How to correctly estimate the effect of advertisement

Introduction of Double Machine Lerning

About me

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- Studying: Econometrics, Causal Inference

Main Theme

How to estimate correctly treatment effects in high-dimensional data(such as advertising data)

Original

- 1. Chernozhukov, Victor, et al. "Double/debiased/neyman machine learning of treatment effects." *American Economic Review* 107.5 (2017): 261-65.
- 2. <u>Belloni, A., Chernozhukov, V., Fernández-Val, I., & Hansen, C. (2017).</u> <u>Program evaluation and causal inference with high-dimensional data.</u> <u>Econometrica, 85(1), 233-298.</u>
- 3. <u>Chernozhukov, Victor, et al. "Double/debiased machine learning for treatment and structural parameters." *The Econometrics Journal* 21.1 (2018): C1-C68.</u>

Problem

- Advertisement data has a lot of variables. (= High-Dimensional Data)
 example: Type of Advertisement , Time , History-data
 - → Under such circumstances, it is known that low-dimensional parameters such as policy effects can not be estimated well by Machine Learning Methods (The model and figure are on the slide later)
- Motivation: How to estimate policy effects correctly?
 - → Use Double Machine Learning(Double ML)!
 - =(FLEXIBLE and SIMPLE Method)
- Double ML Code are on github(https://github.com/VC2015/DMLonGitHub/)
 ...(Double ML package does not exist yet)

Data

Recently

Dataset!

Published

<u>Criteo Uplift Modeling Dataset</u>(EC Site AD Large Dataset for causal inference)

Criteo AL Lab > Dataset > Criteo Uplift Prediction Dataset Criteo Uplift Prediction Dataset By: Criteo Al Lab 31 May 2018 Criteo Uplift Modeling Dataset This dataset is released along with the paper: "A Large Scale Benchmark for Uplift Modeling" Eustache Diemert, Artem Betlei; (Criteo Al Lab), Christophe Renaudin (Criteo), Massih-Reza Amini (LIG, Grenoble INP) This work was published in: AdKDD 2018 Workshop, in conjunction with KDD 2018. When using this dataset, please cite the paper with following bibtex: @inproceedings{Diemert2018, author = {{Diemert Eustache, Betlei Artem} and Renaudin, Christophe and Massih-Reza, Amini}, title={A Large Scale Benchmark for Uplift Modeling}, publisher = {ACM}. booktitle = {Proceedings of the AdKDD and TargetAd Workshop, KDD, London, United Kingdom, August, 20 $year = \{2018\}$

Data

Features are as follows

Fields

Here is a detailed description of the fields (they are comma-separated in the file):

- f0, f1, f2, f3, f4, f5, f6, f7, f8, f9, f10, f11: feature values (dense, float)
- **treatment**: treatment group (1 = treated, 0 = control)
- **conversion**: whether a conversion occured for this user (binary, label)
- visit: whether a visit occured for this user (binary, label)
- **exposure**: treatment effect, whether the user has been effectively exposed (binary)
- Averege Visit Rate and Treatment Ratio is on the right

Key figures

- Format: CSV
- Size: 459MB (compressed)
- Rows: 25,309,483
- Average Visit Rate: .04132
- Average Conversion Rate: .00229
- Treatment Ratio: .846

Model

Partially Linear Model

$$Y = D\theta_0 + g_0(Z) + U, \ E[U|Z,D] = 0$$

Y: Outcome Variable(Whether user visited = visit)

D: Policy Variable (Whether or not an ad has been delivered = treatment)

Z: High-Dimensional vector of covariates(such as history data = f0 f1 ...)

θο: "lift" parameter(Policy effect parameter)

D is expressed by the following equation

$$D = m_0(Z) + V, \ \ E[V|Z] = 0$$

D is conditionally exogenous

Bad ML Estimation

Predict Y using D and Z, then obtain

$$D\hat{ heta}_0 + \hat{g}_0(Z)$$

(Example)

1. Run Random Forest $Y - D\hat{\theta}_0$ on Z and get

$$\hat{g}_0(Z)$$

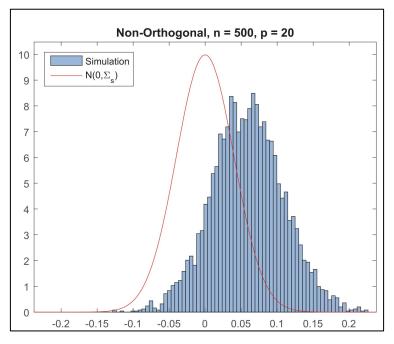
2. $OLS ext{ on } Y - \hat{g}_0(Z) ext{ on } Z ext{ and get}$

$$\hat{\theta}_0$$

3. Repeat 1. and 2. until convergence

Bad ML Estimation

• Prediction performance is Good, but the distribution of $\hat{ heta}_0 - heta_0$ looks like below.



(from paper)

Double ML

1. Predict Y and D using Z by

$$\widehat{E[Y|Z]}$$
 and $\widehat{\mathrm{E[D|Z]}}$

obtained using ML methods(Lasso, Random Forest etc...)

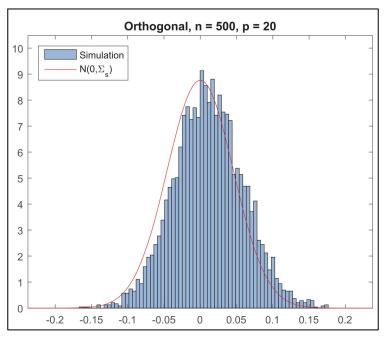
2. Residualize

$$\widehat{W} = Y - \widehat{E[Y|Z]}$$
 and $\widehat{V} = D - \widehat{E[D|Z]}$

3. Regress $\widehat{w}_{\mathrm{on}} \, \widehat{\mathbf{v}}$ and get $\widetilde{\boldsymbol{\theta}}_{0}$

Double ML

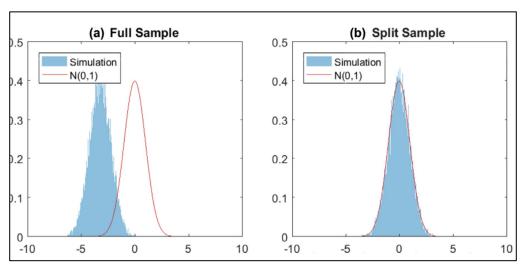
ullet Prediction performance is Good, and the distribution of $ilde{ heta}_0- heta_0$ looks like below.



(from paper)

Sample Splitting

- In Double ML, Sample Splitting is one key Ingredient.
 - → Convergence rate improves with sample splitting(Please refer to the papers for the theoretical content)



(from paper)

Double ML in R

All the code for the simulation is on the github

```
DML2.for.PLM <- function(x, d, y, dreg, yreg, nfold=5) {
 # this implements DML2 algorithm, where there moments are estimated via DML, before constructing
 # the pooled estimate of theta randomly split data into folds
  nobs <- nrow(x)
  foldid <- rep.int(1:nfold,times = ceiling(nobs/nfold))[sample.int(nobs)]</pre>
  I <- split(1:nobs, foldid)</pre>
 # create residualized objects to fill
  ytil <- dtil <- rep(NA, nobs)
  # obtain cross-fitted residuals
  cat("fold: ")
  for(b in 1:length(I)){
    dfit <- dreg(x[-I[[b]],], d[-I[[b]]]) #take a fold out</pre>
    yfit <- yreg(x[-I[[b]],], y[-I[[b]]]) # take a folot out
    dhat <- predict(dfit, x[I[[b]],], type="response") #predict the fold out</pre>
    yhat <- predict(yfit, x[I[[b]],], type="response") #predict the fold out</pre>
    dtil[I[[b]]] <- (d[I[[b]]] - dhat) #record residual</pre>
    ytil[I[[b]]] <- (y[I[[b]]] - yhat) #record residial</pre>
    cat(b," ")
                                         #estimate the main parameter by regressing one residual on the other
  rfit <- lm(ytil ~ dtil)
  coef.est <- coef(rfit)[2]</pre>
                                         #extract coefficient
                                         #record standard error
  se <- sqrt(vcovHC(rfit)[2,2])</pre>
  cat(sprintf("\ncoef (se) = %g (%g)\n", coef.est , se))
  low_ <- coef.est - se*1.96
  upp_ <- coef.est + se*1.96
  return(data.frame(Method = "Double ML", ATE = coef.est, lower_ci = low_, upper_ci = upp_))
```

Since dataset is too large, I made overall sampling first (about 2%)

Introduce Sampling Bias(Since we are using data coming from a randomized

experiment)

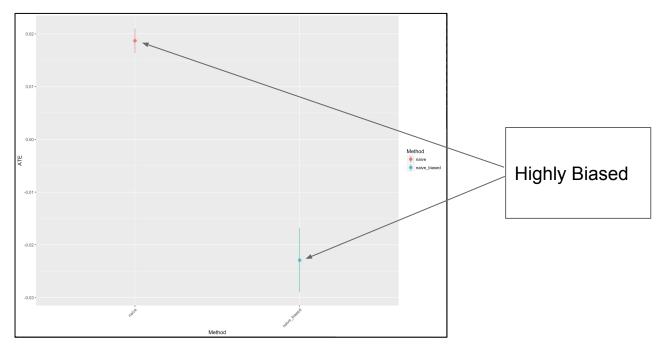
```
#Introduce Sampling bias
pt <- .80 # Drop p% of users who satisfy the following condition
pc <- .95
drop_from_treat <- (df[,"f6"] < 2 | df[,"f0"] > 0.5)
drop_from_control <-(df[,"f6"] > 2 | df[,"f0"] < 0.5)
drop_treat_idx <- which(df[,"W"] == 1 & drop_from_treat)</pre>
drop_control_idx <- which(df[,"W"] == 0 & drop_from_control)</pre>
drop_idx <- unique(c(drop_treat_idx[1:round(pt*length(drop_treat_idx))],</pre>
                      drop_control_idx[1:round(pc*length(drop_control_idx))]))
print(length(drop_idx))
df_mod <- df[-drop_idx,]</pre>
df_mod <- df_mod[sample(nrow(df_mod)),]</pre>
rownames(df_mod) <- NULL
```

Naive ATE Method

 Unbiased estimation is possible with this technique even if there is no sampling bias

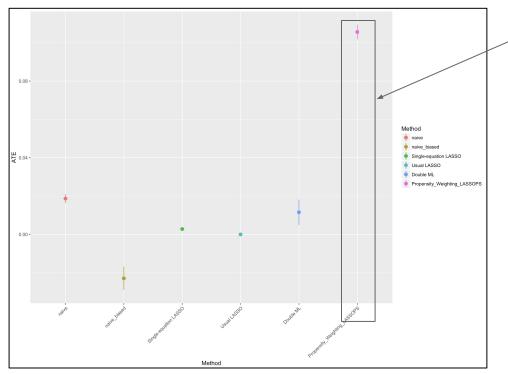
```
mean_df <- dataset %>%
  group_by_(W_var) %>%
 summarise_(y = paste("mean(", outcome_var, ")"),
             y_var = paste("var(", outcome_var, ")"),
             count = "n()") %>%
  mutate(y_var_weight = y_var/(count - 1))
E_y0 = mean_df y[mean_df W == 0]
E_y1 = mean_df y[mean_df W == 1]
tau_hat <- E_y1 - E_y0
```

- Average treatment effect without sampling bias on the left
- Average treatment effect with sampling bias on the right



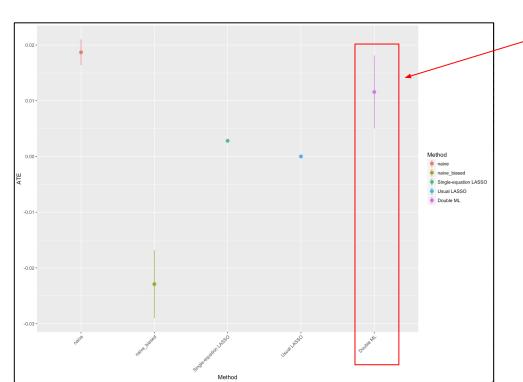
- Use approach: Double ML with Lasso(You can use RandomForest instead of Lasso)
 - → Counter approach: Naive ATE, Single equation Lasso , Usual Lasso , Lasso with Propensity Score Weighting

Result(with Lasso with Propensity Score Weighting)



Lasso with PSW's performance is very bad, so remove this

Result(without Lasso with Propensity Score Weighting)



Double ML's performance is very good!

Conclusion

- Double ML is Flexible and Simple Method for causal inference
- Double ML's performance seems very good with real data