

University of Essex

CE-888-7-SP DATA SCIENCE AND DECISION MAKING

CAUSAL INFERENCE ASSIGNMENT-21a3

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https://github.com/krnbhtt/CE-888

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ABSTRACT

For decades, the term causal inference has been a critical research topic in a variety of domains such as computer science, statistics, economics, and public policy. Furthermore, analysing the causal influence from observational data has recently become an interesting topic of study due to the abundance of available data as well as the low budget needs, when compared to randomised controlled trials. In addition, as a result of the rapidly evolving field of machine learning, various estimation methodologies of causal effects for observational data have formed. Furthermore, this research provides an in-depth examination of the "causal inference" technique below the possible output structure. One of them is the framework of causal inference. Furthermore, the processes are divided into two fundamental types based on whether if they need all three assumptions of the corresponding output framework or not. Furthermore, for each categorization, both the current process of enhanced machine learning and the techniques of classical statistics are discussed and compared. The linear regression model is picked in the regression model and its feature relevance is described. Initially, the commonly used benchmark sets of data and open-source scripts are summarised, which generally open the way for researchers to evaluate and use causal inference methodologies. This research will explain the whole training process, from uploading data to preopening it to train by using various models that predict the outcomes of various scenarios.

INTRODUCTION

Currently, causality, Language and correlation are largely interchangeable, besides they have wide range of explanation. Furthermore, the term correlation denotes a shared link: if the variables show a rising or declining trend, they are connected. Furthermore, the term causality is used to refer to both the cause and the impact, where the cause is somehow essential for the impact, and the impact is somehow intertwined with the cause. Furthermore, causal inference is the way of formulating a conclusion about the "causal relationship" based on the conditions of occurrence of a result. The difference between the correlation inference and the causal inference is that the computations of the reactions of the outcome variable, even when the consequence is modified. Furthermore, the investigator collects observational data by essentially observing the process without any form of intervention. The investigators have no authority over individuals or treatments, and they just monitor the suggested and gathered data based on their observations. However, using observational data, it is simple to deduce the actions, information, and results of what has occurred. Moreover, the concept causal inference refers to the calculation of a causal effect. Also, the problem is whether just monitored outputs can be used to derive the implications of the proposed system's intervention. This is also known as the fundamental issue of causal inference. Also, the primary goal of this report is to identify the fundamental causal inference data sets, as well as the dataset's features and essential performance metrics. Moreover, the whole report is built on developing a prediction machine learning prediction system to predict whether child is healthy / under treatment, family got support or not (IHDP), whether person have got job training or not. This report is remarkable because, by using this, it would be easy to predict whether a family got a treatment/support or not/they are under control (as per IHDP dataset), or the person is currently employed / did he get a training from NSWP or not, and is he employed or not (as per job data). This report will also discuss how to become acquainted with, in addition to well-established types of causal inference estimation methods, as well as a thorough explanation of the important properties. More than that, this report will help to acquire an instinct for how causal inference relates to machine learning problems. Furthermore, in this assignment, I will load, clean, and investigate the data sets which will be discussed in the data section. In this assignment I have used Liner Regressor and Random Forest Regressor models and explained feature importance, why these models are chosen.

Data:

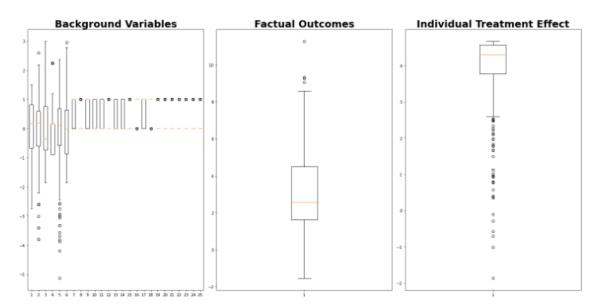
The linear regression model is mostly used to predict the value of a variable based on other factors. While pre-processing the data I found that There are no null values or zero values in both IHDP and job datasets.

```
<class 'pandas.core.frame.DataFrame'>
RangeIndex: 3212 entries, 0 to 3211
Data columns (total 17 columns):
     Column Non-Null Count Dtype
                 3212 non-null
                 3212 non-null
                                      float64
                 3212 non-null
                 3212 non-null
                                      float64
                 3212 non-null
                                      float64
                 3212 non-null
                                      float64
                                      float64
                 3212 non-null
                 3212 non-null
      8
                 3212 non-null
                                      float64
                 3212 non-null
 10
     10
                 3212 non-null
                                      float64
     11
                 3212 non-null
 12
      12
                 3212 non-null
                                      float64
                 3212 non-null
     14
15
 14
                 3212 non-null
                                      float64
 15
                 3212 non-null
                                      float64
 16 16
                 3212 non-null
16 16 3212 non-null float64
dtypes: float64(17)
memory usage: 426.7 KB
<class 'pandas.core.frame.DataFrame'>
RangeIndex: 3212 entries, 0 to 3211
Data columns (total 1 columns):
 # Column Non-Null Count Dtype
                 3212 non-null uint8
dtypes: uint8(1)
memory usage: 3.3 KB

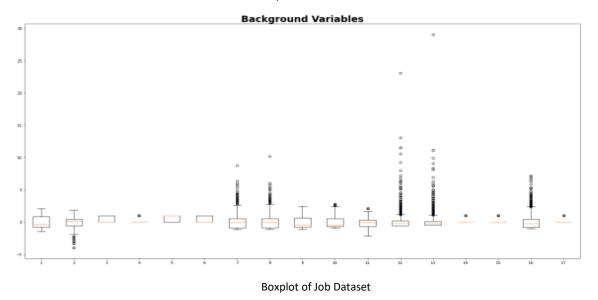
<class 'pandas.core.frame.DataFrame'>

RangeIndex: 3212 entries, 0 to 3211
Data columns (total 1 columns):
# Column Non-Null Count Dtype
                 3212 non-null uint8
 0 0
dtypes: uint8(1)
memory usage: 3.3 KB <class 'pandas.core.frame.DataFrame'>
RangeIndex: 3212 entries, 0 to 3211
Data columns (total 1 columns):
# Column Non-Null Count Dtype
                 3212 non-null uint8
```

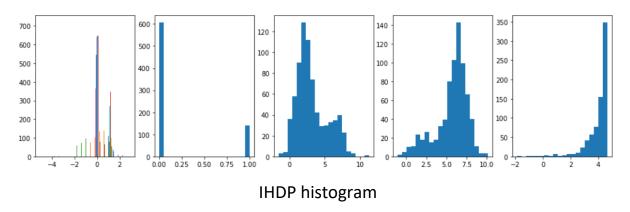
therefore, no pre-processing is needed in regards to encoding and filling Nan rows.



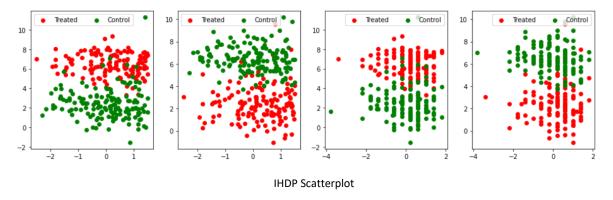
Boxplot of IHDP dataset



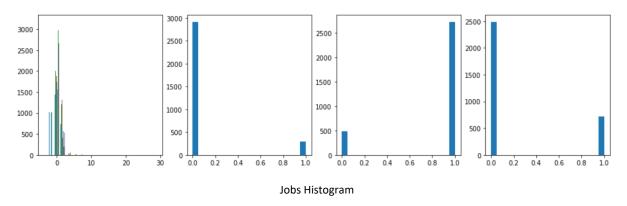
But There does appear to be quite a few outliers in the IHDP and job dataset (as seen by the values lying outside the minimum and maximum of each box and whisker plot), especially in the non-binary variables. But both datasets vast number of outliers in the background variables, thus a similar approach of experimenting with standardization will need to be taken. We could also investigate random forest regression models that should internally deal with any outliers and reduce the chance of our results being skewed or bias in any way.



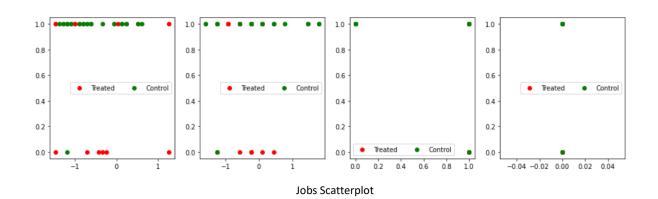
The background variables X, although unbalanced, seem to demonstrate a typical shape given the dataset size, the same goes for both the factual and counter factual outcomes. The treatment variable is clearly represented as a binary, 0 being no treatment and 1 being treatment given. The individual treatment effect seems to be extremely negatively skewed although most of the values are around 4 which seems correct given our average treatment effect of 4.02



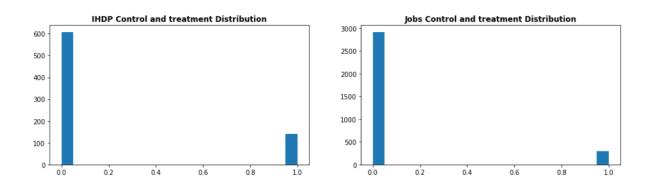
Looking at the above graphs of the first two background variables (factual and counterfactual) measured against their outcomes with both the treated and control groups, we can observe some clear effects of the treatment which hopefully means they can be measured as we train and test our regression models.



Similarly, to IHDP, the background variables in Jobs seem to be relatively unbalanced, however again demonstrate the typical shape for the relative dataset size. All of t, y, and e represent binary values of either 0 or 1 with all graphs being unbalanced in their distributions. What I mean by this is there is a lot more of one result then there is the other (More 1's than 0's or vice versa)



As opposed to IHDP jobs outcomes is recorded as a binary variable thus only plotting scatter points on 0 and 1. It is significantly harder to spot obvious effects given the 4 graphs above, however this could just be a coincidence given the background variables we have chosen to plot.



The above graphs confirm our need to use X-learner. There is clear imbalance towards the treatment and control groups in both datasets, hopefully X-learner should be able to deal with this when calculating our CATE value.

Splitting the Data

I'm looking here to split the datasets with the same ratios of 80/20, where 80% of data will be used as train data and 20% data would be the test data and then create a function that given the parameters that are used to call it, return standardized data.

IHDP_x_train, IHDP_x_test, IHDP_t_train, IHDP_t_test, IHDP_yf_train, IHDP_yf_test, IHDP_ite_train, IHDP_ite_test = train_test_split(IHDP_x, IHDP_t, IHDP_yf, IHDP_ite, test_size=0.2)

Jobs_x_train, Jobs_x_test, Jobs_t_train, Jobs_t_test, Jobs_y_train, Jobs_y_test,
Jobs_e_train, Jobs_e_test = train_test_split(Jobs_x, Jobs_t, Jobs_y, Jobs_e,
test_size=0.2)

Now we can start data modelling and training.

Methodology:

To solve this causal inference problem, I have used linear regression and random forest regressor. The linear regression model is mostly used to predict the value of a variable based on other variables. Moreover, utilising the methods of cross-validation, dimensionality reduction, and regularisation, linear regression may effectively handle overfitting. And the random forest regressor helps to internally deal with any outliers and reduce the chance of our results being skewed or bias in any way. However, for this case Linear regression and random forest regression both are the regressor tools.

```
In [26]: ▶ 1 # Simple Random forest trained using the IHDP features + treatment
                  2 random_forest_IHDP = RandomForestRegressor()
3 Concatenated_XT_train_IHDP = np.concatenate([IHDP_x_train, IHDP_t_train], axis=1)
                  4 random_forest_IHDP.fit(Concatenated_XT_train_IHDP, IHDP_yf_train.flatten())
                  6 # Simple Random forest trained using the Jobs features + treatment
                 7 random_forest_Jobs = RandomForestRegressor()
8 Concatenated_XT_train_Jobs = np.concatenate([Jobs_x_train, Jobs_t_train], axis=1)
9 random_forest_Jobs.fit(Concatenated_XT_train_Jobs, Jobs_y_train.flatten())
    Out[26]: RandomForestRegressor()
           Now we predict Y0 and Y1 given t=0 and t=1 respectively for both datasets
                 1 # t=0 - IHDP
                     Concatenated_XT_zeros_IHDP = np.concatenate([IHDP_x_test, np.zeros_like(IHDP_t_test)], axis=1)
                  3 Y0_IHDP = random_forest_IHDP.predict(Concatenated_XT_zeros_IHDP)
4 # t=1 - IHDP
                  5  Concatenated_XT_ones_IHDP = np.concatenate([IHDP_x_test, np.ones_like(IHDP_t_test)], axis=1)
6  Y1  IHDP = random forest IHDP.predict(Concatenated XT ones IHDP)
                  9 # t=0 - Jobs
                 10 Concatenated_XT_zeros_Jobs = np.concatenate([Jobs_x_test, np.zeros_like(Jobs_t_test)], axis=1)
                 13 Concatenated_XT_ones_Jobs = np.concatenate([Jobs_x_test, np.ones_like(Jobs_t_test)], axis=1)
14 Y1_Jobs = random_forest_Jobs.predict(Concatenated_XT_ones_Jobs)
           Now we can calculate our effect estimates as Y1-Y0
In [28]: N
                 1 #ITEs for IHDP dataset
2 Effect_Estimates_IHDP :
                     Effect_Estimates_IHDP = Y1_IHDP - Y0_IHDP
                     #ITEs for Jobs dataset
                  4 Effect_Estimates_Jobs = Y1_Jobs - Y0_Jobs
```

Simple Random Forest regression Model Building and Training

```
In [29]: H 1 # Functions Taken from Lab 4 def pehe(effect_true, effect_pred):
                                                                                                                                Precision in Estimating the Meterogeneous Treatment Effect (PEHE) 
:param effect_true: true treatment effect value 
:param effect_pred: predicted treatment effect value 
:return: PEHE
                                                                                                                                 return np.abs(np.mean(effect_pred) - np.mean(effect_true))
                                                                                     def abs_ate(effect_true, effect_pred):

Absolute error for the Average Trea

param effect_true: true treatment

param effect_pred: predicted treat

return: absolute error on ATE

return np.sqrt(np.mean((effect_true
                                                                                                                           Absolute error for the Average Treatment Effect (ATE) 
:param effect_true: true treatment effect value 
:param effect_pred: predicted treatment effect value 
:return: absolute error on ATE
                                                                                                                                    return np.sqrt(np.mean((effect_true - effect_pred)**2))
                                                                                     1 # Function taken from D. Machlanski (Unit Supervisor) - https://github.com/dmachlanski def abs_att(effect_pred, yf, t, e):
In [30]: 📕
                                                                                                                                   Absolute error for the Average Treatment Effect on the Treated :param effect_pred: predicted treatment effect value :param yf: factual (observed) outcome :param : treatment status (created/control) :param e: whether belongs to the experimental group :param : shoulte error on ATT
                                                                                 Absolute error to the works and the series of the series o
                                                                                                                                    \begin{array}{lll} \text{att\_true} &=& \text{np.mean}(yf[t > \emptyset]) &-& \text{np.mean}(yf[(1 - t + e) > 1]) \\ \text{att\_pred} &=& \text{np.mean}(effect\_pred[(t + e) > 1]) \\ \end{array} 
                                                                                                                              Computes the risk of the policy defined by predicted effect 
param effect pred: predicted treatment effect value 
param t: treatment status (treated/control) 
param t: whether belongs to the experimental group 
prediction of the property 
                                                                                                                                   If Consider only the cases for which we have experimental data (i.e., e > \theta) t_e = t[e > \theta] yf_e = yf[e > \theta] effect_pred_e = effect_pred[e > \theta]
                                                                                                                                   policy = effect_pred_e > 0.0
treat_overlap = (policy == t_e) * (t_e > 0)
control_overlap = (policy == t_e) * (t_e < 1)
                                                                                                                                    else:
treat_value = np.mean(yf_e[treat_overlap])
                                                                                                                                 if np.sum(control_overlap) == 0:
    control_value = 0
else:
    control_value = np.mean(yf_e[control_overlap])
                                                                                                                                      pit = np.mean(policy)
policy_value = pit * treat_value + (1.0 - pit) * control_value
```

Performance evaluation of random forest regressor model

Hyper parameter Tuning:

Here we have used random search parameters, using 10-fold cross validation.

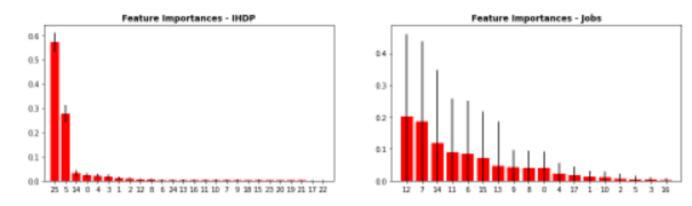
Grid Search IHDP

Using the results, we can construct our parameter grid for grid search with values more in the ballpark of what we're looking for

```
[37]: M 1 Benchmark_IHDP = RandomForestRegressor()
                     # Our parameter Grid uses values centered around the parameters found from the random search
                 10 }
                GridSearch_RF_IHDP = GridSearchCV(estimator = Benchmark_IHDP, param_grid = param_grid, cv = 10, n_jobs = -1, verbose GridSearch_RF_IHDP.fit(Concatenated_XT_train_IHDP, IHDP_yf_train.flatten())

GridSearch_RF_IHDP.best_params_
               Fitting 10 folds for each of 162 candidates, totalling 1620 fits
   Out[37]: {'bootstrap': True, 'max_depth': 8,
                 'max_features': 'auto',
                 'min_samples_leaf': 3,
'min_samples_split': 4,
                'n_estimators': 600}
          Grid Search Jobs
[38]: N 1 Benchmark Jobs = RandomForestRegressor()
2 # Our parameter Grid uses values centered around the parameters found from the random search
                   # Our parameter Grid uses values center
param_grid = {
    'bootstrap': [True],
    'max_depth': [None, 2, 4],
    'min_samples_leaf': [3, 4, 5],
    'min_samples_split': [2, 4, 6],
    'n_estimators': [1800, 2000, 2200]
                13  GridSearch RF Jobs.fit(Concatenated_XT_train_Jobs, Jobs_y_train.flatten())
14  GridSearch_RF_Jobs.best_params_
               Fitting 10 folds for each of 162 candidates, totalling 1620 fits
  Out[38]: {'bootstrap': True,
'max_depth': None,
                 'max features':
                 'min_samples_leaf': 5,
'min_samples_split': 2
'n_estimators': 1800}
```

Now I am performing feature importance using random forest regression method and plot a graph. Here is the graph of importance features of both IHDP and Job datasets.

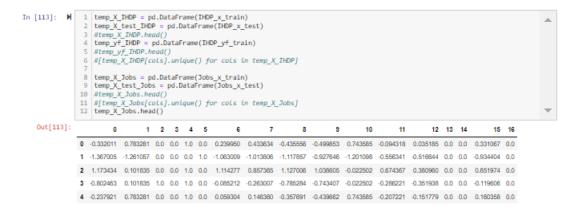


In IHDP dataset we can clearly see that column 25 is a most important feature. Column 25 is our concatenated treatment data. In this dataset feature importance is extremely skewed between column no. 25 & 5

In Jobs dataset feature importance are less skewed than IHDP dataset. We can clearly see that between column 12 & 7. Moreover, we can see that our treatment variable is placed on column 12.

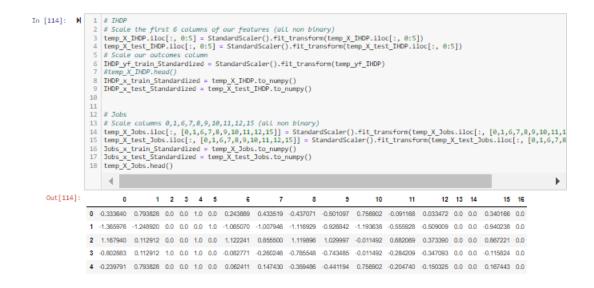
Linear Regressor:

In this section we will start with the simple linear regressor model like random forest regressor. Here every binary feature will be standardized.



In IHDP dataset we can see that column 0-5 require standard scaling. From past model we know that our outcome column requires it.

In jobs dataset there are many columns which requires standard scaling rest are binary, so they do not require it.



Now let's start with building the simple linear regressor model.

So now our effect estimates are built so we can continue with metric evaluation.

```
In [125]: M 1 # Metrics for IHDP dataset
2 results = []
3 results.append(['RF', ATE_IHDP, PEHE_IHDP])
                          results.append(['Optimized RF', ATE_IHDP_Optimized, PEHE_IHDP_Optimized])
results.append(['Is', ATE_IHDP_IR, PEHE_IHDP_LR])
cols = ['Method', 'ATE_IND_IR, PEHE_IHDP_LR])
df_First = pd.DataFrame(results, columns=cols)
df_First.
                       8 df First
     Out[125]:
                               Method ATE test PEHE test
                     0
                                RF 1.305712 0.116149
                      1 Optimized RF 1.254617 0.137322
                     2 LR 2.401287 2.263552
In [126]: W 1 # Metrics for Jobs Dataset
2    results_ = []
3    results_append(['RF', ATT_Jobs, Rpol])
4    results_append(['Optimized RF', ATT_Jobs_Optimized, Rpol_Optimized])
5    results_append(['LR', ATT_Jobs_LR, Rpol_LR])
6    cols_ = ['Method', 'ATT_test', 'Risk Policy']
7    df_Second = pd.DataFrame(results_, columns=cols_)
8    df_Second
                       8 df_Second
     Out[126]:
                             Method ATT test Risk Policy
                      0 RF 0.063641 0.202296
                      1 Optimized RF 0.033940 0.217834
                     2 LR 0.006004 0.224490
```

For hyperparameter tuning, we can execute a grid search using the selection of parameters available to us to find the best performing linear regression model

So, our grid search has been executed so let's move towards performance evaluation side combined with 10-fold cross validation.

```
Concatenated_XT_zeros_IHDP = np.concatenate([IHDP_x_test_Standardized, np.zeros_like(IHDP_t_test)], axis=1)
                          Y0_IHDP = LR_IHDP_Optimized.predict(Concatenated_XT_zeros_IHDP)
# t=1 - IHDP
                       11 LR_Jobs_Optimized = LinearRegression(fit_intercept=True, positive=False)
12 LR_Jobs_Optimized.fit(Concatenated_XT_train_Jobs, Jobs_y_train.flatten())
                     14 # t=9 - Jobs
15 Concatenated_XT_zeros_lobs = np.concatenate([Jobs_x_test_Standardized, np.zeros_like(Jobs_t_test)], axis=1)
16 Y0_Jobs = LR_Jobs_Optimized.predict(Concatenated_XT_zeros_Jobs)
17 # t=1 - Jobs
                     18 Concatenated_XT_ones_Jobs = np.concatenate([Jobs_x_test_Standardized, np.ones_like(Jobs_t_test)], axis=1)
19 Y1_Jobs = LR_Jobs_Optimized.predict(Concatenated_XT_ones_Jobs)
                      21 WITEs for IHDP dataset
                     In [131]: M    1 ATE_IHDP_LR_Optimized = abs_ate(IHDP_ite_test, Effect_Estimates_IHDP_LR_Optimized)
                          PEME_IMOP_LR_Optimized = pehe(IMOP_ite_test, Effect_Estimates_IMOP_LR_Optimized)
ATT_Jobs_LR_Optimized = abs_att(Effect_Estimates_Jobs_LR_Optimized, Jobs_y_test.flatten(), Jobs_t_test.flatten(), Jobs_e
Rpol_LR_Optimized = policy_risk(Effect_Estimates_Jobs_LR_Optimized, Jobs_y_test.flatten(), Jobs_t_test.flatten(), Jobs_e
                                                                                                                                                                                                    Þ
In [132]: 🔰 1 # Metrics for IHDP dataset
                          results = []
results.append(['RF', ATE_IHDP, PEHE_IHDP])
                      results.append(['RF', ATE_INDP, PEHE_INDP])

results.append(['Optimized R', ATE_INDP_Optimized, PEHE_INDP_Optimized])

results.append(['Optimized R', ATE_INDP_LR, PEHE_INDP_LR])

results.append(['Optimized LR', ATE_INDP_LR_Optimized, PEHE_INDP_LR_Optimized])

cols = ['Method', 'ATE test', 'PEHE test']

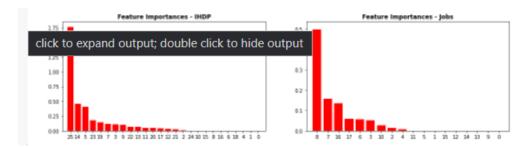
df_First = pd.DataFrame(results, columns=cols)

df_First
     Out[132]:
                              Method ATE test PEHE test
                     0
                          RF 1.305712 0.116149
                     1 Optimized RF 1.254617 0.137322
                    2 LR 2.401287 2.263552
                     3 Optimized LR 2.410418 2.273235
In [134]: M 1 # Metrics for Jobs Dataset
                          results_ = []
results_.append(['RF', ATT_Jobs, Rpol])
                          results_.append([NF, AII_JOBS, MPDI])
results_.append([Optimized RF, ATI_JOBS_Optimized, Rpol_Optimized])
results_.append(['LR', ATI_JOBS_LR, Rpol_LR])
results_.append(['Optimized LR', ATI_JOBS_LR Optimized, Rpol_LR_Optimized])
cols_ = ['Method', 'ATT test', 'Risk Policy']
df_Second = pd.DataFrame(results_, columns=cols_)
df_Second
     Out[134]:
                              Method ATT test Risk Policy
                     0 RF 0.063641 0.202296
                    2 LR 0.006004 0.224490
                     3 Optimized LR 0.006004 0.224490
```

Now finally we can plot and visualise the feature importance of the data, from more useful to less useful. we can recognize feature by their column indices.

```
10 LR_Jobs_Optimized.fit(Concatenated_XT_train_Jobs, Jobs_y_train.flatten())
                                                        | print("Feature Importances - Jobs") | print("Feature Importances - Jobs") | for f in range(Concatenated_XT_train_Jobs.shape[1]): | print("%d. %s (%f)" % (f + 1, "Feature Column: "+ str(indices_Jobs[f]), importances_Jobs[indices_Jobs[f]]))
                                                    Feature Importances - IHDP
1. Feature Column: 25 (1.760401)
2. Feature Column: 14 (0.459039)
3. Feature Column: 5 (0.407445)
4. Feature Column: 23 (0.177242)
                                                    4. Feature Column: 23 (6.177242)
5. Feature Column: 19 (6.141741)
6. Feature Column: 7 (8.118741)
7. Feature Column: 3 (8.118880)
8. Feature Column: 9 (8.181924)
9. Feature Column: 22 (8.066579)
10. Feature Column: 13 (8.063097)
11. Feature Column: 11 (8.043673)
12. Feature Column: 28 (8.046622)
                                                     13. Feature Column: 17 (0.043350)
14. Feature Column: 12 (0.032215)
15. Feature Column: 21 (0.027382)
                                                    15. Feature Column: 21 (0.027382)
16. Feature Column: 2 (0.088747)
17. Feature Column: 24 (0.000000)
18. Feature Column: 15 (0.000000)
19. Feature Column: 15 (0.000000)
20. Feature Column: 8 (0.000000)
21. Feature Column: 6 (0.000000)
22. Feature Column: 6 (0.000000)
23. Feature Column: 18 (0.000000)
                                                     24. Feature Column: 4 (0.000000)
25. Feature Column: 1 (0.000000)
26. Feature Column: 0 (0.000000)
                                                   Feature Importances - Jobs
1. Feature Column: 8 (0.496983)
2. Feature Column: 7 (0.156823)
3. Feature Column: 17 (0.85839)
5. Feature Column: 17 (0.858395)
5. Feature Column: 6 (0.085442)
6. Feature Column: 3 (0.081712)
7. Feature Column: 10 (0.826582)
8. Feature Column: 2 (0.034205)
9. Feature Column: 1 (0.096089)
10. Feature Column: 11 (-0.096089)
11. Feature Column: 5 (-0.086770)
12. Feature Column: 15 (-0.082146)
13. Feature Column: 15 (-0.082146)
14. Feature Column: 12 (-0.082981)
14. Feature Column: 12 (-0.082981)
14. Feature Column: 12 (-0.089981)
                                                    13. Feature Column: 12 (-0.0902140)
14. Feature Column: 12 (-0.09991)
15. Feature Column: 14 (-0.15553)
16. Feature Column: 13 (-0.155559)
17. Feature Column: 9 (-0.267712)
18. Feature Column: 0 (-0.284215)
      In [136]: N

1 fig, axs = plt.subplots(1, 2, figsize=(16, 4))
2 axs[0].bar(range(Concatenated_XT_train_IHDP.shape[1]), importances_IHDP[indices_IHDP], color="r", align="center")
3 axs[0].set_xticks(range(Concatenated_XT_train_IHDP.shape[1]), indices_IHDP)
4 axs[0].set_ylim([0, None])
5 axs[0].set_ylim([0, None])
6 axs[0].set_ylim([0, None])
7 axs[1].bar(range(Concatenated_XT_train_Jobs.shape[1]), importances_Jobs[indices_Jobs], color="r", align="center")
8 axs[1].set_xticks(range(Concatenated_XT_train_Jobs.shape[1]), indices_Jobs)
9 axs[1].set_xlim([-1, Concatenated_XT_train_Jobs.shape[1])]
10 axs[1].set_ylim([0, None])
11 axs[1].set_xlim([-1, Concatenated_XT_train_Jobs.shape[1])]
12 plt.show()
```



In IHDP dataset our feature importance follow similar pattern with the results being extremely skewed and our most important features being, our treatment column, background feature 14, and background feature 5 same as random forest regressor.

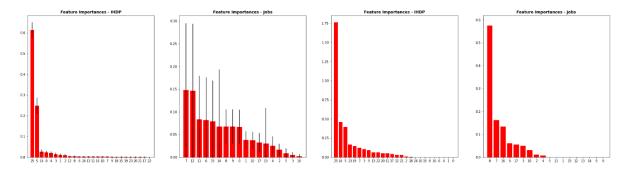
In Jobs dataset The skewness of the feature importance has completely changes after using linear regressions, we now have unbalanced importance, with our previously most importance features being much further down the list, we seemingly have increased importance in regards to the treatment feature however

For propensity Score Re-weighting we are going to use a random forest classifier. First we need to train a classifier to predict propensity scores based on background features. We also need a function that calculates sample weights

based on these propensity scores. Given equation is as follow: $w_i = \frac{l_i}{e(x_i)} + \frac{1-l_i}{1-e(x_i)}$

Now we have sample weights so we can train our regressors using the weights.

Now we will repeat all Random Search, Grid Search, Metric Evaluation, Feature importance (IPSW) and then we will visualise the data again.



Advanced CATE Estimators

```
xl_IHDP.fit(IHDP_yf_train, IHDP_t_train.flatten(), X=IHDP_x_train) xl_Jobs.fit(Jobs_y_train, Jobs_t_train.flatten(), X=Jobs_x_train)
                                 7 xl_te_test_IHDP = xl_IHDP.effect(IHDP_x_test)
8 xl_te_test_Jobs = xl_Jobs.effect(Jobs_x_test)
 In [86]: # # Metrics for IHDP dataset
    results.append(['RF', ATE_IHDP, PEHE_IHDP])
    results.append(['Optimized Rr', ATE_IHDP_Optimized, PEHE_IHDP_Optimized])
    results.append(['LR', ATE_IHDP_LR, PEHE_IHDP_LR])
    results.append(['Optimized LR', ATE_IHDP_LR_Optimized, PEHE_IHDP_LR_Optimized])
    results.append(['Optimized LR', ATE_IHDP_LR_IPSW, PEHE_IHDP_RR_IPSW])
    results.append(['IPSW LR', ATE_IHDP_LR_IPSW, PEHE_IHDP_LR_IPSW])
    results.append(['XL', XL] ate_test , XL_pehe_test])
    cols = ['Method', 'ATE_test', 'PEHE_test']
    df_First = pd.DataFrame(results, columns=cols)
    df_First
         Out[86]:
                                           Method ATE test PEHE test
                              0
                                           RF 1.245153 0.190956
                              1 Optimized RF 1.209269 0.196533
                              2 LR 2.385154 2.274672
                              3 Optimized LR 2.386931 2.276535
                              4 IPSW RF 1.192577 0.171409
                                       IPSW LR 2.385161 2.274679
                              6 XL 0.501730 0.143855
In [87]: # # Metrics for Jobs Dataset
    results_ = []
    results_ append(['RF', ATT_Jobs, Rpol])
    results_append(['Optimized RF', ATT_Jobs_Optimized, Rpol_Optimized])
    results_append(['IR', ATT_Jobs_LR, Rpol_LR])
    results_append(['IPSW RF', ATT_Jobs_LR_Optimized, Rpol_LR_Optimized])
    results_append(['IPSW RF', ATT_Jobs_LR_IPSW, Rpol_RF_IPSW])
    results_append(['IPSW LF', ATT_Jobs_LR_IPSW, Rpol_RF_IPSW])
    results_append(['XL', xL] att_test_xL rpol])
    cols_ = ['Method', 'ATT_test', 'Risk Policy']
    if f Second = pd.DataFrame(results_, columns=cols_)
    df_Second
         Out[87]:
                                            Method ATT test Risk Policy
```

	moulou	ATTION	ruok ruildy
0	RF	0.026039	0.219669
1	Optimized RF	0.014877	0.288724
2	LR	0.038637	0.259259
3	Optimized LR	0.038637	0.259259
4	IPSW RF	0.029692	0.206518
5	IPSW LR	0.029424	0.259259
6	XL	0.088744	0.275864

Conclusion:

According to the final metric table of IHDP dataset, XLearner seems to be the most consistent and highly performing model, with the highest ATE and second best PEHE scores as compared to jobs dataset. I would say that, if I had to offer a suggestion to an organization adopting this specific dataset, I would prefer XLearner because of its capability to counterbalanced control to treated data sets, resulting in perhaps the greatest performance overall.

The basic random forest model appears to be the best performing model for this jobs dataset, offering not just the most precise result of treatment impact on the treated, as well as a relatively low risk policy. However, the standard model's ATT is so precise that I feel it is an anomaly, for example if I reran the code, I do not believe I would get the same result. If I had to provide a recommendation to a firm that was using this data set, I would suggest using the linear regressor. It has incredibly less errors on average across three different different versions, demonstrating consistency in performance. Because optimising the hyperparameters is challenging with this model, there is room for the model to improve.

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