net ~ is_winner, data = mps)
```

Welch Two Sample t-test

```
data: ln.net by is_winner

t = -3.9552, df = 287.65, p-value = 9.636e-05

alternative hypothesis: true difference in means between the second terms of the second term
```

# Homogeneous and Heterogeneous Effects: Estimation Homogeneous effects:

$$\text{Outcome} = \beta_0 + \beta_1 \text{Treatment} + \sum \beta_j X_j + \epsilon$$

Homogeneous effects:

Outcome = 
$$\beta_0 + \beta_1$$
Treatment +  $\sum \beta_j X_j + \epsilon$ 

lm(formula = ln.net ~ is\_winner + is\_labour + is\_london + a
 data = mps)

#### Coefficients:

0.00

Homogeneous effects:

lm\_lin(ln.net ~ is\_winner, covariates = ~ is\_labour + is\_le

Estimate Std. Error

12.111785723 12.42195044 416 0.088075873 0.60390123 416

-0.461346226 0.13861367 416

-0.249106208 0.73457813 416

t val

		Dod. Error	0 141
(Intercept)	1.226687e+01	0.078894901	155.4836617
is_winner	3.459885e-01	0.131207672	2.6369536
is_labour_c	-1.613663e-01	0.152608515	-1.057387
is_london_c	2.427360e-01	0.250214401	0.9701118
age_c	4.740367e-03	0.007031323	0.6741786
<pre>is_winner:is_labour_c</pre>	-9.104022e-01	0.264395760	-3.4433313
<pre>is_winner:is_london_c</pre>	-8.847770e-02	0.426241818	-0.2075763
is_winner:age_c	-4.778657e-05	0.012753800	-0.0037468
	CI Lower	CI Upper	DF

(Intercept)

is\_labour\_c is london c

is\_winner

### CATEs: Conditional ATEs

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- ➤ Sometimes "CACE"
- Inference: not "evidence against TE = 0?", but "evidence against  $CATE_1 = CATE_2$ ?"

Heterogeneous effects:

 $\label{eq:outcome} \text{Outcome} = \beta_0 + \beta_1 \text{Treatment} + \beta_2 \text{Group} + \beta_3 \text{Treatment} \cdot \text{Group} + \epsilon$ 

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Heterogeneous effects:

 $Outcome = \beta_0 + \beta_1 Treatment + \beta_2 Group + \beta_3 Treatment \cdot Group + \epsilon$ 

- $\triangleright$   $\beta_1$  gives TE for Group == 0
- $\triangleright$   $\beta_1 + \beta_3$  gives TE for Group == 1

#### Heterogeneous effects:

```
lm out <- lm(ln.net ~ is_winner * is_labour +</pre>
                is_london + age, data = mps)
coef(lm_out) |> round(3)
```

```
(Intercept)
                       is winner
                                             is labour
     11.959
                           0.780
        age is_winner:is_labour
      0.005
                          -0.914
```

-0.16

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$$\hat{y}_{R_j} = \frac{1}{|R_j|} \sum_{i \in R_j} Y_i$$

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$$Y(0), Y(1) \perp \!\!\!\perp T \mid \mathbf{X}$$

▶ Let  $\{T, R_i\} = \{i : T_i = 1, i \in R_i\}$  (Tr obs in  $R_i$ )

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$$\hat{\bar{\tau}}_{R_j} = \frac{1}{|\{T,R_j\}|} \sum_{\{T,R_j\}} Y_i - \frac{1}{|\{C,R_j\}|} \sum_{\{C,R_j\}} Y_i$$

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So, we can use RF methods to estimate conditional (heterogeneous) treatment effects, CATEs.

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  - ▶ Prediction, estimation of  $\hat{\bar{\tau}}$  uses only  $\mathcal{I}$
- $\blacktriangleright$  Build a random forest (decorrelated deep trees picking from m predictors) of causal trees

```
library(grf)
X <- mp_train |> select(age, is_labour, is_london)
W <- mp_train |> select(is_winner) |>
  unlist() |> as.numeric()
Y <- mp_train |> select(ln.net) |> unlist()
cf out <- causal forest(X, Y, W)
```

```
cf_out
```

```
cf_out
```

("How frequently was i the split feature?")

```
X test <- mp test |> select(age, is labour, is london)
cf pred est var <- predict(cf out, X test,
                            estimate.variance = TRUE)
cf preds <- cf pred est var$predictions
df cf <- tibble(X test,</pre>
                cf te = cf preds,
                cf se = sqrt(cf pred est var$variance.
                te 1se lower = cf te - cf se,
                te 1se upper = cf te + cf se)
```

Avg pred treatment effect in test sample:

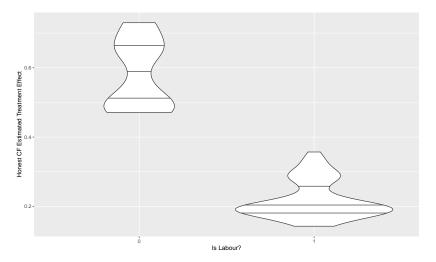
```
mean(cf_preds)
```

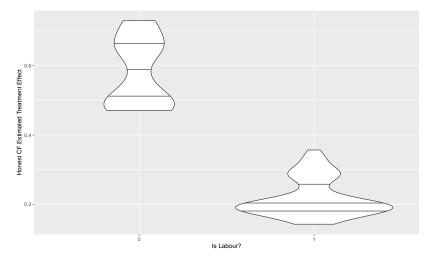
[1] 0.405236

A doubly-robust ATE from training sample:

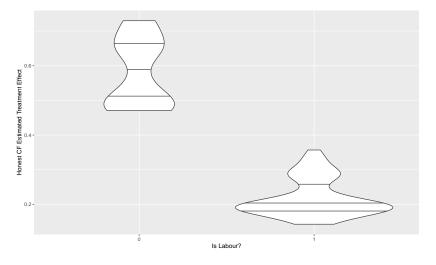
```
average_treatment_effect(cf_out)
```

estimate std.err 0.3465627 0.1715298

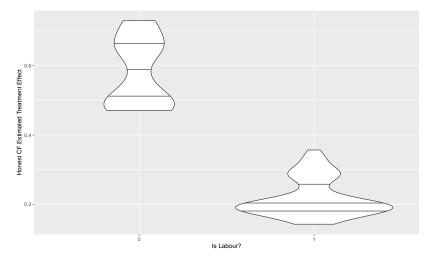




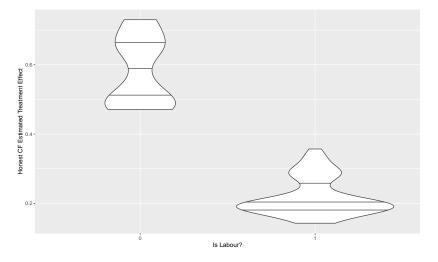
▶ Mean CF TE, Tory: 0.58



▶ Mean CF TE, Tory:  $0.58 \rightsquigarrow £192,000$ 



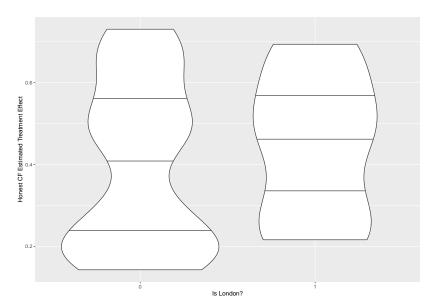
- ► Mean CF TE, Tory: 0.58 → £192,000
- ▶ Mean CF TE, Labour: 0.219



- ► Mean CF TE, Tory: 0.58 → £192,000
- ▶ Mean CF TE, Labour:  $0.219 \rightsquigarrow £60,000$

```
average_treatment effect(
  cf out,
  subset = X$is labour == 0)
estimate std.err
0.7503034 0.2237473
average treatment effect(
  cf out,
  subset = X$is labour == 1)
  estimate std.err
-0.1033197 0.2589494
```

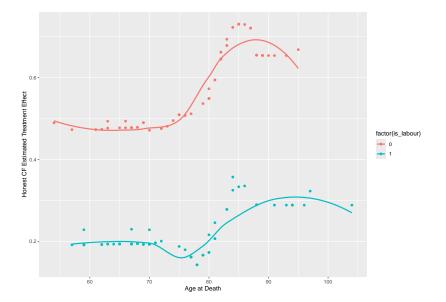
# Example: Causal Forests Results, London



# Example: Causal Forests Results, London

```
average treatment effect(
 cf out,
  subset = X[, "is london"] == 1)
estimate std.err
1.1156002 0.3778383
average treatment effect(
  cf out,
 subset = X[, "is london"] == 0)
estimate std.err
0.2400806 0.1874061
```

# Example: Causal Forests Results, Age





### Feature Selection

▶ Wrappers: pick subset of covars, train on data (estimate model), test on hold-out, score predictions. Keep best-scoring subset.

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- Filters: correlate covars with outcome. Keep strongest.
- ▶ Embeds: select features and estimate model at same time. Penalize using more predictors.

# Regularization Methods

OLS reminder

Minimize SSR:

$$\begin{aligned} & \operatorname{argmin}_{\beta} \sum_{i=1}^{n} \left( y_{i} - \hat{y}_{i} \right)^{2} \\ & \operatorname{argmin}_{\beta} \sum_{i=1}^{n} \left( \mathbf{y} - \mathbf{X} \hat{\beta} \right)^{2} \end{aligned}$$

L1 regularization: the LASSO (Least Absolute Shrinkage and Selection Operator)

$$\operatorname{argmin}_{\beta} \left[ \sum_{i=1}^{n} \left( y_i - \mathbf{X} \hat{\beta} \right)^2 + \lambda \sum_{j=1}^{k} |\hat{\beta}_j| \right]$$

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L2 regularization: Ridge regression

$$\operatorname{argmin}_{\beta} \left[ \sum_{i=1}^{n} \left( y_i - \mathbf{X} \hat{\beta} \right)^2 + \lambda \sum_{j=1}^{k} \hat{\beta}_j^2 \right]$$

Mix L1 and L2: Elastic net

$$\operatorname{argmin}_{\beta} \left( \frac{\sum\limits_{i=1}^{n} \left( y_i - \mathbf{X} \hat{\beta} \right)^2}{2n} + \lambda \left[ \alpha \sum\limits_{j=1}^{k} |\hat{\beta}_j| + \frac{1-\alpha}{2} \sum\limits_{j=1}^{k} \hat{\beta}_j^2 \right] \right)$$

Mix L1 and L2: Elastic net

$$\operatorname{argmin}_{\beta} \left( \frac{\sum\limits_{i=1}^{n} \left( y_i - \mathbf{X} \hat{\boldsymbol{\beta}} \right)^2}{2n} + \lambda \left[ \alpha \sum\limits_{j=1}^{k} |\hat{\beta}_j| + \frac{1-\alpha}{2} \sum\limits_{j=1}^{k} \hat{\beta}_j^2 \right] \right)$$

Regularized trees, ...

How to choose  $\lambda$ ,  $\alpha$ ?

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Cross-validation for  $\lambda$ :

```
df_lasso <- read_csv("../data/01-lasso.csv")

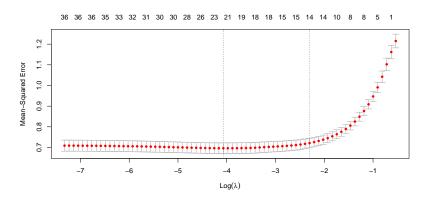
X <- as.matrix(df_lasso[, 2:ncol(df_lasso)])

Y <- as.matrix(df_lasso[, "y"])

library(glmnet)

cv_lasso <- cv.glmnet(X, Y, alpha = 1)</pre>
```

### plot(cv\_lasso)



#### cv\_lasso\$lambda.min

#### [1] 0.0170891

#### Implement:

```
Call: glmnet(x = X, y = Y, alpha = 1, lambda = cv_lasso$1a
```

Df %Dev Lambda 1 21 45.32 0.01709

#### Coefficients:

```
coef_lasso <- coef(lasso_out)</pre>
round(coef_lasso, 3)
37 x 1 sparse Matrix of class "dgCMatrix"
                 s0
(Intercept)
             0.000
x1
              0.112
              0.095
x2
xЗ
              0.086
x4
              0.147
x5
              0.002
              0.063
x6
x7
              0.051
8x
              0.074
x9
              0.042
x10
```

#### Coefficients:

(Intercept)

### round(coef\_lasso[, ], 3)

x18

x24

x30

x36 0.048

0.010

0.000

0.000

0.147	0.086	0.095	0.112	0.000
x10	x9	x8	x7	x6
0.000	0.042	0.074	0.051	0.063
x16	x15	x14	x13	x12
0.000	0.000	0.026	0.000	0.039

x2

x20

x26

x32

-0.015

0.000

0.032

xЗ

x21

x27

x33

0.030

0.000

0.000

x2:

x28

x34

0.000

-0.010

-0.04

x1

x19

x25

x31

0.127

0.000

0.028

#### The LASSO

x3

x4

x5 x6

x7

8x

Implement, alternative  $\lambda$ :

```
lasso_1se <- glmnet(X, Y, alpha = 1,</pre>
                     lambda = cv_lasso$lambda.1se)
coef(lasso_1se)
37 x 1 sparse Matrix of class "dgCMatrix"
                        s0
(Intercept) -0.0003034087
x1
             0.1051188782
x2
             0.0898842045
```

0.0742522801

0.1513883536

0.0603811184

0.0389489143 0.0575738993

#### The LASSO

#### Coefficients:

(Intercent)

0.000

x36 0.030

# round(coef(lasso\_1se)[, ], 3)

	AO	AZ	VI	(Incercebe)
0.	0.074	0.090	0.105	0.000
	х9	8x	x7	х6
0.	0.037	0.058	0.039	0.060
:	x15	x14	x13	x12

15

0.000

	x9	8x	x7	x6
0	0.037	0.058	0.039	0.060
	x15	x14	x13	x12
0	0.000	0.008	0.000	0.029

<del>√</del>1

0.000

x10	x9	8x	x7	x6
0.000	0.037	0.058	0.039	0.060
x16	x15	x14	x13	x12
0.000	0.000	0.008	0.000	0.029
x22	x21	x20	x19	x18
0.000	0.000	0.000	0.039	0.004
<b>∀</b> 28	<b>v</b> 27	<b>v</b> 26	<b>v</b> 25	<b>~</b> 24

0.00	0.000	0.008	0.000	0.029	
x2:	x21	x20	x19	x18	
0.00	0.000	0.000	0.039	0.004	
x28	x27	x26	x25	x24	

0.000	0.000	0.000	0.039	0.004	
x28	x27	x26	x25	x24	
0.000	0.000	0.000	0.000	0.000	
x34	x33	x32	x31	x30	

0.013

0.000

The idea:

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- $\triangleright \approx$  "double robust", "AIPW" estimators
- (different to just "doing LASSO twice" for regularization + shrinkage)

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- 2. Model T = f(X), using LASSO

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- 3. Let  $X_{\text{LASSO}}$  be set of imp covariates identified s.t. each  $\beta_{X_{\text{LASSO}}} > 0$ 4. Model  $Y = T + X_{\text{LASSO}}$

Dear Registered Voter:

#### WHAT IF YOUR NEIGHBORS KNEW WHETHER YOU VOTED?

Why do so many people fail to vote? We've been talking about the problem for years, but it only seems to get worse. This year, we're taking a new approach. We're sending this mailing to you and your neighbors to publicize who does and does not vote.

The chart shows the names of some of your neighbors, showing which have voted in the past. After the August 8 election, we intend to mail an updated chart. You and your neighbors will all know who voted and who did not.

#### DO YOUR CIVIC DUTY - VOTE!

MAPLE DR	Aug 04	Nov 04	Aug 06
9995 JOSEPH JAMES SMITH	Voted	Voted	
9995 JENNIFER KAY SMITH		Voted	
9997 RICHARD B JACKSON		Voted	
9999 KATHY MARIE JACKSON		Voted	
9999 BRIAN JOSEPH JACKSON		Voted	
9991 JENNIFER KAY THOMPSON		Voted	
9991 BOBR THOMPSON		Voted	
9993 BILLS SMITH			
9989 WILLIAM LUKE CASPER		Voted	
9989 JENNIFER SUE CASPER		Voted	
9987 MARIA S JOHNSON	Voted	Voted	

# The Double LASSO for Treatment Effects: Example library(hdm) library(qss) data(social)

```
df_social <- social |>
  mutate(is_male = if_else(sex == "male", 1, 0),
```

age = 2006 - yearofbirth,
<pre>is_neighbors = if_else(messages == "Neighbors" filter(messages %in% c("Neighbors", "Control"))</pre>
<pre>df_social  &gt; select(-yearofbirth)  &gt; head()</pre>
·
sex primary2004 messages primary2006 hhsize is_male

Control

Control

Control

Control

Control

3 3

male

male

male

female

2 female

3

5

```
rlasso_out <- rlassoATE(
  primary2006 ~ age + is_male + primary2004 + hhsize +
    is_neighbors | age + is_male + primary2004 + hhsize,
  data = df_social)</pre>
```

```
summary(rlasso out)
```

```
Estimation and significance testing of the treatment effect
Type: ATE
Bootstrap: not applicable
coeff. se. t-value p-value
```

TE 0.080091 0.002625 30.51 <2e-16 \*\*\*

---

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

```
X <- as.matrix(df_social[, c("age", "is_male", "primary2004</pre>
                            "hhsize", "is neighbors")])
Y <- as.matrix(df social[, "primary2006"])
D <- as.matrix(df social[, "is neighbors"])</pre>
summary(rlassoEffects(X, Y, method = "double selection"))
[1] "Estimates and significance testing of the effect of ta
            Estimate. Std. Error t value Pr(>|t|)
          0.0038449 0.0000681 56.456 < 2e-16 ***
age
is male 0.0086763 0.0018889 4.593 4.36e-06 ***
primary2004 0.1474364 0.0019924 74.000 < 2e-16 ***
hhsize 0.0004260 0.0012618 0.338 0.736
is neighbors 0.0802361 0.0026278 30.534 < 2e-16 ***
```

Is\_neignbors 0.0802361 0.0026278 30.534 < 2e-16 \*\*\*
--Signif. codes: 0 '\*\*\*' 0.001 '\*\*' 0.01 '\*' 0.05 '.' 0.1 '

R packages for Regularization, etc.

- ▶ glmnet
- caret

See also tidymodels, parsnip,  $\dots$ 

Thanks!

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