

GR5243 Project 4 Doubly Robust Estimations(without scale)

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```
In [1]: import pandas as pd
import numpy as np
import time
from matplotlib import style
from matplotlib import pyplot as plt
import seaborn as sns
from sklearn.linear_model import LogisticRegression
from sklearn.linear_model import LinearRegression
from sklearn.model_selection import train_test_split
from sklearn.model_selection import GridSearchCV
from sklearn.preprocessing import StandardScaler
%matplotlib inline

# set seed
random_state = 2021
```

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In [2]: lowdim_data = pd.read_csv('../data/lowDim_dataset.csv')
highdim_data = pd.read_csv('../data/highDim_dataset.csv')
```

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In [3]: def best_param(data, random_state, param_grid, cv=10):
    ...
    Purpose: to find the best parameter "C" (coefficient of regularization strength) fo

    Parameters:
    data - dataset to best tested on
    random_state - set seed
    param_grid - set of parameter values to test on
    cv - number of folds for cross-validation

    ...

    x = data.drop(['A','Y'], axis = 1)
    y = data[['A']].values.ravel()

    x_train, x_test, y_train, y_test = train_test_split(x, y, test_size=0.25, random_st

    model_cv = GridSearchCV(LogisticRegression(penalty='l1', solver = 'liblinear'), para
    model_cv.fit(x_train, y_train)

    print("The best tuned coefficient of regularization strength is", model_cv.best_para
          "with a testing accuracy of", model_cv.score(x_test, y_test))

    return model_cv.best_params_.get('C')
```

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In [4]: def propensity_score(data, C=0.1, plot = True):
```

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'''
Purpose: to estimate propensity score with L1 penalized logistic regression

Parameters:
data - dataset to estimate on
C - coefficient of regularization strength
plot - print out visualization to show distribution of propensity scores

Returns:
1. ps for Propensity Score
2. Visualization plot to show distribution of propensity scores

'''

T = 'A'
Y = 'Y'
X = data.columns.drop([T,Y])

ps_model = LogisticRegression(random_state=random_state, penalty='l1',
                              solver='liblinear').fit(data[X], data[T])

ps = ps_model.predict_proba(data[X])[:,1] # we are interested in the probability of

if plot:
    df_plot = pd.DataFrame({'Treatment':data[T], 'Propensity Score':ps})

    sns.histplot(data=df_plot, x = "Propensity Score", hue = "Treatment", element =
plt.title("Distribution of Propensity Score by Treatment Group", size=20)
plt.show()

return ps

```

```

In [5]: # setting parameters
param_grid = {"C":[0.01,0.05,0.1,0.3,0.5,0.7,1]}

```

Low Dimensional Case

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In [6]: # use 10-fold cross-validation to tune for the best parameter for logistic regression
DR_low_start = time.time()
c_low = best_param(lowdim_data, random_state=random_state, param_grid=param_grid)

```

The best tuned coefficient of regularization strength is 0.3 with a testing accuracy of 0.8

```

In [7]: # calculate propensity score for low dimensional case
PS_low = propensity_score(lowdim_data, C = c_low, plot = False)

```

```

In [8]: # reload data, add propensity score column and divide data into treat and control group
lowdim_data_new = pd.read_csv('../data/lowDim_dataset.csv')
lowdim_data_new['PS_low'] = pd.Series(PS_low, index=lowdim_data_new.index)
lowdim_treat = lowdim_data[lowdim_data['A'] == 1].reset_index(drop = True)
lowdim_control = lowdim_data[lowdim_data['A'] == 0].reset_index(drop = True)

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In [9]: # fit regression models to treat and control group

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xlow_treat = lowdim_treat.drop(['A', 'Y'], axis=1)
y_low_treat = lowdim_treat['Y']
lr_low_treat = LinearRegression().fit(xlow_treat, y_low_treat)

xlow_control = lowdim_control.drop(['A', 'Y'], axis=1)
y_low_control = lowdim_control['Y']
lr_low_control = LinearRegression().fit(xlow_control, y_low_control)

```

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In [10]: # make prediction based on trained models and construct a full dataset
xlow = lowdim_data_new.drop(['A', 'Y', 'PS_low'], axis=1)
lowdim_data_new['mtreat'] = lr_low_treat.predict(xlow)
lowdim_data_new['mcontrol'] = lr_low_control.predict(xlow)

```

```

In [11]: # perform Doubly Robust Estimation algorithm
DR_low_1 = 0
DR_low_0 = 0

for i in range(len(lowdim_data_new)):
    DR_low_1 = DR_low_1 + (lowdim_data_new['A'][i] * lowdim_data_new['Y'][i] - (lowdim_
    DR_low_0 = DR_low_0 + ((1-lowdim_data_new['A'][i]) * lowdim_data_new['Y'][i] + (lowd

DR_low_ETA = (DR_low_1 - DR_low_0)/len(lowdim_data_new)
DR_low_accu = 1 - abs((DR_low_ETA - 2.0901)/2.0901)
DR_low_end = time.time()
DR_low_time = DR_low_end - DR_low_start

```

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In [12]: # print the ETA, accuracy and algorithm running time results
print(f'Doubly robust estimation method for low dimensional dataset:\n ETA = {DR_low_ET

```

```

Doubly robust estimation method for low dimensional dataset:
ETA = 2.090
Accuracy = 1.000
DR running time = 0.749

```

High Dimensional Case

```

In [13]: # use 10-fold cross-validation to tune for the best parameter for Logistic regression
DR_high_start = time.time()
c_high = best_param(highdim_data, random_state=random_state, param_grid=param_grid)

```

The best tuned coefficient of regularization strength is 0.01 with a testing accuracy of 0.71

```

In [14]: # calculate propensity score for high dimensional case
PS_high = propensity_score(highdim_data, C = c_high, plot = False)

```

```

In [15]: # reload data, add propensity score column and divide data into treat and control group
highdim_data_new = pd.read_csv('../data/highDim_dataset.csv')
highdim_data_new['PS_high'] = pd.Series(PS_high, index=highdim_data.index)
highdim_treat = highdim_data[highdim_data.A == 1].reset_index(drop = True)
highdim_control = highdim_data[highdim_data.A == 0].reset_index(drop = True)

```

```
In [16]: # fit regression model to treat and control group
xhigh_treat = highdim_treat.drop(['A','Y'],axis=1)
yhigh_treat = highdim_treat['Y']
lr_high_treat = LinearRegression().fit(xhigh_treat, yhigh_treat)

xhigh_control = highdim_control.drop(['A','Y'],axis=1)
yhigh_control = highdim_control['Y']
lr_high_control = LinearRegression().fit(xhigh_control, yhigh_control)
```

```
In [17]: # make prediction based on trained models and construct a full dataset
xhigh = highdim_data_new.drop(['A','Y','PS_high'],axis=1)
highdim_data_new['mtreat'] = lr_high_treat.predict(xhigh)
highdim_data_new['mcontrol'] = lr_high_control.predict(xhigh)
```

```
In [18]: # perform Doubly Robust Estimation algorithm
DR_high_1 = 0
DR_high_0 = 0

for i in range(len(highdim_data_new)):
    DR_high_1 = DR_high_1 + (highdim_data_new['A'][i] * highdim_data_new['Y'][i] - (highdim_data_new['A'][i] * DR_high_0))
    DR_high_0 = DR_high_0 + ((1-highdim_data_new['A'][i]) * highdim_data_new['Y'][i] + (highdim_data_new['A'][i] * DR_high_1))

DR_high_ETA = (DR_high_1 - DR_high_0)/len(highdim_data_new)
DR_high_accu = 1 - abs((DR_high_ETA - (-54.8558))/(-54.8558))
DR_high_end = time.time()
DR_high_time = DR_high_end - DR_high_start
```

```
In [19]: # print the ETA, accuracy and algorithm running time result
print(f'Doubly robust estimation method for high dimensional dataset:\n ETA = {DR_high_ETA}, Accuracy = {DR_high_accu}, DR running time = {DR_high_time}
```

```
Doubly robust estimation method for high dimensional dataset:
ETA = -56.622
Accuracy = 0.968
DR running time = 85.069
```