# **Data Loading and Preprocessing**

```
In [1]: import pandas as pd
        import numpy as np
        df = pd.read csv('../data/compas-scores-two-vears.csv')
        df.columns
        Index(['id', 'name', 'first', 'last', 'compas screening date', 'sex', 'dob',
Out[1]:
                'age', 'age cat', 'race', 'juv fel count', 'decile score',
                'juv misd count', 'juv other count', 'priors count',
                'days b screening arrest', 'c jail in', 'c jail out', 'c case number',
                'c offense date', 'c arrest date', 'c days from compas',
                'c charge degree', 'c charge desc', 'is recid', 'r case number',
                'r charge degree', 'r days from arrest', 'r offense date',
                'r charge desc', 'r jail in', 'r jail out', 'violent recid',
                'is violent recid', 'vr case number', 'vr charge degree',
                'vr offense date', 'vr charge desc', 'type of assessment',
                'decile score.1', 'score text', 'screening date',
                'v type of assessment', 'v decile score', 'v score text'.
               'v screening date', 'in custody', 'out custody', 'priors count.1',
                'start', 'end', 'event', 'two vear recid'],
              dtvpe='object')
        compas df = df
In [2]:
        races = ['African-American','Caucasian']
        compas df=compas df[compas df.race.isin(races)]
        compas df.head()
        #use only specific columns for future prediction
        compas df filtered = compas df.loc[:, ['race']].join(compas df.select dtypes(include=['int64']))
        #drop id and duplicate columns
        compas df filtered = compas df filtered.drop(['id', 'decile score.1', 'priors count.1'], axis=1)
        compas df filtered.head()
```

Out[2]:		race	age	juv_fel_count	decile_score	juv_misd_count	juv_other_count	priors_count	is_recid	is_violent_recid	v_decile_score	start	end	eveı
	1	African- American	34	0	3	0	0	0	1	1	1	9	159	
	2	African- American	24	0	4	0	1	4	1	0	3	0	63	
	3	African- American	23	0	8	1	0	1	0	0	6	0	1174	
	6	Caucasian	41	0	6	0	0	14	1	0	2	5	40	
	8	Caucasian	39	0	1	0	0	0	0	0	1	2	747	
4														

```
In [3]: compas_df_filtered["race"].replace(['African-American', 'Caucasian'],[0, 1], inplace=True)

In [4]: y = compas_df_filtered["two_year_recid"]
    X = compas_df_filtered.drop(["two_year_recid"], axis=1)
    from sklearn.model_selection import train_test_split
    X_train, X_test, y_train, y_test=train_test_split(X,y,test_size=0.2, random_state=50, stratify=y)
    print(X_train.shape, y_train.shape, X_test.shape, y_test.shape)
    X_train, X_valid, y_train, y_valid=train_test_split(X_train,y_train,test_size=0.2, random_state=67, stratify=y_train)
    print(X_train.shape, y_train.shape, X_valid.shape, y_valid.shape)

(4920, 13) (4920,) (1230, 13) (1230,)
```

### **Baseline**

(3936, 13) (3936,) (984, 13) (984,)

```
Out[5]:
In [7]: index_afram= (X_test["race"]==0)
         X_test_afram=X test[index afram]
         y test afram=y test[index afram]
         X test cau=X test[-index afram]
         y test cau=y test[-index afram]
In [8]: index afram= (X train["race"]==0)
         X train afram=X train[index afram]
         y train afram=y train[index afram]
         X train cau=X train[-index afram]
         v train cau=v train[-index afram]
In [9]: #calibration
         print(clf.score(X test afram, y test afram))
         print(clf.score(X test cau,y test cau))
         print("calibration score: ",abs(clf.score(X test afram,y test afram)-clf.score(X test cau,y test cau)))
         0.9807692307692307
         0.9860557768924303
         calibration score: 0.005286546123199565
         A5
In [25]: import torch as t
         import torch.nn as nn
```

### **Data Preprocessing**

0.9829268292682927

```
In [26]: data_filtered_afram = compas_df_filtered[compas_df_filtered['race']==0]
    data_filtered_cau = compas_df_filtered[compas_df_filtered['race']==1]

#X_afram = data_filtered_afram[["race", "age", "priors_count", "juv_count", 'score_text', "sex", "c_charge_degree", "decile_score"]]

#y_afram = data_filtered_afram[['two_year_recid']]

X_afram = data_filtered_afram.drop(["two_year_recid"]]

X_cau = data_filtered_cau.drop(["two_year_recid"]]

X_cau = data_filtered_cau.drop(["two_year_recid"]]

X_cau = data_filtered_cau[["two_year_recid"]]
```

```
#X_cau = data_filtered_cau[["race","age","priors_count","juv_count",'score_text',"sex","c_charge_degree","decile_score"]]
#y_cau = data_filtered_cau[['two_year_recid']]

In [27]: X_train_afram, X_test_afram, y_train_afram, y_test_afram = train_test_split(X_afram, y_afram, test_size=0.2)
X_train_cau, X_test_cau, y_train_cau, y_test_cau = train_test_split(X_cau, y_cau, test_size=0.2)
```

#### **Data Preprocessing using Tensor**

```
In [28]: def tensorX(X):
    return t.tensor(np.array(X)).to(t.float32)

def tensorY(X,Y):
    return t.from_numpy(np.array(Y).astype('float32')).reshape(X.shape[0], 1)

In [29]: X_train_afram = tensorX(X_train_afram)
    X_test_afram = tensorX(X_test_afram)
    X_train_cau = tensorX(X_train_cau)
    X_train_afram = tensorX(X_test_cau)

    y_train_afram = tensorY(X_train_afram,y_train_afram)
    y_test_afram = tensorY(X_test_afram,y_test_afram)
    y_train_cau = tensorY(X_train_cau,y_train_cau)
    y_test_cau = tensorY(X_test_cau,y_test_cau)
```

### Algorithm: Prejudice Remover Regularizer Loss Function

```
In [30]: #update LogisticRegression function
    class LogisticRegression(nn.Module):
        def __init__(self,data):
            super(LogisticRegression, self).__init__()
             self.w = nn.Linear(data.shape[1], out_features=1, bias=True)
             self.sigmod = nn.Sigmoid()
        def forward(self, x):
             w = self.w(x)
             output = self.sigmod(w)
             return output
```

```
In [31]: #Define the Prejudice Remover Regularizer Loss Function
    class PRLoss():
        def __init__(self, eta=1.0):
            super(PRLoss, self).__init__()
        self.eta = eta
```

```
def PI(self,output_afram,output_cau):
    N_afram = t.tensor(output_afram.shape[0])
    N_cau = t.tensor(output_cau.shape[0])
# calculate P[y|s]
P_ys = t.stack((t.sum(output_afram),t.sum(output_cau)),axis=0) / t.stack((N_afram,N_cau),axis=0)
# calculate P[y]
P_y = t.sum(t.cat((output_afram,output_cau),0)) / (X_train_afram.shape[0]+X_train_cau.shape[0])
# calculate PI
PI_sly1 = output_afram * (t.log(P_ys[1]) - t.log(P_y))
PI_sly0 = (1- output_afram) *(t.log(1-P_ys[1]) - t.log(1-P_y))
PI_s0y1 = output_cau * (t.log(P_ys[0]) - t.log(P_y))
PI_s0y0 = (1- output_cau)* (t.log(1-P_ys[0]) - t.log(1-P_y))
PI = t.sum(PI_sly1) + t.sum(PI_sly0) + t.sum(PI_s0y1) + t.sum(PI_s0y0)
PI = self.eta * PI
return PI
```

#### Algorithm:Logistical regression with Prejudice Remover Regularizer

```
In [32]:
         #This function has a hyperparameter eta which is the size of the regulating term in the loss function.
         #The fit method will give us the accuracy and caliberation of the fitted model
         class LogisticRegressionWithPRR():
             def init (self, eta=0.0, epochs=10, lr = 0.01):
                  super(LogisticRegressionWithPRR, self). init ()
                 self.eta = eta
                 self.epochs = epochs
                 self.lr = lr
             def fit(self,X train afram, Y train afram, X train cau, Y train cau, X test afram, Y test afram, X test cau, Y test cau):
                 #LoaisticRearession model
                 model afram = LogisticRegression(X train afram)
                 model cau = LogisticRegression(X train cau)
                 criterion = nn.BCELoss()
                 PI = PRLoss(eta=self.eta)
                 epochs = self.epochs
                 #L2 regularization (non-zero weight decay)
                 optimizer = t.optim.Adam(list(model afram.parameters())+ list(model cau.parameters()), self.lr, weight decay=1e-5)
                 for epoch in range(epochs):
                     #train
                     model afram.train()
                     model_cau.train()
```

```
#zero out the aradients
                     optimizer.zero grad()
                     #compute Loss
                     output afram = model afram(X train afram)
                     output cau = model cau(X train cau)
                     log loss = criterion(output afram, y train afram)+ criterion(output cau, y train cau)
                     PI loss = PI.PI(output afram.output cau)
                     loss = PI loss +log loss
                     loss.backward()
                     optimizer.step()
                 #eval
                 model afram.eval()
                 model cau.eval()
                 #calculate accuracy
                 #Accuracy is the average of correctly predicted labels of two groups
                 #Caliberation is the difference of accuracy bewtween the two groups
                 v pred afram = (model afram(X test afram) >= 0.5)
                 v pred cau = (model cau(X test cau) \geq 0.5)
                 #sum of correct prediction/total num
                 accuracy afram = t.sum(v pred afram == v test afram) / v test afram.shape[0]
                 accuracy cau = t.sum(y pred cau == y test cau) / y test cau.shape[0]
                 accuracy = (accuracy afram + accuracy cau) / 2
                 calibration = t.abs(accuracy afram - accuracy cau)
                 return accuracy.item(), calibration.item(), accuracy afram.item(), accuracy cau.item()
         PR = LogisticRegressionWithPRR(eta = 860, epochs = 100, lr = 1e-04)
         PR.fit(X train afram, X train cau, Y train cau, X test afram, Y test cau, Y test cau)
Out[41]: (0.43259480595588684,
          0.15643197298049927,
          0.5108107924461365,
          0.3543788194656372)
```

## **Tuning Hyperparameter eta**

```
In [42]: #Tuning the hyperparameter to find the best model
    eta=np.linspace(0, 100, num=1000)
    hist_acc=np.zeros(1000)
    hist_cal=np.zeros(1000)
```

```
hist afram = np.zeros(1000)
         hist cau = np.zeros(1000)
         for i in range(1000):
             PR = LogisticRegressionWithPRR(eta = eta[i], epochs = 100, lr = 1e-04)
             results = PR.fit(X train afram, Y train afram, X train cau, Y train cau, X test afram, Y test afram, X test cau, Y test cau)
             hist acc[i]=results[0]
             hist cal[i]=results[1]
             hist afram[i]=results[2]
             hist cau[i]=results[3]
In [43]: clf accuracy = np.max(hist acc)
         print("Accuracy:")
         np.max(hist acc)
         Accuracy:
         0.7574337720870972
Out[43]:
In [44]: A5 cal score = np.min(hist cal)
         print("Calibration Score:")
         hist cal[np.argmax(hist acc)] #calibration decreases
         Calibration Score:
         0.0851324200630188
Out[44]:
In [45]: #clf_accuracy = hist_acc[np.argmin(hist cal)]
         print("Accuracy:")
         hist acc[np.argmin(hist cal)]
         Accuracy:
         0.6058223247528076
Out[45]:
In [46]: #A5_cal_score = np.min(hist cal)
         print("Calibration Score:")
         np.min(hist cal) #minimum calibration
         Calibration Score:
         0.0018687844276428223
Out[46]:
In [47]: #np.sort(hist_cal)[0:5]
         hist cal new = hist cal
         np.sort(hist cal new)[0:5]
         array([0.00186878, 0.00220457, 0.00247699, 0.00727418, 0.00727418])
Out[47]:
```

#### **A6**

```
import numpy as np
import pandas as pd
from sklearn.model_selection import train_test_split
from sklearn.linear_model import LogisticRegression
from sklearn.metrics import confusion_matrix
import matplotlib.pyplot as plt
```

#### **Data Loading and Preprocessing**

```
In [97]: compas_df= pd.read_csv('../data/compas-scores-two-years.csv')
#Use the data of African American and Caucasian

races = ['African-American','Caucasian']

compas_df=compas_df[compas_df.race.isin(races)]

compas_df.head()

#use only specific columns for future prediction

compas_df_filtered = compas_df.loc[:, ['race']].join(compas_df.select_dtypes(include=['int64']))

#drop id and duplicate columns

compas_df_filtered = compas_df_filtered.drop(['id','decile_score.1', 'priors_count.1'], axis=1)

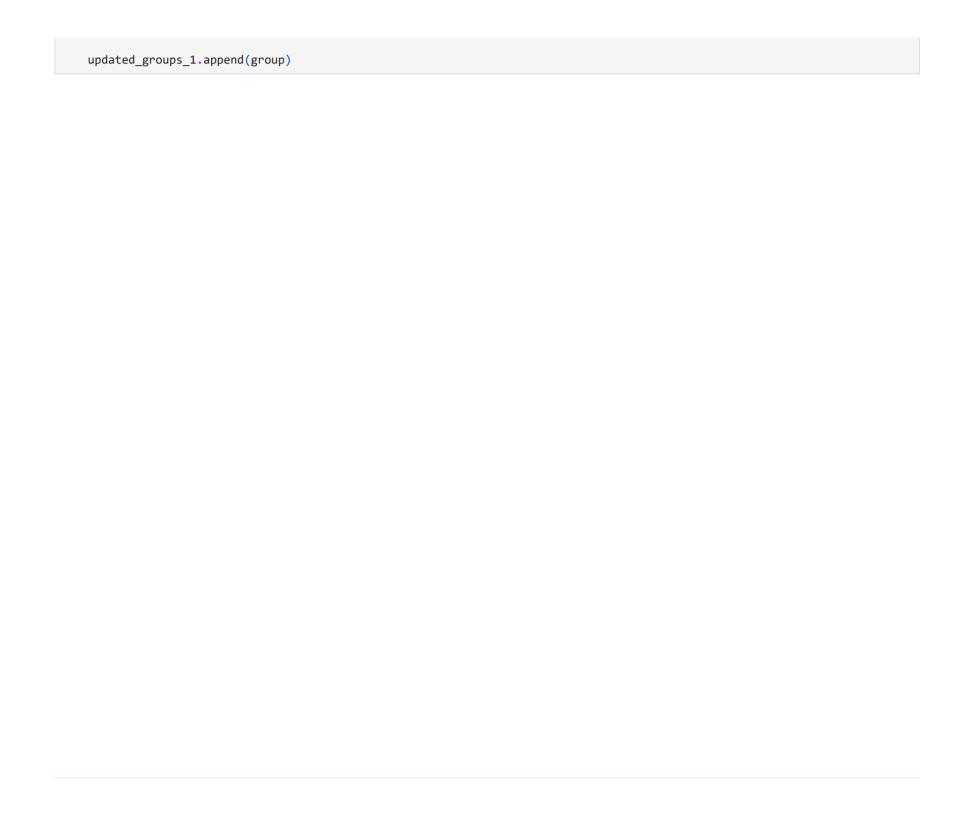
#compas_df_filtered = df[["two_year_recid", "race", "age", "priors_count", "juv_count", 'score_text', "sex", "c_charge_degree", "decile#compas_df_filtered.head()
```

## Implemntaion of algorithms

```
In [98]: #for the algorithms below, we apply on the whole dataset
          X=compas df filtered
          s=X.race
          e=X.decile score
          y=X.two year recid
          def PARTITION(X, e):
              unique values = e.unique() # find unique values of e
              groups = {} # initialize empty dictionary to store resulting DataFrames
              for val in unique values:
                  group df = X[e == val] # filter rows in X where e equals current unique value
                  groups[val] = group df # store resulting DataFrame in dictionary with key = unique value
              return groups # return dictionary of resulting DataFrames
In [99]: resulting groups = PARTITION(X, e)
          first group df = resulting groups[1]
          #print(first group df.head)
          unique values = e.unique()
          print(unique values)
          [3 4 8 6 1 10 5 9 2 7]
          def delta(X, gender):
In Γ100...
              # calculate the baseline probabilities
              p pos base = X.two year recid.mean()
              # calculate probabilities conditioned on gender
              p pos gender = X[X.race == gender].two year recid.mean()
              # calculate the number of gender people in X
              G = len(X[X.race == gender])
              # calculate delta
              delta val = G * abs(p pos gender - p pos base)
              return int(delta_val)
```

### Algorithm: Local massaging

```
#Algorithm 1: Local massaging
In [102...
          resulting groups = PARTITION(X, e)
          updated groups 1 = []
          for group in resulting groups.values():
              males = group[group.race == 'Caucasian']
              females = group[group.race == 'African-American']
              # learn a ranker H for this group
              H = LogisticRegression(random state=42.max iter=500)
              H.fit(group.drop(['race','two year recid'], axis=1), group.two year recid)
              # rank and relabel males
              males females = pd.concat([males,females])
              males females['proba'] = H.predict proba(males females.drop(['two year recid','race'], axis=1))[:, 1]
              males females = males females.sort values(by='proba')
              delta_males = int(delta(group, 'Caucasian'))
              males to relabel = males females[(males females['race'] == 'Caucasian') & (males females['two year recid'] == 0) & (males females['two year recid'] == 0) **
              group.loc[males to relabel.index, 'two year recid'] = 1 - males to relabel.two year recid
              # rank and relabel females
              delta_females = int(delta(group, 'African-American'))
              females to relabel = males females[(males females['race'] == 'African-American') & (males females['two year recid'] == 1)
              group.loc[females to relabel.index, 'two year recid'] = 1 - females to relabel.two year recid
```



```
C:\Users\zhang\Anaconda3\lib\site-packages\pandas\core\indexing.py:1773: SettingWithCopyWarning:
A value is trying to be set on a copy of a slice from a DataFrame.
Try using .loc[row indexer.col indexer] = value instead
See the caveats in the documentation: https://pandas.pydata.org/pandas-docs/stable/user guide/indexing.html#returning-a-view-v
ersus-a-copy
  self. setitem single column(ilocs[0], value, pi)
C:\Users\zhang\Anaconda3\lib\site-packages\pandas\core\indexing.pv:1773: SettingWithCopyWarning:
A value is trying to be set on a copy of a slice from a DataFrame.
Try using .loc[row indexer,col indexer] = value instead
See the caveats in the documentation: https://pandas.pvdata.org/pandas-docs/stable/user_guide/indexing.html#returning-a-view-v
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A value is trying to be set on a copy of a slice from a DataFrame.
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A value is trying to be set on a copy of a slice from a DataFrame.
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C:\Users\zhang\Anaconda3\lib\site-packages\pandas\core\indexing.py:1773: SettingWithCopyWarning:
A value is trying to be set on a copy of a slice from a DataFrame.
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  self. setitem single column(ilocs[0], value, pi)
C:\Users\zhang\Anaconda3\lib\site-packages\pandas\core\indexing.py:1773: SettingWithCopyWarning:
A value is trying to be set on a copy of a slice from a DataFrame.
Try using .loc[row indexer, col indexer] = value instead
See the caveats in the documentation: https://pandas.pydata.org/pandas-docs/stable/user guide/indexing.html#returning-a-view-v
```

```
ersus-a-copy
            self. setitem single column(ilocs[0], value, pi)
          C:\Users\zhang\Anaconda3\lib\site-packages\pandas\core\indexing.pv:1773: SettingWithCopyWarning:
          A value is trying to be set on a copy of a slice from a DataFrame.
          Try using .loc[row indexer.col indexer] = value instead
          See the caveats in the documentation: https://pandas.pydata.org/pandas-docs/stable/user guide/indexing.html#returning-a-view-v
          ersus-a-copy
            self. setitem single column(ilocs[0], value, pi)
          C:\Users\zhang\Anaconda3\lib\site-packages\pandas\core\indexing.py:1773: SettingWithCopyWarning:
          A value is trying to be set on a copy of a slice from a DataFrame.
          Try using .loc[row indexer.col indexer] = value instead
          See the caveats in the documentation: https://pandas.pydata.org/pandas-docs/stable/user guide/indexing.html#returning-a-view-v
          ersus-a-copy
            self. setitem single column(ilocs[0], value, pi)
          C:\Users\zhang\Anaconda3\lib\site-packages\pandas\core\indexing.pv:1773: SettingWithCopyWarning:
          A value is trying to be set on a copy of a slice from a DataFrame.
          Try using .loc[row indexer,col indexer] = value instead
          See the caveats in the documentation: https://pandas.pvdata.org/pandas-docs/stable/user_guide/indexing.html#returning-a-view-v
          ersus-a-copy
            self. setitem single column(ilocs[0], value, pi)
          updated df 1 = pd.concat(updated groups 1)
In Γ103...
          updated df 1 = updated df 1.sort index()
          print(sum(updated df 1['two year recid']!=compas df filtered['two year recid']))
          #67 Labels have been changed
          67
```

## Algorithm: Local preferential sampling

```
In [104...
#Algorithm 2: Local preferential sampling
resulting_groups = PARTITION(X, e)
updated_groups_2 = []
for group in resulting_groups.values():
    males = group[group.race == 'Caucasian']
    females = group[group.race == 'African-American']

# Learn a ranker H for this group
H = LogisticRegression(random_state=42,max_iter=500)
H.fit(group.drop(['race','two_year_recid'], axis=1), group.two_year_recid)

# rank delete, and relabel males
males_females = pd.concat([males, females])
```

```
males females['proba'] = H.predict proba(males females.drop(['two year recid', 'race'], axis=1))[:, 1]
          males females = males females.sort values(by='proba')
          delta males = int(0.5*delta(group, 'Caucasian'))
          males to delete = males females[(males females['race'] == 'Caucasian') & (males females['two year recid'] == 1) & (male
          males to duplicate = males females[(males females['race'] == 'Caucasian') & (males females['two year recid'] == 0) & (males
          group = group[~group.index.isin(males to delete.index)]
          group = pd.concat([group, males to duplicate], ignore index=True)
          # rank delete.and relabel females
          delta females = int(0.5*delta(group, 'African-American'))
          females to delete = males females[(males females['race'] == 'African-American') & (males females['two year recid'] == 0) &
          females to duplicate = males females[(males females['race'] == 'African-American') & (males females['two year recid'] == 1
          group = group[~group.index.isin(females to delete.index)]
          group = pd.concat([group, females to duplicate], ignore index=True)
          updated groups 2.append(group)
updated df 2 = pd.concat(updated groups 2)
#Now we have 3 data frames.compas df filtered is the original one, updated df 1 is the one of algorithm1 updated df 1 is the or
```

```
# split the data into train, validation and test set

# Split data into training and validation/test sets
train_df, val_test_df = train_test_split(compas_df_filtered, test_size=0.2, random_state=42)

# Split validation/test set into validation and test sets
val_df, test_df = train_test_split(val_test_df, test_size=0.5, random_state=42)

# Split data into training and validation/test sets
train_df_algo1, val_test_df_algo1 = train_test_split(updated_df_1, test_size=0.2, random_state=42)

# Split validation/test set into validation and test sets
val_df_algo1, test_df_algo1 = train_test_split(val_test_df_algo1, test_size=0.5, random_state=42)

# Split data into training and validation/test sets
train_df_algo2, val_test_df_algo2 = train_test_split(updated_df_2, test_size=0.2, random_state=42)

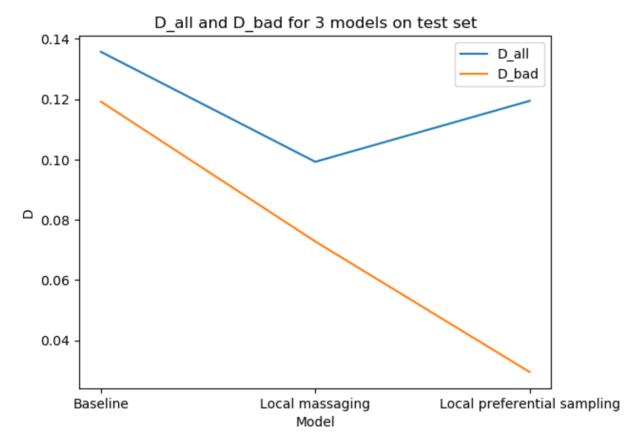
# Split validation/test set into validation and test sets
val_df_algo2, test_df_algo2 = train_test_split(val_test_df_algo2, test_size=0.5, random_state=42)
```

```
X train1 = train df.drop(['is recid', 'race'], axis=1)
In [106...
          y train1 = train df['is recid']
          X train2 = train df algo1.drop(['is recid', 'race'], axis=1)
          y_train2 = train_df_algo1['is_recid']
          X train3 = train df algo2.drop(['is recid', 'race', 'proba'], axis=1)
          y train3 = train df algo2['is recid']
          # Fit logistic regression model for X train1
          logreg1 = LogisticRegression()
          logreg1.fit(X train1, y train1)
          # Fit logistic regression model for X train2
          logreg2 = LogisticRegression()
          logreg2.fit(X train2, y train2)
          # Fit logistic regression model for X train3
          logreg3 = LogisticRegression()
          logreg3.fit(X train3, y train3)
```

```
C:\Users\zhang\Anaconda3\lib\site-packages\sklearn\linear model\ logistic.pv:818: ConvergenceWarning: lbfgs failed to converge
          (status=1):
          STOP: TOTAL NO. of ITERATIONS REACHED LIMIT.
          Increase the number of iterations (max iter) or scale the data as shown in:
              https://scikit-learn.org/stable/modules/preprocessing.html
          Please also refer to the documentation for alternative solver options:
              https://scikit-learn.org/stable/modules/linear model.html#logistic-regression
            extra warning msg= LOGISTIC SOLVER CONVERGENCE MSG.
          C:\Users\zhang\Anaconda3\lib\site-packages\sklearn\linear model\ logistic.py:818: ConvergenceWarning: lbfgs failed to converge
          (status=1):
          STOP: TOTAL NO. of ITERATIONS REACHED LIMIT.
          Increase the number of iterations (max iter) or scale the data as shown in:
              https://scikit-learn.org/stable/modules/preprocessing.html
          Please also refer to the documentation for alternative solver options:
              https://scikit-learn.org/stable/modules/linear model.html#logistic-regression
            extra warning msg= LOGISTIC SOLVER CONVERGENCE MSG,
          C:\Users\zhang\Anaconda3\lib\site-packages\sklearn\linear model\ logistic.py:818: ConvergenceWarning: lbfgs failed to converge
          (status=1):
          STOP: TOTAL NO. of ITERATIONS REACHED LIMIT.
          Increase the number of iterations (max iter) or scale the data as shown in:
              https://scikit-learn.org/stable/modules/preprocessing.html
          Please also refer to the documentation for alternative solver options:
              https://scikit-learn.org/stable/modules/linear_model.html#logistic-regression
            extra warning msg= LOGISTIC SOLVER CONVERGENCE MSG,
          LogisticRegression()
Out[106]:
In [107... #Get results for validation set
          # For X val1
          X val1 = val df.drop(['is recid', 'race'], axis=1)
          y val1 = val df['is recid']
          y pred val1 = logreg1.predict(X val1)
          # For X val2
          X val2 = val df algo1.drop(['is recid', 'race'], axis=1)
          y val2 = val df algo1['is recid']
          y pred val2 = logreg2.predict(X val2)
          # For X val3
          X val3 = val df algo2.drop(['is recid', 'race', 'proba'], axis=1)
          y val3 = val df algo2['is recid']
          y pred val3 = logreg3.predict(X val3)
```

```
#Get results for test set
In [108...
          # For X test1
          X test1 = test df.drop(['is recid', 'race'], axis=1)
          v test1 = test df['is recid']
          y pred test1 = logreg1.predict(X test1)
          # For X test2
          X test2 = test df algo1.drop(['is recid', 'race'], axis=1)
          y test2 = test df algo1['is recid']
          v pred test2 = logreg2.predict(X test2)
          # For X test3
          X test3 = test df algo2.drop(['is recid','race','proba'], axis=1)
          v test3 = test df algo2['is recid']
          v pred test3 = logreg3.predict(X test3)
In [109... # Make a copy of the original test dataframes
          test df1 = test df.copy()
          test df2 = test df algo1.copy()
          test df3 = test df algo2.copv()
          test df1["y pred"] = y pred test1
          test df2["y pred"] = y pred test2
          test df3["y pred"] = y pred test3
In [110... # Define function to compute D all and D bad
          def compute D(test df, y pred):
              # Compute D all
              D all = test df.loc[test df['race'] == 'African-American', y pred].mean() - \
                      test df.loc[test df['race'] == 'Caucasian', y pred].mean()
              # Compute D bad
              decile_scores = list(range(1, 6))
              D bad = 0
              for score in decile scores:
                  P_score = test_df.loc[(test_df['race'] == 'African-American') & (test_df['decile_score'] == score), y_pred].mean()
                  P_score -= test_df.loc[(test_df['race'] == 'Caucasian') & (test_df['decile_score'] == score), y_pred].mean()
                  P_y = test_df.loc[test_df['decile_score'] == score, y_pred].mean()
                  D bad += ((P score) * (P y))
```

```
return D all, D bad
          # Compute D all and D bad for the three logistic regression models on the three test sets
          D all 1, D bad 1 = compute D(test df1, 'two year recid')
          D all 2, D bad 2 = compute D(test df2, 'two year recid')
          D all 3, D bad 3 = compute D(test df3, 'two year recid')
          # Calculate D all and D bad for each model on the test set
In [111...
          D all test = [D all 1,D all 2,D all 3]
          D bad test = [D bad 1, D bad 2, D bad 3]
          plt.plot(['Baseline', 'Local massaging', 'Local preferential sampling'], D all test, label='D all')
          plt.plot(['Baseline', 'Local massaging', 'Local preferential sampling'], D bad test, label='D bad')
          # Add axis Labels and title
          plt.xlabel('Model')
          plt.ylabel('D')
          plt.title('D all and D bad for 3 models on test set')
          # Add Leaend
          plt.legend()
          # Show the plot
          plt.show()
          print(D all test)
          print(D bad test)
```



[0.13573754620665374, 0.09924529132195037, 0.11945715462632062] [0.11922600741704245, 0.07281751731787592, 0.02948165449549718]

# **Comparision plots:**

```
from sklearn.metrics import accuracy_score
import matplotlib.pyplot as plt

# Define function to compute calibration
def compute_calibration(df):
    african_american = df[df['race'] == 'African-American']
    caucasian = df[df['race'] == 'Caucasian']
    african_american_accuracy = accuracy_score(african_american['two_year_recid'], african_american['y_pred'])
    caucasian_accuracy = accuracy_score(caucasian['two_year_recid'], caucasian['y_pred'])
    return abs(african_american_accuracy - caucasian_accuracy)

# Compute accuracy and calibration for test set 1
```

```
accuracy test1 = accuracy score(test df1['two year recid'], test df1['y pred'])
calibration test1 = compute calibration(test df1)
baseline cal=calibration test1*len((e.unique()))
# Compute accuracy and calibration for test set 2
accuracy test2 = accuracy score(test df2['two year recid'], test df2['y pred'])
calibration test2 = compute calibration(test df2)
# Compute accuracy and calibration for test set 3
accuracy test3 = accuracy score(test df3['two year recid'], test df3['v pred'])
calibration test3 = compute calibration(test df3)
A5 acc = hist acc[158]
A5 calibration = np.sort(hist cal new)[2]
# Plot results
accuracies = [accuracy test1.A5 acc. accuracy test2. accuracy test3]
calibrations = [baseline cal, A5 calibration.calibration test2, calibration test3]
models = ['Baseline', 'A5', 'A6 LM', 'A6 LPS']
accs = accuracies
cals = calibrations
fig, axs = plt.subplots(nrows=1, ncols=2, figsize=(10, 5))
axs[0].bar(models, accuracies)
axs[0].set title('Accuracy on test sets')
axs[1].bar(models, calibrations)
axs[1].set title('Calibration on test sets')
plt.show()
```

