Assignment 02

Name:- Sk Fardeen Hossain

Roll No.:- 2021CSB023

G-Suite Id:- 2021csb023.sk@students.iiests.ac.in

Department:- Computer Science and Technology

Question 01

Download Cancer Wisconsin (Diagnostic) Data Set (already in the needed format). The dataset is used to recognize 2 types of cancer to be predicted (benign or malignant).

```
In [54]: ## Install and import the pandas library
         !pip install pandas
        Requirement already satisfied: pandas in /home/fardeen/Desktop/MachineLearningLab/venv/lib/python3.12/site-packages (2.2.2)
        Requirement already satisfied: numpy>=1.26.0 in /home/fardeen/Desktop/MachineLearningLab/venv/lib/python3.12/site-packages (from pandas) (2.0.1)
        Requirement already satisfied: python-dateutil>=2.8.2 in /home/fardeen/Desktop/MachineLearningLab/venv/lib/python3.12/site-packages (from pandas)
        (2.9.0.post0)
        Requirement already satisfied: pytz>=2020.1 in /home/fardeen/Desktop/MachineLearningLab/venv/lib/python3.12/site-packages (from pandas) (2024.1)
        Requirement already satisfied: tzdata>=2022.7 in /home/fardeen/Desktop/MachineLearningLab/venv/lib/python3.12/site-packages (from pandas) (2024.1)
        Requirement already satisfied: six>=1.5 in /home/fardeen/Desktop/MachineLearningLab/venv/lib/python3.12/site-packages (from python-dateutil>=2.8.2->p
        andas) (1.16.0)
In [55]: import pandas as pd
         DATASET PATH = "../ML DRIVE/Assignment02/data.csv"
         cancer dataframe = pd.read csv(DATASET PATH)
         print(cancer dataframe.columns)
         cancer dataframe.head()
        Index(['id', 'diagnosis', 'radius mean', 'texture mean', 'perimeter mean',
               'area mean', 'smoothness mean', 'compactness mean', 'concavity mean',
               'concave points mean', 'symmetry mean', 'fractal dimension mean',
               'radius se', 'texture se', 'perimeter se', 'area se', 'smoothness se',
               'compactness se', 'concavity se', 'concave points se', 'symmetry se',
               'fractal dimension se', 'radius worst', 'texture worst',
               'perimeter worst', 'area worst', 'smoothness worst',
               'compactness worst', 'concavity worst', 'concave points worst',
               'symmetry worst', 'fractal dimension worst', 'Unnamed: 32'],
              dtype='object')
```

Out[55]:		id	diagnosis	radius_mean	texture_mean	perimeter_mean	area_mean	smoothness_mean	compactness_mean	concavity_mean	concave points_mean	•••	texture_worst per	·im
	0	842302	М	17.99	10.38	122.80	1001.0	0.11840	0.27760	0.3001	0.14710		17.33	
	1	842517	М	20.57	17.77	132.90	1326.0	0.08474	0.07864	0.0869	0.07017		23.41	
	2	84300903	М	19.69	21.25	130.00	1203.0	0.10960	0.15990	0.1974	0.12790		25.53	
	3	84348301	М	11.42	20.38	77.58	386.1	0.14250	0.28390	0.2414	0.10520		26.50	
	4	84358402	М	20.29	14.34	135.10	1297.0	0.10030	0.13280	0.1980	0.10430		16.67	

5 rows × 33 columns

In [56]: ## Pre-processing the data
 cancer_dataframe = cancer_dataframe.drop('Unnamed: 32',axis=1) ### Dropping the column with null values
 cancer_dataframe = cancer_dataframe.drop('id',axis=1) ### Dropping the id column
 cancer_dataframe

Out[56]:	diag	nosis	radius_mean	texture_mean	perimeter_mean	area_mean	smoothness_mean	compactness_mean	concavity_mean	concave points_mean	symmetry_mean	radius_worst
	0	М	17.99	10.38	122.80	1001.0	0.11840	0.27760	0.30010	0.14710	0.2419	25.380
	1	М	20.57	17.77	132.90	1326.0	0.08474	0.07864	0.08690	0.07017	0.1812	24.990
	2	М	19.69	21.25	130.00	1203.0	0.10960	0.15990	0.19740	0.12790	0.2069	23.570
	3	М	11.42	20.38	77.58	386.1	0.14250	0.28390	0.24140	0.10520	0.2597	14.910
	4	М	20.29	14.34	135.10	1297.0	0.10030	0.13280	0.19800	0.10430	0.1809	22.540
	•••			•••								
5	564	М	21.56	22.39	142.00	1479.0	0.11100	0.11590	0.24390	0.13890	0.1726	25.450
į	565	М	20.13	28.25	131.20	1261.0	0.09780	0.10340	0.14400	0.09791	0.1752	23.690
į	566	М	16.60	28.08	108.30	858.1	0.08455	0.10230	0.09251	0.05302	0.1590	18.980
į	567	М	20.60	29.33	140.10	1265.0	0.11780	0.27700	0.35140	0.15200	0.2397	25.740
į	568	В	7.76	24.54	47.92	181.0	0.05263	0.04362	0.00000	0.00000	0.1587	9.456

569 rows × 31 columns

Question 02

Implement Logistic Regression using scikit-learn package.

- a. Use 'newton-cg', 'lbfgs', 'liblinear' solver for the regression model.
- b. For each solver use 'l1', 'l2', 'none' penalty to train the model.
- c. Split the dataset in to 80:10:10 percent (train: validation: test) (use seed = 5 for splitting).

- d. Train the model initially with 80% training data and create a table for the coefficients of all the features.
- e. Fine-tune the model with the remaining validation partition of the dataset (consisting of 10% of the original dataset) and create a table for the updated coefficients of all the features.

```
In [57]: ## Install the scikit-learn package
         !pip install scikit-learn
        Requirement already satisfied: scikit-learn in /home/fardeen/Desktop/MachineLearningLab/venv/lib/python3.12/site-packages (1.5.1)
        Requirement already satisfied: numpy>=1.19.5 in /home/fardeen/Desktop/MachineLearningLab/venv/lib/python3.12/site-packages (from scikit-learn)
        (2.0.1)
        Requirement already satisfied: scipy>=1.6.0 in /home/fardeen/Desktop/MachineLearningLab/venv/lib/python3.12/site-packages (from scikit-learn) (1.1
        4.0)
        Requirement already satisfied: joblib>=1.2.0 in /home/fardeen/Desktop/MachineLearningLab/venv/lib/python3.12/site-packages (from scikit-learn)
        Requirement already satisfied: threadpoolctl>=3.1.0 in /home/fardeen/Desktop/MachineLearningLab/venv/lib/python3.12/site-packages (from scikit-learn)
        (3.5.0)
In [58]: ## Separating the class variable from the dataset
         import pandas as pd
         y = cancer dataframe[['diagnosis']]
         print(y.head()) # --> M implies Malignant B implies Benign
         print(y.value counts())
          diagnosis
        0
                  Μ
                  Μ
        2
                  Μ
        3
                  Μ
                  М
        diagnosis
                     357
                     212
        Μ
        Name: count, dtype: int64
In [59]: ## Rest of the features will be independent variables
         X = cancer dataframe.drop('diagnosis',axis=1)
         X.head()
```

Out[59]:	radius_mean	texture_mean	perimeter_mean	area_mean	smoothness_mean	compactness_mean	concavity_mean	concave points_mean	symmetry_mean	fractal_dimension_mean	•••
	0 17.99	10.38	122.80	1001.0	0.11840	0.27760	0.3001	0.14710	0.2419	0.07871	
	1 20.57	17.77	132.90	1326.0	0.08474	0.07864	0.0869	0.07017	0.1812	0.05667	
	2 19.69	21.25	130.00	1203.0	0.10960	0.15990	0.1974	0.12790	0.2069	0.05999	
	3 11.42	20.38	77.58	386.1	0.14250	0.28390	0.2414	0.10520	0.2597	0.09744	
	4 20.29	14.34	135.10	1297.0	0.10030	0.13280	0.1980	0.10430	0.1809	0.05883	

5 rows × 30 columns

Standardization of feature values to keep less bias on feature value thus maintaining uniformity and treating features on an equal scale

Out[60]:	radius_mea		texture_mean	perimeter_mean	area_mean	smoothness_mean	compactness_mean	concavity_mean	concave points_mean	symmetry_mean	fractal_dimension_mean	
	0	0.521037	0.022658	0.545989	0.363733	0.593753	0.792037	0.703140	0.731113	0.686364	0.605518	
	1	0.643144	0.272574	0.615783	0.501591	0.289880	0.181768	0.203608	0.348757	0.379798	0.141323	
	2	0.601496	0.390260	0.595743	0.449417	0.514309	0.431017	0.462512	0.635686	0.509596	0.211247	
	3	0.210090	0.360839	0.233501	0.102906	0.811321	0.811361	0.565604	0.522863	0.776263	1.000000	
	4	0.629893	0.156578	0.630986	0.489290	0.430351	0.347893	0.463918	0.518390	0.378283	0.186816	

5 rows × 30 columns

Splitting the dataset into training, testing and validation data

```
In [61]: from sklearn.model_selection import train_test_split

X_train, X_rem, y_train, y_rem = train_test_split(X, y, test_size=0.2, random_state=5)

X_valid, X_test, y_valid, y_test = train_test_split(X_rem, y_rem, train_size=0.5, random_state=5)

# print(X_train.info())
# print(X_train.info())
```

```
# print(X valid.info())
         # print(y_train.info())
         # print(y test.info())
         # print(y valid.info())
In [70]: ## Utility Function to create and vary the Logistic Regression model
         import numpy as np
         from sklearn.linear model import LogisticRegression
         from sklearn.metrics import accuracy score, f1 score, recall score, precision score
             @brief:- Function that creates a logisitic regression model and returns the accuracy and coefficient of all the features
             @params:- X train -> training data independent variable
                       y_train -> training data class variable
                       X valid -> validation data independent variable
                       y valid -> validation data class variable
                       solver -> solver used for fitting the model and increase convergence speed
                       penalty -> l1/l2 regularization to prevent over-fitting
                       max iter -> Max iterations for fitting the model
             @return:- Model coefficients(before and after finetuning), accuracy score, precision score, recall score, flscore
         1.1.1
         def utilLogRegInfo(
             X train:
                              'pd.Dataframe',
             y train:
                              'pd.Dataframe',
             X valid:
                              'pd.Dataframe',
             y valid:
                              'pd.Dataframe',
             X test:
                              'pd.Dataframe',
             y test:
                              'pd.Dataframe',
             solver:
                              'str',
             penalty:
                              'str'='none',
             inv reg strength:'float'=1.0,
             max iter:
                              'int' =10000) -> 'list':
             if penalty=='none':
                 model = LogisticRegression(
                     solver = solver,
                     max iter= max iter,
                     penalty=None,
                     C=inv reg strength
                 ).fit(X_train,y_train['diagnosis'])
                 coef_train = model.coef_
                 y pred before finetune = model.predict(X valid)
                 accuracy before fine tune = accuracy score(y valid,y pred before finetune)
                 precision_before_fine_tune = precision_score(y_valid,y_pred_before_finetune,average='macro').item()
                 recall before fine tune = recall score(y valid,y pred before finetune,average='macro').item()
                 Flscore before fine tune = fl score(y valid,y pred before finetune,average='macro').item()
```

print(X test.info())

```
model.fit(np.vstack([X train, X valid]), np.vstack([y train, y valid]))
                 coef finetuned = model.coef
                 y pred after finetune = model.predict(X test)
                 accuracy after fine tune = accuracy score(y test,y pred after finetune)
                 precision after fine tune = precision score(y test,y pred after finetune,average='macro').item()
                 recall after fine tune = recall score(y test,y pred after finetune,average='macro').item()
                 Flscore_after_fine_tune = fl_score(y_test,y_pred_after_finetune,average='macro').item()
             else:
                 model = LogisticRegression(
                     solver = solver,
                     penalty = penalty,
                     max iter= max iter,
                     C=inv reg strength
                 ).fit(X train,y train['diagnosis'])
                 coef train = model.coef
                 y pred before finetune = model.predict(X valid)
                 accuracy_before_fine_tune = accuracy_score(y_valid,y_pred_before_finetune)
                 precision before fine tune = precision score(y valid,y pred before finetune,average='macro').item()
                 recall before fine tune = recall score(y valid,y pred before finetune,average='macro').item()
                 Flscore before fine tune = fl score(y valid,y pred before finetune,average='macro').item()
                 model.fit(np.vstack([X train, X valid]), np.vstack([y train, y valid]))
                 coef finetuned = model.coef
                 y pred after finetune = model.predict(X test)
                 accuracy after fine tune = accuracy score(y test,y pred after finetune)
                 precision after fine tune = precision score(y test,y pred after finetune,average='macro').item()
                 recall_after_fine_tune = recall_score(y_test,y_pred_after_finetune,average='macro').item()
                 Flscore after fine tune = fl score(y test,y pred after finetune,average='macro').item()
             return [solver, penalty, inv reg strength, accuracy before fine tune, precision before fine tune, recall before fine tune,
                     Flscore before fine tune,accuracy after fine tune,precision after fine tune,recall after fine tune,Flscore after fine tune]+\
                     coef_train.tolist()[0] + coef_finetuned.tolist()[0]
In [71]: newton cg reg l2 = utilLogRegInfo(X train,y train,X valid,y valid,X test,y test,solver='newton-cg',penalty='l2')
         newton cg reg none = utilLogRegInfo(X train,y train,X valid,y valid,X test,y test,solver='newton-cg')
         lbfgs_reg_l2 = utilLogRegInfo(X_train,y_train,X_valid,y_valid,X_test,y_test,solver='lbfgs',penalty='l2')
         lbfqs reg none = utilLogRegInfo(X train,y train,X valid,y valid,X test,y test,solver='lbfqs')
         liblinear reg l1 = utilLogRegInfo(X train,y train,X valid,y valid,X test,y test,solver='liblinear',penalty='l1')
         liblinear reg l2 = utilLogRegInfo(X train,y train,X valid,y valid,X test,y test,solver='liblinear',penalty='l2')
         solvers report = pd.DataFrame(
```

columns= ['solver', 'penalty']+[f"{col} coeff." for col in X_train.columns] + [f"{col} finetuned coeff. " for col in X_train.columns],

data = [newton cg reg l2[:2]+newton cg reg l2[11:],newton cg reg none[:2]+newton cg reg none[11:],

lbfgs reg l2[:2]+lbfgs reg l2[11:],lbfgs reg none[:2]+lbfgs reg none[11:],

liblinear_reg_l1[:2]+liblinear_reg_l1[11:],
liblinear_reg_l2[:2]+liblinear_reg_l2[11:]]

solvers_report

O + I	F 7 1 1	
UHIT	/	

	solver	penalty	radius_mean coeff.	texture_mean coeff.	perimeter_mean coeff.	area_mean coeff.	smoothness_mean coeff.	compactness_mean coeff.	concavity_mean coeff.	concave points_mean coeff.	•••	finetuned coeff.	finetu co
0	newton- cg	l2	1.763553	1.586001	1.732097	1.497118	0.581991	0.393523	1.438618	2.084012		2.344268	2.345
1	newton- cg	none	-3.832666	-16.951656	-33.221431	-32.735619	22.313882	-74.121615	77.898319	58.902442		-112.911015	77.34€
2	lbfgs	l2	1.760413	1.587441	1.728953	1.498279	0.581363	0.391197	1.437480	2.088940		2.347741	2.342
3	lbfgs	none	-107.628970	-251.288585	-134.686577	-6.218266	54.224016	-311.087162	279.523186	451.847827		43.177874	50.340
4	liblinear	l1	0.000000	0.000000	0.000000	0.000000	0.000000	0.000000	0.000000	4.964401		13.411096	5.432
5	liblinear	l2	0.732458	0.838540	0.795956	1.056228	-0.396246	0.377031	1.863024	2.470633		1.729162	1.622

6 rows × 62 columns

Question 03

For every solver vary the 'l1' penalty over the range (0.1, 0.25, 0.75, 0.9) and compare the coefficients of the features.

```
In [73]: ## Only liblinear and saga solver supports l1 penalty
         # liblinear
         liblinear l1 penalty = pd.DataFrame(
             columns = ['solver','penalty','inverse_regularization_strength'] + [f"{col}_coeff. " for col in X_train.columns] +
                         [f"{col} finetuned coeff. " for col in X_train.columns],
             data = [ utilLogRegInfo(X train,y train,X valid,y valid,X test,y test, solver='liblinear',inv reg strength=penalty,penalty='l1')[:3]
                     + utilLogRegInfo(X_train,y_train,X_valid,y_valid,X_test,y_test, solver='liblinear',inv_reg_strength=penalty,penalty='l1')[11:]
                     for penalty in [0.1,0.25,0.75,0.9]
         saga_l1_penalty = pd.DataFrame(
             columns = ['solver', 'penalty', 'inverse_regularization_strength'] + [f"{col}_coeff. " for col in X_train.columns] +
                         [f"{col} finetuned coeff. " for col in X_train.columns],
             data = [ utilLogRegInfo(X_train,y_train,X_valid,y_valid,X_test,y_test,solver='saga',inv_reg_strength=penalty,penalty='l1')[:3] +
                      utilLogRegInfo(X_train,y_train,X_valid,y_valid,X_test,y_test, solver='saga',inv_reg_strength=penalty,penalty='l1')[11:]
                     for penalty in [0.1,0.25,0.75,0.9]
         frames = [liblinear_l1_penalty, saga_l1_penalty]
```

```
l1_penalty_report = pd.concat(frames)
l1_penalty_report
```

solver penalty inverse regularization strength radius mean coeff. texture mean coeff. perimeter mean coeff. area mean coeff. smoothness mean coeff. compactness m

Out[731:

0 liblinear	l1	0.10	0.0	0.0	0.0	0.0	0.0
1 liblinear	l1	0.25	0.0	0.0	0.0	0.0	0.0
2 liblinear	l1	0.75	0.0	0.0	0.0	0.0	0.0
3 liblinear	l1	0.90	0.0	0.0	0.0	0.0	0.0
0 saga	l1	0.10	0.0	0.0	0.0	0.0	0.0
1 saga	l1	0.25	0.0	0.0	0.0	0.0	0.0
2 saga	l1	0.75	0.0	0.0	0.0	0.0	0.0

0.0

0.0

0.0

0.0

8 rows × 63 columns

saga

3

Question 04

Using the test split (of 10%) to show the accuracy, precision, recall, F1-score of the regression model.

0.90

a. Show the output for every possible combination of solver-penalty (newton- cg-l1, newton-cg-l2, newton-cg-none,...)

0.0

- b. Show the output for both the situations: one before performing fine-tuning on the model (with validation data split) and one after performing the fine-tuning on the model (with validation split).
- c. Comment on the improvement, if any.

l1

Previously we have performed the comparative accuracy, recall, precision and f1-score analysis alongwith coefficients of independent variables for variated penalty on solvers like newton-cg, liblinear, lbfgs both before and after fine-tuning, we will here also perform the same analysis on all the possible solvers

```
In [74]: newton_cg_reg_l2 = utilLogRegInfo(X_train,y_train,X_valid,y_valid,X_test,y_test,solver='newton-cg',penalty='l2')[:11]
    newton_cg_reg_none = utilLogRegInfo(X_train,y_train,X_valid,y_valid,X_test,y_test,solver='newton-cg')[:11]
    lbfgs_reg_l2 = utilLogRegInfo(X_train,y_train,X_valid,y_valid,X_test,y_test,solver='lbfgs',penalty='l2')[:11]
    lbfgs_reg_none = utilLogRegInfo(X_train,y_train,X_valid,y_valid,X_test,y_test,solver='lbfgs')[:11]
    liblinear_reg_l1 = utilLogRegInfo(X_train,y_train,X_valid,y_valid,X_test,y_test,solver='liblinear',penalty='l2')[:11]
    sag_reg_l2 = utilLogRegInfo(X_train,y_train,X_valid,y_valid,X_test,y_test,solver='sag',penalty='l2')[:11]
    sag_reg_none = utilLogRegInfo(X_train,y_train,X_valid,y_valid,X_test,y_test,solver='sag')[:11]
    saga_reg_l1 = utilLogRegInfo(X_train,y_train,X_valid,y_valid,X_test,y_test,solver='saga',penalty='l1')[:11]
    saga_reg_l2 = utilLogRegInfo(X_train,y_train,X_valid,y_valid,X_test,y_test,solver='saga',penalty='l1')[:11]
    saga_reg_l2 = utilLogRegInfo(X_train,y_train,X_valid,y_valid,X_test,y_test,solver='saga',penalty='l2')[:11]
    saga_reg_none = utilLogRegInfo(X_train,y_train,X_valid,y_valid,X_test,y_test,solver='saga',penalty='l2')[:11]
    saga_reg_none = utilLogRegInfo(X_train,y_train,X_valid,y_valid,X_test,y_test,solver='saga')[:11]
```

]: _	so	olver	penalty	inverse_regularization_strength	accuracy_before_finetuning	precision_before_finetuning	recall_before_finetuning	F1_score_before_finetuning	accuracy_afte
	o new	vton- cg	l2	1.0	0.964912	0.968750	0.962963	0.964640	
	1 new	vton- cg	none	1.0	0.964912	0.964815	0.964815	0.964815	
	2 l	lbfgs	l2	1.0	0.964912	0.968750	0.962963	0.964640	
	3 l	lbfgs	none	1.0	0.964912	0.964815	0.964815	0.964815	
	4 libli	near	l1	1.0	0.964912	0.968750	0.962963	0.964640	
	5 libli	near	l2	1.0	0.947368	0.954545	0.944444	0.946779	
	6	sag	l2	1.0	0.964912	0.968750	0.962963	0.964640	
	7	sag	none	1.0	0.964912	0.964815	0.964815	0.964815	
	8 :	saga	l1	1.0	0.964912	0.968750	0.962963	0.964640	
	9 :	saga	l2	1.0	0.964912	0.968750	0.962963	0.964640	
1	0 :	saga	none	1.0	0.964912	0.964815	0.964815	0.964815	

Improvements

1. Cross-Validation

K-Fold Cross-Validation: Using cross-validation to ensure our model is not overfitting to the training data and to get a better estimate of its performance on unseen data.

2. Advanced Optimization Techniques

Grid Search or Random Search: Grid Search or Random Search to systematically search for the best hyperparameters. Bayesian Optimization: Bayesian optimization for more efficient hyperparameter tuning.

3. Ensemble of Logistic Regression Models

Model Averaging: Training multiple Logistic Regression models with different hyperparameters and average their predictions to reduce variance.

4. Regularization Techniques

Elastic Net: Combining L1 and L2 regularization to benefit from both techniques.

5. Dimensionality Reduction

Dimensionality Reduction: Using various techniques like PCA to focus more on significant features that aids in processing and overall accuracy.