In [2]: #Import Dataset csvData = pd.read_csv('diabetes.csv') csvData

Out[2]:

	Pregnancies	Glucose	BloodPressure	SkinThickness	Insulin	BMI	DiabetesPedigreeFunction
0	6	148	72	35	0	33.6	0.62
1	1	85	66	29	0	26.6	0.38
2	8	183	64	0	0	23.3	0.67
3	1	89	66	23	94	28.1	0.16
4	0	137	40	35	168	43.1	2.28
763	10	101	76	48	180	32.9	0.17
764	2	122	70	27	0	36.8	0.34
765	5	121	72	23	112	26.2	0.24
766	1	126	60	0	0	30.1	0.34
767	1	93	70	31	0	30.4	0.3

768 rows × 9 columns

```
In [3]: #Sort Dataset
x = csvData.iloc[:,0:-1].values
y = csvData.iloc[:,-1].values
```

```
In [4]: #Train-Test Split
    from sklearn.model_selection import train_test_split
    xt, xv, yt, yv = train_test_split(x, y, train_size = 0.8, test_size = 0.2, ran
```

```
In [5]: #Clean the Dataset
    from sklearn.preprocessing import MinMaxScaler, StandardScaler
    # scaler = StandardScaler() #MinMaxScaler gave better results here
    scaler = MinMaxScaler()
    xt = scaler.fit_transform(xt)
    xv = scaler.fit_transform(xv)
```

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```
In [6]: #Perform the Training
        from sklearn.linear_model import LogisticRegression
        training_montage = LogisticRegression(random_state=1337)
        training_montage.fit(xt,yt);
In [7]: #Test the Model
        p = training_montage.predict(xv)
In [8]: #Evaluate the model metrics
        from sklearn import metrics
        print("Model Accuracy: {:.3f}%".format(metrics.accuracy_score(yv,p)*100))
        print("Model Precision: {:.3f}%".format(metrics.precision_score(yv,p)*100))
                              {:.3f}%".format(metrics.recall_score(yv,p)*100))
        print("Model Recall:
        Model Accuracy:
                         77.273%
        Model Precision: 53.488%
        Model Recall:
                         60.526%
In [9]: #Analyze using the Confusion Matrix
        from sklearn.metrics import confusion_matrix
        import seaborn as sns
        classes = ['Not Diabetes', 'Diabetes']
        figure, axis = plt.subplots()
        ticks = np.arange(len(classes))
```

sns.heatmap(pd.DataFrame(confusion_matrix(yv, p)), annot=True, cmap="Greens",

Confusion Matrix

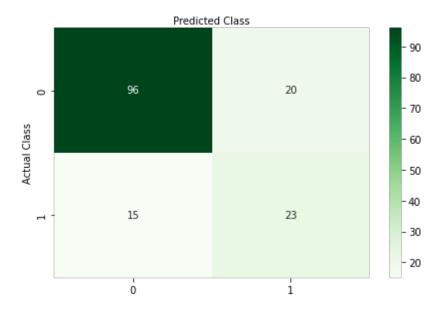
plt.xticks(ticks, classes)
plt.yticks(ticks, classes)

plt.ylabel('Actual Class')
plt.xlabel('Predicted Class');

plt.tight_layout()

axis.xaxis.set_label_position("top")

plt.title('Confusion Matrix', y=1.1)



In []:

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In [2]: #Import Dataset csvData = pd.read_csv('diabetes.csv') csvData

Out[2]:		Pregnancies	Glucose	BloodPressure	SkinThickness	Insulin	ВМІ	DiabetesPedigreeFunction
_	0	6	148	72	35	0	33.6	0.62
	1	1	85	66	29	0	26.6	0.3!
	2	8	183	64	0	0	23.3	0.6
	3	1	89	66	23	94	28.1	0.16
	4	0	137	40	35	168	43.1	2.28
,	763	10	101	76	48	180	32.9	0.17
	764	2	122	70	27	0	36.8	0.34
,	765	5	121	72	23	112	26.2	0.24
,	766	1	126	60	0	0	30.1	0.34
ı	767	1	93	70	31	0	30.4	0.3

768 rows × 9 columns

```
In [3]: #Sort Dataset
x = csvData.iloc[:,0:-1].values
y = csvData.iloc[:,-1].values
```

```
In [18]: #Clean the Dataset
    from sklearn.preprocessing import MinMaxScaler, StandardScaler
    scaler = StandardScaler()
    # scaler = MinMaxScaler() #StandardScaler gave better results here
    x = scaler.fit_transform(x)
```

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```
In [19]: #Perform the Training with K=5
         from sklearn.model_selection import KFold
         from sklearn.linear_model import LogisticRegression
         from sklearn.model_selection import cross_val_score
         kcup = KFold(n_splits=5, random_state=1337, shuffle=True)
         model = LogisticRegression(random_state=1337)
         results = cross_val_score(model,x,y,cv=kcup)
         print("K=5 | Accuracy: {:.3f}% ({:.3f}%)".format(results.mean()*100, results.s
         K=5 | Accuracy: 77.734% (2.167%)
In [17]: #Perform the Training with K=10
         kcup = KFold(n_splits=10, random_state=1337, shuffle=True)
         model = LogisticRegression(random_state=1337)
         results = cross_val_score(model,x,y,cv=kcup)
         print("K=10 | Accuracy: {:.3f}% ({:.3f}%)".format(results.mean()*100, results.
         K=10 | Accuracy: 76.811% (3.108%)
 In [ ]:
```

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In [2]: #Load and Build the Dataset

from sklearn.datasets import load_breast_cancer

loaded = load_breast_cancer()

labels = np.reshape(loaded.target, (len(loaded.target),1))

inputs = pd.DataFrame(loaded.data)

names = np.append(loaded.feature_names, 'label')

dataset = pd.DataFrame(np.concatenate([inputs,labels],axis=1))

dataset.columns = names

dataset

Out[2]:

	mean radius	mean texture	mean perimeter	mean area	mean smoothness	mean compactness	mean concavity	mean concave points	mea symmetr
0	17.99	10.38	122.80	1001.0	0.11840	0.27760	0.30010	0.14710	0.241
1	20.57	17.77	132.90	1326.0	0.08474	0.07864	0.08690	0.07017	0.181
2	19.69	21.25	130.00	1203.0	0.10960	0.15990	0.19740	0.12790	0.206
3	11.42	20.38	77.58	386.1	0.14250	0.28390	0.24140	0.10520	0.259
4	20.29	14.34	135.10	1297.0	0.10030	0.13280	0.19800	0.10430	0.180
564	21.56	22.39	142.00	1479.0	0.11100	0.11590	0.24390	0.13890	0.172
565	20.13	28.25	131.20	1261.0	0.09780	0.10340	0.14400	0.09791	0.175
566	16.60	28.08	108.30	858.1	0.08455	0.10230	0.09251	0.05302	0.159
567	20.60	29.33	140.10	1265.0	0.11780	0.27700	0.35140	0.15200	0.239
568	7.76	24.54	47.92	181.0	0.05263	0.04362	0.00000	0.00000	0.158

569 rows × 31 columns

```
In [3]: #Sort Dataset
x = dataset.iloc[:,0:-1].values
y = dataset.iloc[:,-1].values
```

```
In [4]: #Train-Test Split
    from sklearn.model_selection import train_test_split
    xt, xv, yt, yv = train_test_split(x, y, train_size = 0.8, test_size = 0.2, ran
```

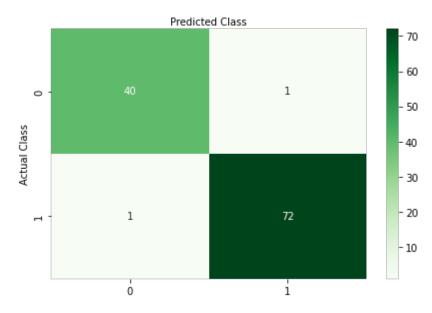
```
In [5]: #Clean the Dataset
        from sklearn.preprocessing import MinMaxScaler, StandardScaler
        scaler = StandardScaler()
         # scaler = MinMaxScaler() #StandardScaler gave better results here
        xt = scaler.fit_transform(xt)
        xv = scaler.fit_transform(xv)
In [6]: #Perform the Training
        from sklearn.linear_model import LogisticRegression
        training_montage = LogisticRegression(random_state=1337)
        training_montage.fit(xt,yt);
In [7]: #Test the Model
        p = training_montage.predict(xv)
In [8]: #Evaluate the model metrics
        from sklearn import metrics
        print("Model Accuracy: {:.3f}%".format(metrics.accuracy_score(yv,p)*100))
        print("Model Precision: {:.3f}%".format(metrics.precision_score(yv,p)*100))
        print("Model Recall: {:.3f}%".format(metrics.recall_score(yv,p)*100))
        Model Accuracy: 98.246%
```

Model Precision: 98.630% Model Recall: 98.630%

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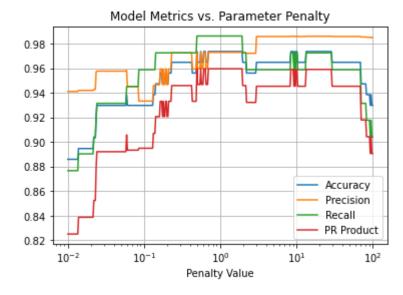
```
In [9]: #Analyze using the Confusion Matrix
    from sklearn.metrics import confusion_matrix
    import seaborn as sns
    classes = ['Not Diabetes','Diabetes']
    figure, axis = plt.subplots()
    ticks = np.arange(len(classes))
    plt.xticks(ticks, classes)
    plt.yticks(ticks, classes)
    sns.heatmap(pd.DataFrame(confusion_matrix(yv, p)), annot=True, cmap="Greens",
    axis.xaxis.set_label_position("top")
    plt.tight_layout()
    plt.title('Confusion Matrix', y=1.1)
    plt.ylabel('Actual Class')
    plt.xlabel('Predicted Class');
```

Confusion Matrix



```
In [10]: #Reevaluate using a variety of weight penalties
lambdas = np.logspace(-2,2,num=400)
acc_log = []
prc_log = []
rec_log = []
for lam in lambdas:
    model = LogisticRegression(penalty='l1',C=lam,solver='liblinear',random_st model.fit(xt,yt)
    p = model.predict(xv)
    rec_log.append(metrics.recall_score(yv,p))
    prc_log.append(metrics.precision_score(yv,p))
acc_log.append(metrics.accuracy_score(yv,p))
```

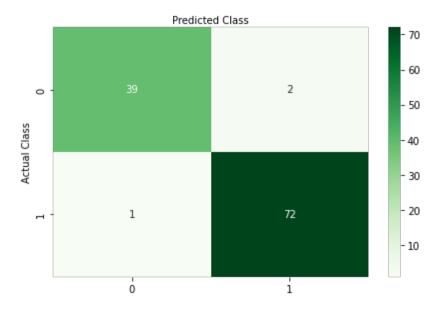
```
In [11]: #Plot the results
    plt.semilogx(lambdas,acc_log,label='Accuracy')
    plt.semilogx(lambdas,prc_log,label='Precision')
    plt.semilogx(lambdas,rec_log,label='Recall')
    PRval = np.multiply(prc_log,rec_log)
    plt.semilogx(lambdas,PRval,label='PR Product')
    plt.grid()
    plt.xlabel('Penalty Value')
    plt.title('Model Metrics vs. Parameter Penalty')
    plt.legend();
```



```
In [12]: #According to the plot, lambda = 1 is about the best it gets
         model = LogisticRegression(penalty='l1',C=1,solver='liblinear',random_state=13
         model.fit(xt,yt)
         p = model.predict(xv)
         print("Model Accuracy: {:.3f}%".format(metrics.accuracy_score(yv,p)*100))
         print("Model Precision: {:.3f}%".format(metrics.precision_score(yv,p)*100))
         print("Model Recall:
                                {:.3f}%".format(metrics.recall_score(yv,p)*100))
         classes = ['Not Diabetes', 'Diabetes']
         figure, axis = plt.subplots()
         ticks = np.arange(len(classes))
         plt.xticks(ticks, classes)
         plt.yticks(ticks, classes)
         sns.heatmap(pd.DataFrame(confusion_matrix(yv, p)), annot=True, cmap="Greens",
         axis.xaxis.set_label_position("top")
         plt.tight_layout()
         plt.title('Confusion Matrix', y=1.1)
         plt.ylabel('Actual Class')
         plt.xlabel('Predicted Class');
```

Model Accuracy: 97.368% Model Precision: 97.297% Model Recall: 98.630%

Confusion Matrix



In []:

```
In [1]:
    """
    ECGR 5105 - Intro to Machine Learning
    Homework 2 - Part 4
    Phillip Harmon
    """
    import numpy as np
    import matplotlib.pyplot as plt
    import pandas as pd
```

In [2]: #Load and Build the Dataset

from sklearn.datasets import load_breast_cancer

loaded = load_breast_cancer()

labels = np.reshape(loaded.target, (len(loaded.target),1))

inputs = pd.DataFrame(loaded.data)

names = np.append(loaded.feature_names, 'label')

dataset = pd.DataFrame(np.concatenate([inputs,labels],axis=1))

dataset.columns = names

dataset

Out[2]:

		mean radius	mean texture	mean perimeter	mean area	mean smoothness	mean compactness	mean concavity	mean concave points	mea symmeti
	0	17.99	10.38	122.80	1001.0	0.11840	0.27760	0.30010	0.14710	0.241
	1	20.57	17.77	132.90	1326.0	0.08474	0.07864	0.08690	0.07017	0.181
	2	19.69	21.25	130.00	1203.0	0.10960	0.15990	0.19740	0.12790	0.206
	3	11.42	20.38	77.58	386.1	0.14250	0.28390	0.24140	0.10520	0.259
	4	20.29	14.34	135.10	1297.0	0.10030	0.13280	0.19800	0.10430	0.180
56	64	21.56	22.39	142.00	1479.0	0.11100	0.11590	0.24390	0.13890	0.172
56	65	20.13	28.25	131.20	1261.0	0.09780	0.10340	0.14400	0.09791	0.175
56	66	16.60	28.08	108.30	858.1	0.08455	0.10230	0.09251	0.05302	0.159
56	67	20.60	29.33	140.10	1265.0	0.11780	0.27700	0.35140	0.15200	0.239
56	88	7.76	24.54	47.92	181.0	0.05263	0.04362	0.00000	0.00000	0.158

569 rows × 31 columns

In [3]: #Sort Dataset x = dataset.iloc[:,0:-1].values y = dataset.iloc[:,-1].values

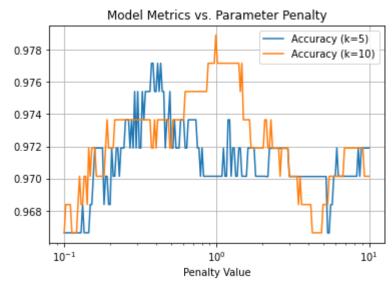
```
In [4]: #Clean the Dataset
    from sklearn.preprocessing import MinMaxScaler, StandardScaler
    scaler = StandardScaler()
    # scaler = MinMaxScaler() #StandardScaler gave better results here
    x = scaler.fit_transform(x)
```

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```
In [5]: #Perform the Training with K=5
        from sklearn.model_selection import KFold
        from sklearn.linear_model import LogisticRegression
        from sklearn.model_selection import cross_val_score
        kcup = KFold(n_splits=5, random_state=1337, shuffle=True)
        model = LogisticRegression(random_state=1337)
        results = cross_val_score(model,x,y,cv=kcup)
        print("K=5 | Accuracy: {:.3f}% ({:.3f}%)".format(results.mean()*100, results.s
        K=5 | Accuracy: 97.539% (1.290%)
In [6]: #Perform the Training with K=10
        kcup = KFold(n_splits=10, random_state=1337, shuffle=True)
        model = LogisticRegression(random_state=1337)
        results = cross_val_score(model,x,y,cv=kcup)
        print("K=10 | Accuracy: {:.3f}% ({:.3f}%)".format(results.mean()*100, results.
        K=10 | Accuracy: 97.716% (1.122%)
In [7]: #Reevaluate using a variety of weight penalties
        lambdas = np.logspace(-1,1,num=200)
        k5_acc_log = []
        kcup = KFold(n_splits=5, random_state=1337, shuffle=True)
        for lam in lambdas:
            model = LogisticRegression(penalty='l1',C=lam,solver='liblinear',random_st
            results = cross_val_score(model,x,y,cv=kcup)
            k5_acc_log.append(results.mean())
        k10_acc_log = []
        kcup = KFold(n_splits=10, random_state=1337, shuffle=True)
        for lam in lambdas:
            model = LogisticRegression(penalty='l1',C=lam,solver='liblinear',random_st
            results = cross_val_score(model,x,y,cv=kcup)
            k10_acc_log.append(results.mean())
```

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```
In [8]: #Plot the results
plt.semilogx(lambdas,k5_acc_log,label='Accuracy (k=5)')
plt.semilogx(lambdas,k10_acc_log,label='Accuracy (k=10)')
plt.grid()
plt.xlabel('Penalty Value')
plt.title('Model Metrics vs. Parameter Penalty')
plt.legend();
```



```
In [9]: #Best k=5 weight is about 0.35
kcup = KFold(n_splits=5, random_state=1337, shuffle=True)
model = LogisticRegression(penalty='l1',C=0.35,solver='liblinear',random_state
results = cross_val_score(model,x,y,cv=kcup)
print("K=5 | Accuracy: {:.3f}% ({:.3f}%)".format(results.mean()*100, results.s)
K=5 | Accuracy: 97.541% (0.653%)
```

```
In [10]: #Best k=10 weight is about 1
kcup = KFold(n_splits=10, random_state=1337, shuffle=True)
model = LogisticRegression(penalty='l1',C=1,solver='liblinear',random_state=13
results = cross_val_score(model,x,y,cv=kcup)
print("K=10 | Accuracy: {:.3f}% ({:.3f}%)".format(results.mean()*100, results.
```

K=10 | Accuracy: 97.892% (1.312%)

In []: