Part 3: Comparing causal model evaluations on the welfare question

Austin Murphy, Fred Lu 6/3/2020

· Models for estimating CATE

```
# Downloading script
# df <- readr::read_csv(file = "https://raw.githubusercontent.com/gsbDBI/ExperimentData/master/Welfare/ProcessedD
        ata/welfarenolabel3.csv", na = character())
# readr::write_csv(df, here::here("data/welfare.csv"))
# Read in provided welfare data
df <- readr::read_csv(file = here::here("data","welfare.csv"), na = character())</pre>
# Read in additional survey data
black assistance <- readr::read_csv(file = here::here("data","black assistance.csv"))</pre>
df <-
 df %>%
  dplyr::left_join(black_assistance)
# Specify outcome, treatment, and covariate variable names to use
outcome_variable_name <- "y" # or y_race for the other outcome variable
treatment variable name <- "w"
covariate_names <- c("partyid", "polviews", "age", "educ", "year", "attblack")</pre>
\verb|all_variable_name| <- \verb|c(outcome_variable_name|, treatment_variable_name|, covariate_names|)|
df <- df[, which(names(df) %in% all variables names)]</pre>
# Rename variables
names(df)[names(df) == outcome_variable_name] <- "Y"</pre>
names(df)[names(df) == treatment_variable_name] <- "W"</pre>
dim(df)
```

```
## [1] 36501 8
```

```
colnames(df)
```

```
## [1] "year" "age" "educ" "partyid" "polviews" "W" "Y"
## [8] "attblack"
```

head(df)

```
## # A tibble: 6 x 8
## year age educ partyid polviews
                               W
                                    Y attblack
## <int> <int> <int> <int> <int> <int> 
                  3
## 1 1986 28 14
                         4 0 0
                                        0.333
                  6
        54 16
44 16
## 2 1986
                           6
                               0
                                    1
                                       0.5
## 3 1986
                    0
                           2
                               0
                                    1
                                       0.75
                                  0
        77 14
                   0
## 4 1986
                                       0.5
## 5 1986
                                       0.5
## 6 1986
         47
            10
                                       0.5
```

```
# Filter out missing data
df <-
    df %>%
    filter_all(all_vars(. != -999)) %>%
    filter(partyid != 7)

# polviews has some values that are 4.1220088 for no reason... remove those
is.wholenumber <-
    function(x, tol = .Machine$double.eps^0.5) abs(x - round(x)) < tol
df <- df[is.wholenumber(df$polviews),]
head(df)</pre>
```

```
## # A tibble: 6 x 8
  year age educ partyid polviews
                              W
                                  Y attblack
## <int> <int> <int> <int>
                       <dbl> <int> <int>
                       4 0 0
                 3
## 1 1986 28 14
                                     0.333
## 2 1986 54 16
## 3 1986 44 16 0
                                 1
                                     0.75
                         2 0
## 4 1986
         77
             14
                   0
                                  0
                                      0.5
## 5 1986
         44
             14
                    0
                              0
                                  0
                                      0.5
            10
## 6 1986 47
                                 0
                   1
                                      0.5
```

Recall that the ATE is -0.33 on this dataset. Earlier we used BART to estimate heterogeneous treatment effects. We now try some comparison methods. (Note that the negative sign is based on the convention of W=1 being treatment. In our report and the original paper the opposite is more interpretable.)

```
mean(df$Y[df$W == 1]) - mean(df$Y[df$W == 0])

## [1] -0.3333626
```

Models for estimating CATE

X-learner

Before fitting the X-learner, first generate, as simple comparisons, model fits with the S-learner and T-learner strategies. We would anticipate that the X-learner would give similar results to BART, both of which would be more effective than S or T learner.

```
testidx <- base::sample(nrow(df), 5000)
df_train <- df[-testidx, ]
df_test <- df[testidx, ]

X <- df_train[,covariate_names]
W <- df_train$\forall W
Y <- df_train$\forall Y
num.trees <- 1000

# s learner
slearn <- regression_forest(df_train[,c(covariate_names,'W')], Y, num.trees=num.trees)

# t learner
n_train <- dim(df_train)[1]
tf0 <- regression_forest(X[W==0,], Y[W==0], num.trees=num.trees)

tf1 <- regression_forest(X[W==1,], Y[W==1], num.trees=num.trees)</pre>
```

Now estimate HTEs via the X-learner strategy, using regression forests for the nuisance components.

```
# Compute the 'imputed treatment effects' using the other group
D1 <- Y[W==1] - predict(tf0, X[W==1,])$predictions
D0 <- predict(tf1, X[W==0,])$predictions - Y[W==0]
# Compute the cross estimators
xf0 <- regression_forest(X[W==0,], D0, num.trees=num.trees)</pre>
xf1 <- regression_forest(X[W==1,], D1, num.trees=num.trees)</pre>
# Predict treatment effects, making sure to always use OOB predictions where appropriate
xf.preds.0 <- rep(0, n_train)</pre>
xf.preds.0[W==0] <- predict(xf0)$predictions
xf.preds.0[W==1] \leftarrow predict(xf0, X[W==1,])$predictions
xf.preds.1 <- rep(0, n train)
xf.preds.1[W==0] <- predict(xf0)$predictions</pre>
xf.preds.1[W==1] \leftarrow predict(xf0, X[W==1,])$predictions
# Estimate the propensity score
propf <- regression forest(X, W, num.trees=num.trees)</pre>
ehat <- predict(propf)$predictions</pre>
# Finally, compute the X-learner prediction
tauhat_xl <- (1 - ehat) * xf.preds.1 + ehat * xf.preds.0</pre>
```

Causal Forest

Train a causal forest and compute (out-of-bag) CATE estimates on the training set.

```
cf <- causal_forest(
    X = df_train[, covariate_names],
    Y = df_train$Y,
    W = df_train$W,
    num.trees=1000
)

oob_pred <- predict(cf, estimate.variance=TRUE)</pre>
```

BART model

Fit the BART model again, to enable test set comparisons with the other models

```
## ****Into main of pbart
## ****Data:
## data:n,p,np: 14738, 7, 10000
## y1,yn: 0, 0
## x1,x[n*p]: 1986.000000, 0.500000
## xp1,xp[np*p]: 2002.000000, 0.250000
## *****Number of Trees: 50
## *****Number of Cut Points: 14 ... 6
## *****burn and ndpost: 1000, 1000
## ****Prior:mybeta,alpha,tau: 2.000000,0.950000,0.212132
## ****binaryOffset: -0.626675
## ****Dirichlet:sparse,theta,omega,a,b,rho,augment: 0,0,1,0.5,1,7,0
## ****nkeeptrain,nkeeptest,nkeeptreedraws: 1000,1000,1000
## ****printevery: 100
## ****skiptr,skipte,skiptreedraws: 1,1,1
##
## MCMC
## done 0 (out of 2000)
## done 100 (out of 2000)
## done 200 (out of 2000)
## done 300 (out of 2000)
## done 400 (out of 2000)
## done 500 (out of 2000)
## done 600 (out of 2000)
## done 700 (out of 2000)
## done 800 (out of 2000)
## done 900 (out of 2000)
## done 1000 (out of 2000)
## done 1100 (out of 2000)
## done 1200 (out of 2000)
## done 1300 (out of 2000)
## done 1400 (out of 2000)
## done 1500 (out of 2000)
## done 1600 (out of 2000)
## done 1700 (out of 2000)
## done 1800 (out of 2000)
## done 1900 (out of 2000)
## time: 98s
## check counts
## trcnt,tecnt: 1000,1000
```

```
# Process BART predictions to get in same format (treatment effect for each entry)
pred_w0 = bart$prob.test.mean[1:5000]
pred_w1 = bart$prob.test.mean[5001:10000]
tauhat_bart_test <- pred_w1 - pred_w0</pre>
```

```
X_test <- df_test[,covariate_names]</pre>
tauhat s test <- predict(slearn, df test[,c(covariate names,'W')])</pre>
tauhat_t_test <- numeric(nrow(df_test))</pre>
\label{tauhat_test} {\tt tauhat_t_test[which(df_test$W==1)] <- predict(tf1, X_test[df_test$W==1, ])} {\tt spredictions}
\label{tauhat_t_test} {\tt tauhat_t_test[which(df_test\$W==0)] <-\ predict(tf1,\ X\_test[df_test\$W==0,\ ])} \ {\tt spredictions}
X_test <- df_test[,covariate_names]</pre>
ehat.test <- predict(propf, X_test)$predictions</pre>
xf.preds.1.test <- predict(xf1, X_test)$predictions</pre>
xf.preds.0.test <- predict(xf0, X_test)$predictions</pre>
tauhat_xl_test <- (1 - ehat.test) * xf.preds.1.test + ehat.test * xf.preds.0.test</pre>
test_pred <- predict(cf, as.matrix(df_test[,covariate_names]), estimate.variance=TRUE)</pre>
tauhat_cf_test <- test_pred$predictions</pre>
# Compute Y-star
p <- mean(df_test$W)</pre>
Y_star \leftarrow ((df_test$W - p)/(p*(1-p)))*df_test$Y
{\it \# Compute the sample average treatment effect to use as a baseline comparison}
tauhat_sample_ate <- with(df_train, mean(Y[W==1]) - mean(Y[W==0]))</pre>
\# Compute test mse for all methods
mse <- data.frame(</pre>
  Sample_ATE_Loss = (Y_star - tauhat_sample_ate)^2,
  BART Loss = (Y star - tauhat bart test)^2,
  Causal_Forest_Loss = (Y_star - tauhat_cf_test)^2,
  X_Learner_Loss = (Y_star - tauhat_xl_test)^2,
  S_Learner_Loss = (Y_star - tauhat_s_test$predictions)^2,
T_Learner_Loss = (Y_star - tauhat_t_test)^2)
print('Mean')
```

```
## [1] "Mean"
```

colMeans(mse)

```
## Sample_ATE_Loss BART_Loss Causal_Forest_Loss X_Learner_Loss
## 1.0007973 0.9887360 0.9909984 0.9903493
## S_Learner_Loss T_Learner_Loss
## 1.6265117 1.2076632
```

```
print('SD')
```

```
## [1] "SD"
```

```
apply(mse, 2, sd)
```

```
## Sample_ATE_Loss BART_Loss Causal_Forest_Loss X_Learner_Loss
## 1.536451 1.547347 1.544625 1.545251
## S_Learner_Loss T_Learner_Loss
## 2.602909 1.956386
```

As expected, BART, causal forest, and X-learner outperform S learner and T learner. Note that BART, causal forest, and X-learner perform very similarly.

```
Y.forest.test = regression_forest(X = as.matrix(df_test[covariate_names]), Y = df_test$Y)
Y.hat.test = predict(Y.forest.test)$predictions
W.forest.test = regression_forest(X = as.matrix(df_test[covariate_names]), Y = df_test$W)
W.hat.test = predict(W.forest.test)$predictions

mse_rloss <- data.frame(
    Sample_ATE_Loss = (df_test$Y - Y.hat.test - (df_test$W - W.hat.test) * tauhat_sample_ate)^2,
    BART_Loss = (df_test$Y - Y.hat.test - (df_test$W - W.hat.test) * tauhat_bart_test)^2,
    Causal_Forest_Loss = (df_test$Y - Y.hat.test - (df_test$W - W.hat.test) * tauhat_of_test)^2,
    X_Learner_Loss = (df_test$Y - Y.hat.test - (df_test$W - W.hat.test) * tauhat_xl_test)^2,
    S_Learner_Loss = (df_test$Y - Y.hat.test - (df_test$W - W.hat.test) * tauhat_s_test$predictions)^2,
    T_Learner_Loss = (df_test$Y - Y.hat.test - (df_test$W - W.hat.test) * tauhat_xl_test)^2
)
print('Mean')</pre>
```

```
## [1] "Mean"
```

```
colMeans(mse_rloss)
```

```
## Sample_ATE_Loss BART_Loss Causal_Forest_Loss X_Learner_Loss
## 0.1611311 0.1588524 0.1591312 0.1589986

## S_Learner_Loss T_Learner_Loss
## 0.2652901 0.1589986
```

```
print('SD')
```

```
## [1] "SD"
```

```
apply(mse_rloss, 2, sd)
```

```
## Sample_ATE_Loss BART_Loss Causal_Forest_Loss X_Learner_Loss
## 0.1873270 0.1973481 0.1947029 0.1955534
## S_Learner_Loss T_Learner_Loss
## 0.3441243 0.1955534
```

The same trend is observed using R-loss as the performance evaluation.