```
In [17]: import sklearn
         from sklearn.base import BaseEstimator
         from sklearn.preprocessing import scale
         from sklearn import model selection
         from sklearn.decomposition import PCA
         from sklearn.linear model import LinearRegression
         from sklearn.cross decomposition import PLSRegression, PLSSVD
         from sklearn.linear model import LinearRegression as lr
         from sklearn.linear model import ElasticNetCV, Ridge
         from sklearn.model selection import GridSearchCV
         from sklearn.model selection import KFold
         from sklearn.metrics import mean squared error
         from sklearn import datasets, linear model
         from sklearn.model selection import train test split
         from sklearn.preprocessing import PolynomialFeatures as pf
         from sklearn.preprocessing import LabelEncoder as le
         from sklearn.preprocessing import KBinsDiscretizer as kb
         from sklearn.preprocessing import StandardScaler
         from sklearn.decomposition import PCA
         from sklearn.cross decomposition import PLSRegression
         from sklearn.metrics import mean squared error as MSE
         from sklearn.metrics import make scorer
         from sklearn.pipeline import make pipeline, Pipeline
         from collections import Counter
         import tabulate as tb
         import numpy as np
         import matplotlib.pyplot as plt
         import matplotlib.colors
         import seaborn
         import scipy as sp
         from scipy.interpolate import CubicSpline, UnivariateSpline
```

```
import collections
         import os
         import os.path
         import random
         import re
         import glob
         import pandas as pd
         import requests
         import json
         import math
         from patsy import dmatrix
         %matplotlib inline
In [9]: gss tr = pd.read csv('gss train.csv')
         gss te = pd.read csv('gss test.csv')
In [59]: np.random.seed(51)
In [60]: #gss tr = pd.read csv('gss train.csv')
         #gss te = pd.read csv('gss test.csv')
         assert set(gss te.columns) == set(gss tr.columns)
         collect = {}
         for col in gss te.columns:
             pred test, pred train = list(gss_te[col]), list(gss_tr[col])
             label e = le().fit(pred test+pred train)
             gss te[col] = label e.transform(pred test)
             gss tr[col] = label e.transform(pred train)
             collect[col] = label e.classes
```

## **Egalitarianism and income**

(20 points) Perform polynomial regression to predict egalit\_scale as a function of income06. Use and plot 10-fold cross-validation to select the optimal degree d for the polynomial based on the MSE. Plot the resulting polynomial fit to the data, and also graph the average marginal effect (AME) of income06 across its potential values. Be sure to provide substantive interpretation of the results.

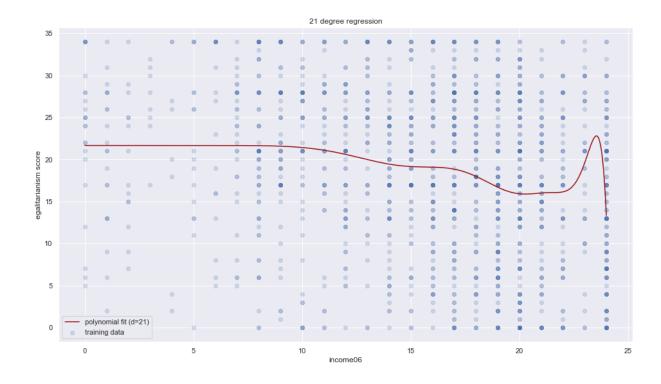
```
best_d = cv.best_params_['poly__degree']

In [106]: income_plot = np.linspace(income_train.min(), income_train.max(), 10000
    ).reshape(-1, 1)
    plot_y = best.predict(income_plot)

plt.scatter(income_train, y_train, label='training data', alpha=0.2)
    plt.plot(income_plot, plot_y, label=f'polynomial fit (d={best_d})', c=p
    lt.cm.Maroons(0.9))

plt.legend()

plt.title(f'{best_d} degree regression')
    plt.xlabel('income06')
    plt.ylabel('egalitarianism score')
```



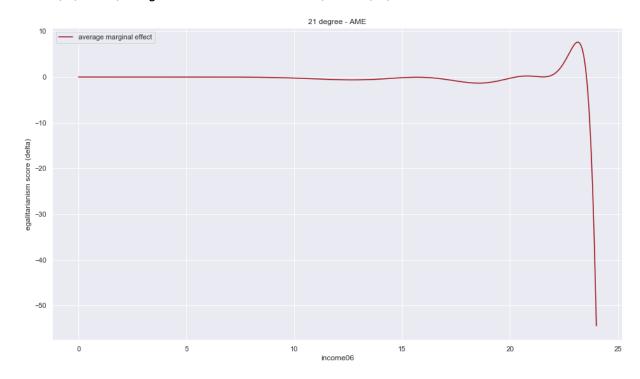
• I ran the polynomial model for degrees ranging from 1-50, and the lowest MSE value corresponded to a degree of 21, for 10-fold cross validation. However, we can see from the plot that this curve does not fit the data very well.

```
In [108]: sp = float(income_plot[1]-income_plot[0])
    plot_ame = np.gradient(plot_y.reshape(-1), sp)
    plt.plot(income_plot, plot_ame, label=f'average marginal effect', c=plt
    .cm.Maroons(0.9))
    plt.legend()

plt.title(f'{best_d} degree - AME')
```

```
plt.xlabel('income06')
plt.ylabel('egalitarianism score (delta)')
```

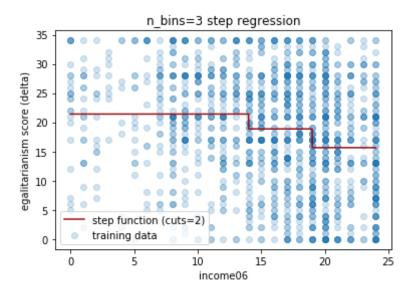
Out[108]: Text(0, 0.5, 'egalitarianism score (delta)')



- The MSE for degree = 21 was roughly 80, and the AME is zero up until roughly degree = 23 (suggests little to no effect). Around this point, we can see that egalitarism score decreases as income goes up. However, as mentioned previously, the polynomial form (despite varying degrees) does not fit this data very well.
- The computed AME = roughly -.36, though that could be explained by the suggest shift in trajectory described.

(20 points) Fit a step function to predict egalit\_scale as a function of income06, and perform 10-fold cross-validation to choose the optimal number of cuts. Plot the fit and interpret the results.

```
In [43]: | step pipe = Pipeline([('cut', kb()), ('lr', lr())])
         params = {'cut_n_bins': range(2, 20)} #can't do 1, threw error
         cv = GridSearchCV(step pipe, params, n jobs=-1, scoring='neg mean squar
         ed error', cv=10)
In [44]: cv.fit(income train, y train)
         best = cv.best estimator
         best bins = cv.best params ['cut n bins']
In [53]: income plot = np.linspace(income train.min(), income train.max(), 10000
         ).reshape(-1, 1)
         plot y = best.predict(income plot)
         plt.scatter(income train, y train, label='training data', alpha=0.2)
         plt.plot(income plot, plot y, label=f'step function (cuts={best cuts-
         1)', c=plt.cm.\overline{R}eds(0.9))
         plt.title(f'n bins={best bins} step regression')
         plt.legend()
         plt.xlabel('income06')
         plt.ylabel('egalitarianism score (delta)')
         #MSE:
         #89.90076400132222
         #84.95884373090617
         #88.77919769150931
         #87.40146010254689
         #84.50891974424688
         #102.57410227823715
         #85.44463506276448
         #98.27130464359588
         #90.82594483834163
         #89.9863648235786
Out[53]: Text(0, 0.5, 'egalitarianism score (delta)')
```



• We can see that 3 bins = the optimal number of cuts (2). The MSE for 3 bins is still quite high (~89), and quite high across the board (see previous cell) across the range of cuts that I chose (across 10-fold cv), suggesting that a step function might not fit our data well (it's an overly restrictive approach for data of this spread). We can also see that there seem to be trends that we can isolate, bin to bin. In other words, egalitarian scores seem to vary across the three bins - it differs across low, medium, and high income.

(20 points) Fit a natural regression spline to predict egalit\_scale as a function of income06. Use 10-fold cross-validation to select the optimal number of degrees of freedom, and present the results of the optimal model.

```
In [54]: y_train = gss_tr['egalit_scale']
    x_train = gss_tr['income06'].values.reshape(-1, 1)
    y_test = gss_te['egalit_scale']
    x_test = gss_te['income06'].values.reshape(-1, 1)
In [58]: track = {'mean_squared_error': [], 'mods': [], 'k': []}
```

```
cv 10 = KFold(n splits=10)
splits = cv 10.split(x train, y train)
for train, test in splits:
    x tr n = x train[train].flatten()
   y tr n = y train[train].values.flatten()
   x flat = x train[test].flatten()
   y flat = y train[test].values.flatten()
    res = pd.DataFrame(data={'feat':x tr n, 'predict':y tr n}, index=pd
.RangeIndex(len(x tr n)))
    res = res.groupby('feat', as index=False)['predict'].mean()
    x ch = res['feat'].values
    y ch = res['predict'].values
    spline = CubicSpline(x ch, y ch, bc type='natural')
    y = sp(x flat)
    mean squared error = np.mean((y egal - y flat) ** 2)
    print(mean squared error)
    track['mean squared_error'].append(mean_squared_error)
    track['mods'].append(spline)
best = np.argmin(track['mean squared error'])
print(best)
best = track['mods'][best]
plot x = np.linspace(0, 25, 10000).reshape(-1, 1)
plot y = best(plot x)
#MSE printed below:
88.90076400132222
83.95884373090617
88.77919769150931
86.40146010254689
05 50001074424600
```

```
85.508919/4424088
101.57410227823715
86.44463506276448
97.27130464359588
89.82594483834163
75.9863648235786
9
```

```
In [57]: plt.plot(plot_x, plot_y, c='C3')
   plt.scatter(x_train, y_train, c='C2', alpha=.2)
   plt.ylim(0, 35)
   plt.xlabel('income06')
   plt.ylabel('egalitarism')
   plt.title('2 Cuts Step Function Regression')
   plt.show()
```



• The MSE is quite high here as well - coming in at around 89. The optimal # of cuts is 2 here, like we obtained using the step function. The flexible nature of the spline tells us more about how income and egalitarianism vary across all three bins, but we can see that despite smoothing, this model does not fit the data very well.

## Egalitarianism and everything

(20 points total) Estimate the following models using all the available predictors (be sure to perform appropriate data pre-processing (e.g., feature standardization) and hyperparameter tuning (e.g. lambda for PCR/PLS, lambda and alpha for elastic net). Also use 10-fold cross-validation for each model to estimate the model's performance using MSE): a. (5 points) Linear regression b. (5 points) Elastic net regression c. (5 points) Principal component regression d. (5 points) Partial least squares regression

a)

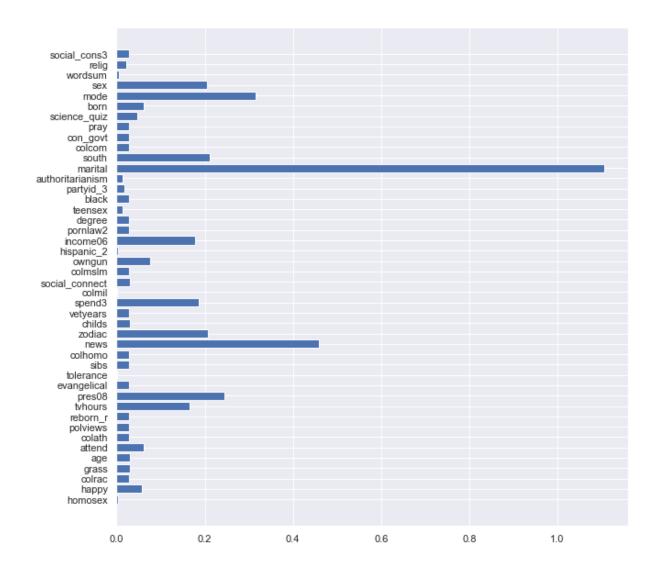
```
In [67]: x 1, x 2 = StandardScaler(), StandardScaler()
         x_1, x_2 = x_1.fit(x_tr), x_2.fit(x_test)
         x train shape, x test shape = x = 1.transform(x train), x 2.transform(x t
         est)
         (1481, 44) (1481, 1)
In [68]: linreg = GridSearchCV(lr(), {}, scoring='neg mean squared error', cv=10
         linreq.fit(x train shape, y tr)
         best linreg = linreg.best estimator
         linreg err = MSE(y test, best linreg.predict(x test shape))
         #Best linear regression MSE: 63.928057088260516
         b)
In [ ]: elnet = ElasticNetCV(l1 ratio=[.1, .5, .7, .9, .95, .99, 1], n alphas=1
         0, cv=10)
         y train = y train.reshape(-1,)
         elnet.fit(x train shape, y train)
         elnet err = MSE(y test, elnet.predict(x test shape))
```

```
#Best ElasticNet MSE: 62.56902435370069, lambda = 0.19753166246833653
         \#alpha = 0.5
         c)
In [ ]: pc reg = Pipeline([('pca', PCA()), ('ridge', Ridge())])
         grid = {'pca n components':np.arange(2, 24, 2), 'ridge alpha':[0.01,
         0.05]+list(np.arange(0.1, 1, 10))}
         pc cv = GridSearchCV(pc reg, grid, scoring='neg mean squared error', cv
         =10. refit=True)
         pc cv.fit(x train shape, y train)
In [ ]: best = pc cv.best estimator
         opt n = pc cv.best params ['pca n components']
         opt lambda = pc cv.best params ['ridge alpha']
         pc err = MSE(v test, best.predict(x test shape))
         # Best PCR MSE: 62.32194871839432, lambda = 0.05
         d)
In [72]: partial ls = PLSRegression()
         partial lscv = GridSearchCV(partial ls, grid={'n components':np.arange(
         2, 21, 2)}, scoring='neg_mean_squared_error', cv=10)
         partial lscv.fit(x train shape, y train)
         best = partial lscv.best estimator
In [ ]: opt n = partial lscv.best params ['n components']
         partial lscv err = MSE(y test, best.predict(x test shape))
         #Best PLS MSE: 63.927709350623815
           1. (20 points) For each final tuned version of each model fit, evaluate feature importance by
```

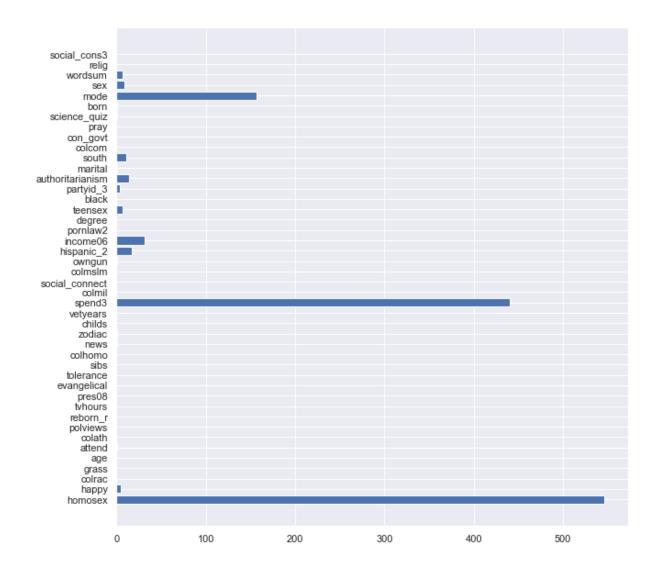
generating feature interaction plots. Upon visual presentation, be sure to discuss the substantive results for these models and in comparison to each other (e.g., talk about

feature importance, conditional effects, how these are ranked differently across different models, etc.).

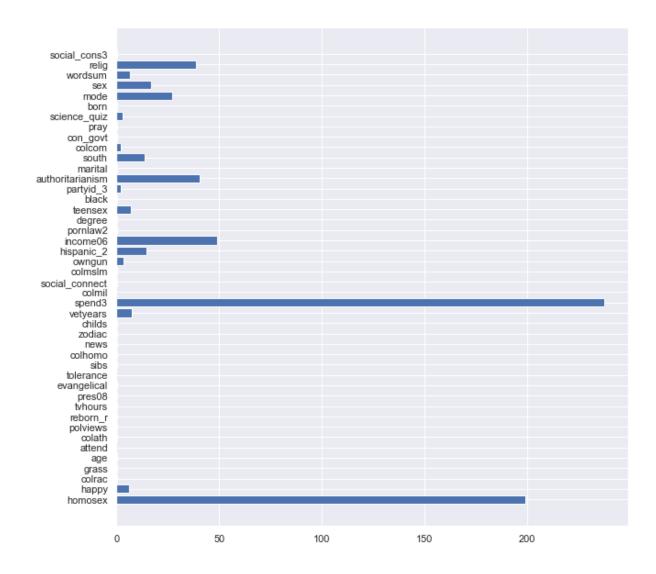
```
In [113]: from scipy import stats
          def feature imp(mod, data, y, x test, y test, header):
              assert y in data.columns
              columns = set(data.columns)
              columnss.remove(v)
              yname, value = [], []
              for i, column in enumerate(columnss):
                  if y == column: continue
                  yname.append(column)
                  x amend = np.zeros(x test.shape)
                  x_{amend}[:, i] = x_{test}[:, i]
                  y amend = model.predict(x amend)
                  y amend = y amend.reshape(y_test.shape)
                  value.append(stats.kruskal(y_amend, y_test)[0])
              plt.figure(figsize=(10, 10))
              plt.barh(np.arange(len(yname)), value)
              plt.yticks(np.arange(len(yname)), yname)
In [114]: feature imp(best linreg, gss tr, 'egalit scale', x test shape, y test,
           'Linear Regression')
```



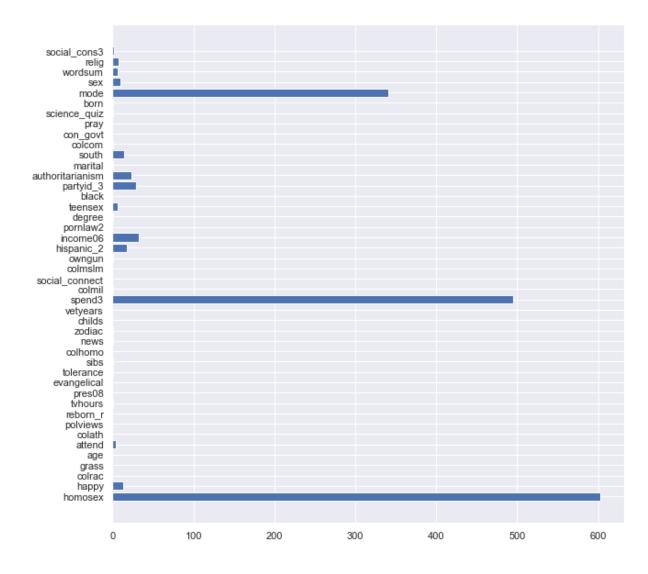
```
In [115]: feature_imp(elnet, gss_tr, 'egalit_scale', x_test, y_test, 'Elastic Ne
t')
```



```
In [116]: feature_int_plot(best_pca, gss_tr, 'egalit_scale', x_test, y_test, 'PC
R')
```



```
In [118]: feature_imp(best_pls, gss_tr, 'egalit_scale', x_test, y_test, 'PLS')
```



• The different models used reach similar conclusions (regarding feature importance). We can see that 'homosex' and 'spend3' have a very high H score across all of the models (except linear regression), implying that they strongly predict egalitarianism scores. Interestingly, 'marital' is a strong predictor of egalitarianism in the linear regression, but not in any of the other models. Additionally, many of the features in the linear regression model seem to be interacting together to inform egalitarianism scores (though 'marital' is the most influential,

followed by 'news'), while the other models only have a couple of predominant/influential predictors (for example, 'mode', 'spend3', and 'homosex' in partial least squares). The models besides linear regression seem to have done a better job of isolating the most relevant features that predict egalitarianism ('spend3', and 'homosex' across elastic net, pcr, and pls; 'mode' isn't quite as weighted through pcr).

In [ ]: