## PS7

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### 0.1 MACS 30150 Problem Set 7

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```
In [1]: import pandas as pd
    import numpy as np
    import matplotlib.pyplot as plt
    import statsmodels.api as sm
    from scipy.interpolate import LSQUnivariateSpline as lsq
    from sklearn.linear_model import LogisticRegression as LR
    from sklearn.cross_validation import train_test_split
    from sklearn.model_selection import train_test_split as tts
    from sklearn.model_selection import LeaveOneOut, KFold
    from sklearn.metrics import classification report
```

C:\ProgramData\Anaconda3\lib\site-packages\sklearn\cross\_validation.py:41: DeprecationWarning: "This module will be removed in 0.20.", DeprecationWarning)

#### 1 Problem 1

#### 1.1 1-(a)

Firstly, let us begin by reading in the data. We can see that, as the question elaborates, there are three levels to our dependent variable cultivar.

I first report the coefficients  $\beta_{1,0}$ ,  $\beta_{1,1}$ ,  $\beta_{1,2}$ ,  $\beta_{1,3}$ ,  $\beta_{1,4}$ ,  $\beta_{2,0}$ ,  $\beta_{2,1}$ ,  $\beta_{2,2}$ ,  $\beta_{2,3}$  and  $\beta_{2,4}$  for the model that uses the entire dataset (as opposed to the training and test split). For j = 1, we have the following (rounded to the nearest hundredth):

$$(\beta_{1,0}, \beta_{1,1}, \beta_{1,2}, \beta_{1,3}, \beta_{1,4}) = (-23.84, 1.58, -0.14, 1.56, 0.11)$$

For j = 2, we have the following (rounded to the nearest hundredth):

$$(\beta_{2,0},\beta_{2,1},\beta_{2,2},\beta_{2,3},\beta_{2,4}) = (23.37,-1.37,-0.51,0.49,-1.15)$$

Now let us subset the dataset into training and test datasets. As the total number of observations is 176 and we take 25 percent of it as the test dataset, the number of observations for test data is 44.

After the test-train split, let us train the model. The coefficients are slightly different from the ones from the model using the entire dataset (yet having the same sign), and they are as follows. For j = 1, we have the following (rounded to the nearest hundredth):

$$(\beta_{1,0},\beta_{1,1},\beta_{1,2},\beta_{1,3},\beta_{1,4}) = (-24.03,1.70,-0.27,1.22,0.02)$$

For j = 2, we have the following (rounded to the nearest hundredth):

$$(\beta_{2,0}, \beta_{2,1}, \beta_{2,2}, \beta_{2,3}, \beta_{2,4}) = (22.78, -1.47, -0.33, 0.66, -0.92)$$

Let us now produce the classification report as directed by the question. From the below table, it can be seen that the error rates (i.e. 1 - precision) are 0.13, 0.00, and 0.00 for j = 1, 2, and 3 respectively. The most accurately predicted category (or the category that the model is best at predicting), in terms of precision, recall, and F1 score is j = 3. However, j = 3 is not the class with the most observations.

```
In [6]: ## fitting
        y_pred = mlogit_train.predict(X_test)
        ## classification report
        print(classification_report(y_test, y_pred))
             precision
                           recall f1-score
                                               support
          1
                  0.87
                             1.00
                                       0.93
                                                    13
          2
                  1.00
                             0.90
                                       0.95
                                                    21
                  1.00
                             1.00
                                       1.00
                                                    10
avg / total
                  0.96
                             0.95
                                       0.96
                                                    44
```

Let us now calculate the *MSE*. Note that the mean squared error (*MSE*) for categorical variables can be written as follows:

$$\frac{1}{N} \sum_{i=1}^{N} \mathbf{1}_{y_i \neq \hat{y}_i}$$

in which N is the number of observations, and  $\mathbf{1}_{y_i \neq \hat{y}_i}$  is the indicator function that equals to 1 when  $y_i \neq \hat{y}_i$  (i.e. the predicted and actual values are not the same) and equals to 0 when  $y_i = \hat{y}_i$  (i.e. the predicted and actual values are the same).

I use the below function  $mse_cat$  to calculate the MSE; the value turns out to be approximately 0.04545.

#### 1.2 1-(b)

Let us now perform the leave-one-out cross-validation (LOOCV). To do so, I define the function cv\_loo\_mlogit for using LOOCV with multinomial logistic regression defined in the question.

This will return the  $CV_{loo}$  value and the actual and predicted dependent variable values that will be used for calculating the classification report (and error rate, accordingly).

```
In [8]: def cv_loo_mlogit(Xvals, yvals, rtn_vectors=True):
            print("(1) Printing when actual and predicted values aren't same...")
            N = yvals.shape[0]
            xN = Xvals.shape[1]
            add_up = 0
            y_pred_vec = np.zeros(N)
            for i in range(N):
                if i == 0:
                    X_loo = Xvals[1:]
                    y_loo = yvals[1:]
                elif i == N-1:
                    X_{loo} = Xvals[0:N-1]
                    y_{loo} = yvals[0:N-1]
                else:
                    X_loo_front, X_loo_back = Xvals[0:i], Xvals[i+1:]
                    y_loo_front, y_loo_back = yvals[0:i], yvals[i+1:]
                    X_loo = np.vstack((X_loo_front, X_loo_back))
                    y_loo = np.hstack((y_loo_front, y_loo_back))
                mlogit_loo = LR(random_state=0, solver='lbfgs',
                                multi_class='multinomial').fit(X_loo, y_loo)
                X_leftout = np.reshape(Xvals[i], (1, xN))
                loo_pred = mlogit_loo.predict(X_leftout)
                y_pred_vec[i] = loo_pred
                if loo_pred[0] != yvals[i]:
                    add_up += 1
                    print('Observ. #{}:'.format(i),
                          'Actual Value: {}, Predicted Value: {}'.format(yvals[i], loo_pred[0]
            loo = add_up / N
            print()
            print("(2) CV_LOO = {}".format(loo))
            if rtn_vectors:
                return loo, yvals, y_pred_vec
            else:
                return loo
```

Produced below are the results for the  $CV_{loo}$  as well as the cross validation table. The error rates for each category are 0.1, 0.09, and 0.04 for j = 1, 2 and 3 respectively. Error rates have decreased

for j = 1, but for j = 2 and j = 3 the said rates have increased in comparison to part (a). *LOOCV* (or  $CV_{loo}$ ) estimate is found to be approximately 0.07955.

# (1) Printing when actual and predicted values aren't same...

Observ. #23: Actual Value: 1, Predicted Value: 2 Observ. #25: Actual Value: 1, Predicted Value: 2 Observ. #37: Actual Value: 1, Predicted Value: 2 Observ. #38: Actual Value: 1, Predicted Value: 2 Observ. #61: Actual Value: 2, Predicted Value: 3

Observ. #62: Actual Value: 2, Predicted Value: 1 Observ. #66: Actual Value: 2, Predicted Value: 1 Observ. #71: Actual Value: 2, Predicted Value: 1

Observ. #72: Actual Value: 2, Predicted Value: 1 Observ. #83: Actual Value: 2, Predicted Value: 3

Observ. #123: Actual Value: 2, Predicted Value: 1 Observ. #130: Actual Value: 3, Predicted Value: 2 Observ. #134: Actual Value: 3, Predicted Value: 2

Observ. #138: Actual Value: 3, Predicted Value: 1

## (2) $CV_LOO = 0.07954545454545454$

	precision	recall	f1-score	support
1	0.90	0.93	0.92	59
2	0.91	0.90	0.91	71
3	0.96	0.93	0.95	46
avg / total	0.92	0.92	0.92	176

#### 1.3 1-(c)

Let us now perform the k-folds cross validation where k=4. To do so, I define the function  $cv_kf_mlogit$  for using the k-folds cross validation with multinomial logistic regression defined in the question. This will return the  $CV_{kf}$  value and the actual and predicted dependent variable values that will be used for calculating the classification report (and error rate, accordingly).

```
In [10]: def cv_kf_mlogit(Xvals, yvals, k=4, rtn_vectors=True):
    ## creating the splits
    kf = KFold(n_splits=k, shuffle=True, random_state=10)
    kf.get_n_splits(Xvals)
```

```
## creating the vector of KFolds MSE
MSE_vec_kf = np.zeros(k)
k_{ind} = int(0)
for train_idx, test_idx in kf.split(Xvals):
    ## splitting training and test sets
    X_tr_kf, X_test_kf = Xvals[train_idx], Xvals[test_idx]
    y_tr_kf, y_test_kf = yvals[train_idx], yvals[test_idx]
    ## making the prediction
    mlogit_kf = LR(random_state=0, solver='lbfgs',
                   multi_class='multinomial').fit(X_tr_kf, y_tr_kf)
    y_pred_kf = mlogit_kf.predict(X_test_kf)
    ## indicator function for categorical variables
    MSE_indicator_kf = (y_pred_kf != y_test_kf).mean()
    ## MSE vector for the specific k-fold
    MSE_vec_kf[k_ind] = MSE_indicator_kf
    print('Fold #{}:'.format(k_ind+1), 'MSE is {}'.format(MSE_indicator_kf))
    ## creating the vectors of actual and test yvals
    if k_ind == 0:
        y_act, y_pred = y_test_kf, y_pred_kf
    else:
        y_act = np.hstack((y_act, y_test_kf))
        y_pred = np.hstack((y_pred, y_pred_kf))
    k_ind += 1
cv_kf = MSE_vec_kf.mean()
print('----')
print('CV_KF is {}'.format(cv_kf))
if rtn_vectors:
   return cv_kf, y_act, y_pred
else:
   return cv_kf
```

Produced below are the results for the  $CV_{kf}$  as well as the cross validation table. The error rates for each category are 0.13, 0.09, and 0.04 for j=1,2 and 3 respectively. In comparison to part (a), error rate is approximately similar for j=1, but for j=2 and j=3 the said rates have increased. In comparison to part (b), error rates have increased for j=1, but remain similar for j=2 and j=3.  $CV_{kf}$  estimate (or k-fold estimate) is found to be approximately 0.09091.

```
print()
         print(classification_report(y_actual_kfolds, y_prediction_kfolds))
Fold #1: MSE is 0.1590909090909091
Fold #2: MSE is 0.11363636363636363
Fold #3: MSE is 0.0454545454545456
Fold #4: MSE is 0.0454545454545456
CV KF is 0.09090909090909091
                                             support
            precision
                          recall f1-score
          1
                  0.87
                            0.93
                                      0.90
                                                  59
                  0.91
                            0.87
                                      0.89
          2
                                                  71
          3
                  0.96
                            0.93
                                      0.95
                                                  46
```

## 2 Problem 2

avg / total

0.91

## 2.1 2-(a)

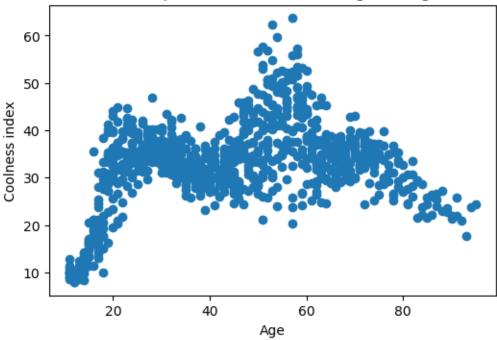
In the below code chunk, I have read in the CoolIndex.txt data and have successfully produced the scatterplot.

0.91

176

0.91





## 2.2 2-(b)

For the OLS regression (to create the stepwise function mentioned in the question), let us first create the dummy variables for each age bin with the age  $\in$  [11, 22) being the base case.

We now can create the representations for X (regressors) and y (dependent variable) for the said OLS regression.

```
In [14]: X = cool[['22-40', '40-59', '59-77', '77-95']]
    X = sm.add_constant(X)
    y = cool[[1]]
```

In the part below, the OLS regression results are produced.

#### OLS Regression Results

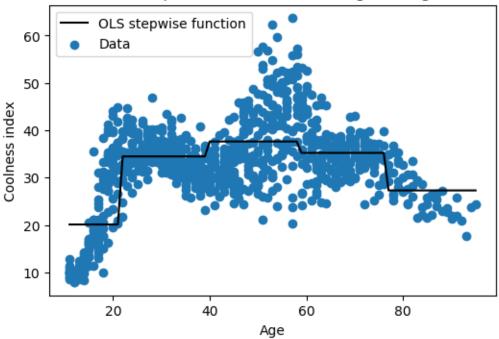
Dep. Variable: Model: Method: Date: Time: No. Observations:	1 OLS Least Squares Tue, 26 Feb 2019 21:33:52 956	R-squared: Adj. R-squared: F-statistic: Prob (F-statistic) Log-Likelihood: AIC:	0.429 0.427 178.7 : 3.73e-114 -3214.5 6439.
Df Residuals:	951	BIC:	6463.
Df Model:	4		
Covariance Type:	nonrobust		
CO6	f std err	t P> t	[0.025 0.975]
const 20.102	25 0.562 3	5.746 0.000	18.999 21.206
22-40 14.373	3 0.708 2	0.287 0.000	12.983 15.764
40-59 17.532	26 0.704 2	4.899 0.000	16.151 18.915
59-77 15.122	9 0.743 2	0.356 0.000	13.665 16.581
77-95 7.193	1.092	6.590 0.000	5.052 9.336
Omnibus: Prob(Omnibus): Skew: Kurtosis:	80.102 0.000 0.714 3.719	Durbin-Watson: Jarque-Bera (JB): Prob(JB): Cond. No.	1.236 101.718 8.17e-23 6.80

#### Warnings:

[1] Standard Errors assume that the covariance matrix of the errors is correctly specified.

We can now use the above OLS regression results to fit the values of *X* and produce the stepwise function as shown in the graph below. Since a 73-year old falls under the category of age bin 59-77, we add the constant term 20.1025 and the coefficient for the said age bin to find out that, under this model, a 73-year old's predicted coolness is approximately 35.2254.





#### $2.3 \quad 2-(c)$

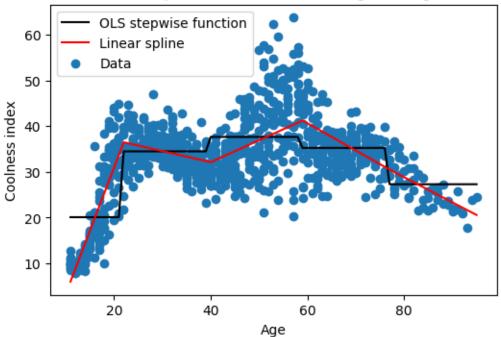
Let us first prepare the set up for the linear spline. Note that to use the function lsq (which is the shorthand I use to refer to LSQUnivariateSpline throughout this problem set), we need to have a vector of ages where the elements are *strictly* increasing. Therefore let us take the mean of coolness for each age that appears in the dataset, and use this to create the spline.

Below code chunk executes the linear spline.

```
In [20]: spl = lsq(x=age, y=cool_means, t=knots, k=1)
```

Below graph showcases the data points, linear spline and the OLS stepwise function created in part (b). It is visible that, while not very smooth, linear spline resembles the data more so than the OLS stepwise function.

## Scatterplot of Coolness Index against Age



Finally, the predicted coolness from the linear spline is approximately 32.8678 for a 73-year-old.

```
In [22]: spl(73)
Out[22]: array(32.86784862349653)
```

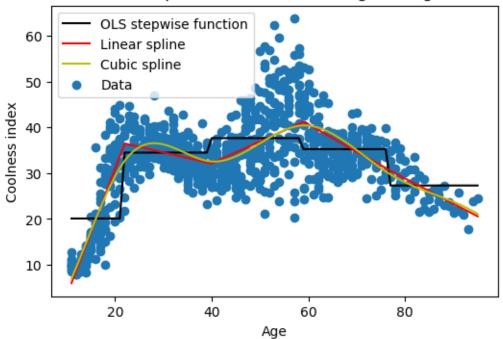
### 2.4 2-(d)

Below code chunk executes the cubic spline.

```
In [23]: spl_cube = lsq(x=age, y=cool_means, t=knots, k=3)
```

Below graph showcases the data points, cubic spline, linear spline from part (c) and the OLS stepwise function created in part (b). It is visible that cubic spline resembles the data more smoothly than the other two.

## Scatterplot of Coolness Index against Age



Finally, the predicted coolness from the cubic spline is approximately 32.6423 for a 73-year-old.

```
In [25]: spl_cube(73)
Out[25]: array(32.642301066279764)
```