DML Applications

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- Experimental design: treatment is random, so the propensity score is known and bias is minimized
- · Observational design: bias can arise from misspecification of either model

Causal Model Choice

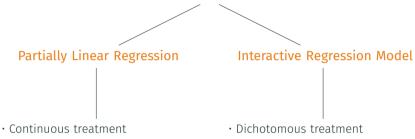
Partially Linear Regression

- Continuous treatment
- No heterogeneous treatment effects
- No functional form assumption for controls
- · Command: DoubleMLPLR

Interactive Regression Model

- · Dichotomous treatment
- Allows heterogeneous treatment effects
- No functional form assumption for controls
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· Clustered Data: DML assumes i.i.d.; assign whole clusters to folds and use cluster-robust SEs (Command: DoubleMLClusterData).

Experimental design

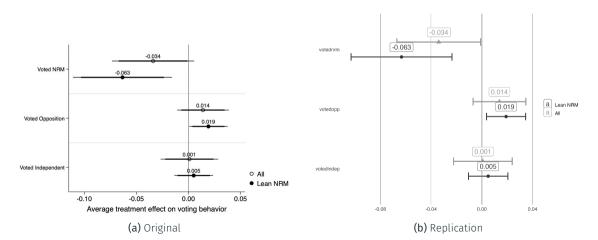
Platas and Raffler (2021): Closing the Gap

Experiment on the effects of providing information in dominant party regimes.

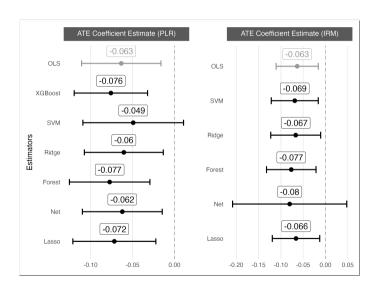
- · Information asymmetry in dominant party regimes (in favor of the regime).
- Can provision of information about all parties (especially opposition ones), reduce the asymmetry in terms of voter knowledge, opposition likability or vote intentions.
- 11 constituencies in Uganda. Within each constituency randomly assign some villages to a screening of parliamentary candidate debates.

Main relationship:

- · Vote intention \sim Received Informational Treatment (0/1)
- \cdot Vote (Incumbent/Opposition/Indep.) \sim Treatment + Treatment x Covariates.



DML application for experiment



Robustness and Application Notes

- The result we replicated have a binary treatment and a control. The classification versions of the machine learning models will likely perform better here.
- For the DoubleMLIRM function, treatment is assumed to be binary, so the learner used to model the propensity score HAS to be a classification learner, not a regression one (e.g. use classif.cv_glmnet instead of regr.cv_glmnet).
- Just make a mental note that such differences in data type matter for exploring pairs of learners when using IRM vs. PLR.

Observational design

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Design

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- · Condition on observables.

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Additional:

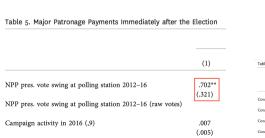
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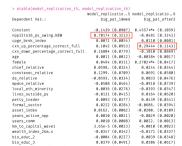
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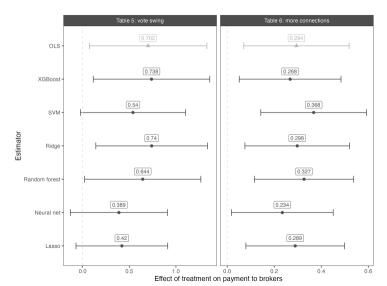




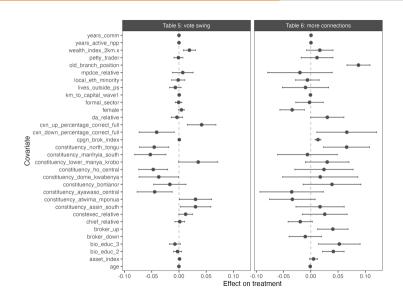


DML application for observational study

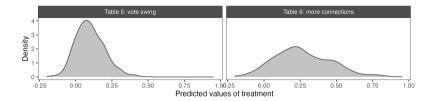
Five-fold cross-fitting and 20 repetitions



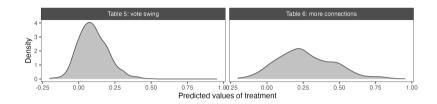
Balance of covariates

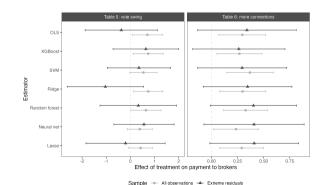


General propensity approach



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Conclusions

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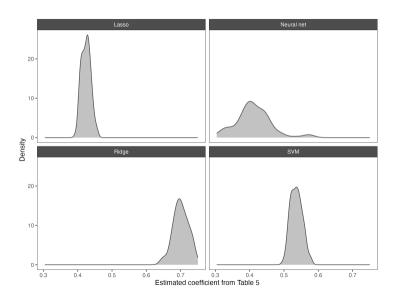
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- Random assignment ensures the propensity score model is correct, eliminating misspecification bias.
- Binary vs. continuous treatments rely on different estimators (AIPW vs. PLR).
- Standard DML assumes i.i.d.; clustering requires adjustments.

Appendix

Repeating estimation 100 times (Brierley and Nathan, 2022)



DML with continuous treatment and i.i.d. SEs

```
_ R Code -
data dml <- data |>
  # turn into data table
  as.data.table() |>
  # create the DML data object
  DoubleMLData$new(y col = outcome,
                   d_{cols} = treatment.
                   x cols = c(covariates, dummies))
# define learner as random forest
learner <- lrn("regr.ranger")</pre>
# obtain two clones of the learner
ml l sim <- learner$clone()</pre>
ml m sim <- learner$clone()</pre>
dml object <- data dml |>
  # specify the learners
  DoubleMLPLR$new(ml l = ml l sim, ml m = ml m sim,
                   # use 5-fold cross-fitting and 20 rounds
                  n_folds=5, n_folds=20)
# fit the model
dml object$fit()
# extract the coefficients and standard errors
coefficients <- tibble(estimate = dml object$coef, se = dml object$se)
```

DML with continuous treatment and clustering

```
R Code -
data dml <- data |>
  # turn into data.table
  as.data.table() |>
 # create the DML data object (note the change in function)
  DoubleMLClusterData$new(y col = outcome,
                           d cols = treatment,
                           x_{cols} = c(covariates, dummies),
                           # kev change
                           cluster cols = clusters)
# define learner as random forest
learner <- lrn("regr.ranger")</pre>
# obtain two clones of the learner
ml l sim <- learner$clone()</pre>
ml m sim <- learner$clone()</pre>
dml object <- data dml |>
  # specify the learners
  DoubleMLPLR$new(ml_l = ml_l_sim, ml_m = ml_m_sim,
                   # use 5-fold cross-fitting and 20 rounds
                   n \text{ folds=5}, n \text{ rep=20}
# fit the model
dml object$fit()
# extract the coefficients and standard errors
coefficients <- tibble(estimate = dml_object$coef, se = dml object$se)</pre>
```

DML Interactive Regression Model and Clustering

```
_____ R Code _____
data dml <- data |>
  # turn into data table
  as.data.table() |>
  # create the DML data object (note the change in function)
  DoubleMLClusterData$new(y col = outcome,
                          d cols = treatment,
                          x cols = c(covariates, dummies).
                          # kev change
                          cluster cols = clusters)
# define learner as random forest
learner <- lrn("classif.ranger")</pre>
# obtain two clones of the learner
ml g = learner$clone() # outcome model
ml m = learner$clone() # treatment model-must be classification learner
dml object <- data dml |>
  # specify the learners
  DoubleMLIRM$new(dml cl obj, ml_g, ml_m, n_folds=5)
# fit the model
dml object$fit()
# extract the coefficients and standard errors
coefficients <- tibble(estimate = dml object$coef, se = dml object$se)
```