Machine Learning Evaluation Metrics

Part of the Knowledge Journal Project

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Libraries

```
In [1]: # Numpy
        import numpy as np
        # Pyplot is a state-based interface to matplotlib. It provides a MATLAB-like way
        import matplotlib.pyplot as plt
        # Display plots that are the output of running code cells
        %matplotlib inline
        # Pandas
        import pandas as pd
        # Train and Test Subsets
        from sklearn.model_selection import train_test_split
        # Breast Cancer Wisconsin Dataset from Scikit-Learn Toy datasets
        from sklearn.datasets import load_breast_cancer
        # MLP Classifier
        from sklearn.neural_network import MLPClassifier
        # Accuracy Metric
        from sklearn.metrics import accuracy_score
        # Precision Metric
        from sklearn.metrics import precision_score
        # Recall Metric
        from sklearn.metrics import recall_score
        # F1 Metric
        from sklearn.metrics import f1 score
        # Confusion Matrix
        from sklearn.metrics import confusion matrix
        from sklearn.metrics import ConfusionMatrixDisplay
        # Cross-validation Metric
        from sklearn.model_selection import cross_val_score
```

Breast Cancer Wisconsin Dataset

```
In [2]: data = load_breast_cancer()
```

```
In [3]: \# X = data.data
        # y = data.target
        X, y = load breast cancer(return X y=True)
In [4]: print('Features:\n')
        for feature in data.feature_names:
            print('\t', feature)
       Features:
                mean radius
                mean texture
                mean perimeter
                mean area
                mean smoothness
                mean compactness
                mean concavity
                mean concave points
                mean symmetry
                mean fractal dimension
                radius error
                texture error
                perimeter error
                area error
                smoothness error
                compactness error
                concavity error
                concave points error
                symmetry error
                fractal dimension error
                worst radius
                worst texture
                worst perimeter
                worst area
                worst smoothness
                worst compactness
                worst concavity
                worst concave points
                worst symmetry
                worst fractal dimension
In [5]: print('Targets:\n\t', data.target_names)
       Targets:
                ['malignant' 'benign']
In [6]: # print(data.DESCR)
        Training and Test Subsets
```

```
In [7]: X_train, X_test, y_train, y_test = train_test_split(X, y, test_size=0.3, shuffle=
In [8]: X_train.shape
Out[8]: (398, 30)
In [9]: X_test.shape
```

```
Out[9]: (171, 30)
In [10]: y_train.shape
Out[10]: (398,)
In [11]: y_test.shape
Out[11]: (171,)
```

Model Training

In [12]: clf = MLPClassifier(random_state=None, max_iter=50*X_train.shape[0]).fit(X_train

Model Prediction

In [13]: y_pred = clf.predict(X_test)

Model Evaluation with Accuracy

"All correct predictions"

$$\frac{TP{+}TN}{TP{+}FP{+}TN{+}FN}$$

In [14]: accuracy_score(y_test, y_pred)

Out[14]: 0.9064327485380117

Model Evaluation with Precision

"It's really true when it says so."

$$\frac{TP}{TP+FP}$$

In [15]: precision_score(y_test, y_pred, average='binary')

Out[15]: 0.9405940594059405

Model Evaluation with Recall

"How much predicts positive compare to all real positives."

$$\frac{TP}{TP+FN}$$

```
In [16]: recall_score(y_test, y_pred, average='binary')
Out[16]: 0.9047619047619048
```

Model Evaluation with Specificity

"How much predicts negative compare to all real negatives."

```
\frac{TN}{TN+FP}
```

```
In [17]: tn, fp, fn, tp = confusion_matrix(y_test, y_pred).ravel()
    specificity = tn / (tn+fp)
    specificity
```

Out[17]: np.float64(0.9090909090909091)

Model Evaluation with F1

"Harmonic mean of precision and recall."

```
\frac{2 \cdot Precision \cdot Recall}{Precision + Recall}
```

```
In [18]: f1_score(y_test, y_pred, average='binary')
Out[18]: 0.9223300970873787
```

Model Evaluation with Confusion Matrix

```
In [19]: # Code from this cell in reference 15.

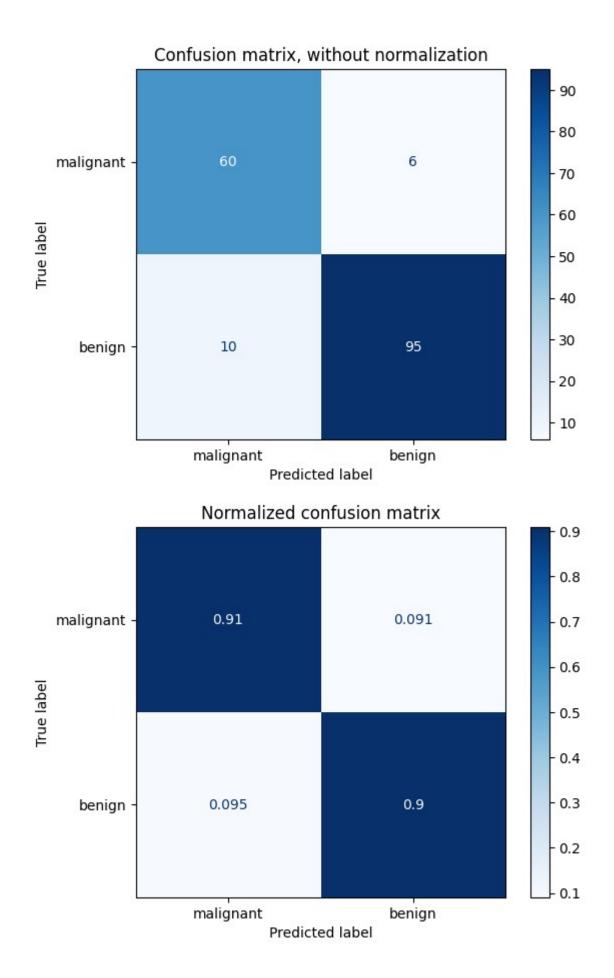
np.set_printoptions(precision=2)
class_names = data.target_names

titles_options = [
    ("Confusion matrix, without normalization", None),
    ("Normalized confusion matrix", "true"),
]
```

```
for title, normalize in titles_options:
     disp = ConfusionMatrixDisplay.from_estimator(
         clf,
         X_test,
         y_test,
         display_labels=class_names,
         cmap=plt.cm.Blues,
         normalize=normalize,
     disp.ax_.set_title(title)
     tn, fp, fn, tp = confusion_matrix(y_test, y_pred).ravel()
     print("tn={} \nfp={} \n".format(tn, fp, fn, tp))
     print(title)
     print(disp.confusion_matrix)
 plt.show()
tn=60
fp=6
fn=10
tp=95
Confusion matrix, without normalization
[[60 6]
[10 95]]
tn=60
fp=6
fn=10
tp=95
```

Normalized confusion matrix

[[0.91 0.09] [0.1 0.9]]



Model Evaluation with Cross-validation

```
In [20]: scores = cross_val_score(clf, X, y, cv=5)
    scores

Out[20]: array([0.92, 0.93, 0.89, 0.93, 0.94])

In [21]: np.mean(scores)

Out[21]: np.float64(0.92094395280236)

In [22]: np.std(scores)

Out[22]: np.float64(0.018297531703108907)
```

References

- 1. Python DOC
- 2. Numpy DOC
- 3. matplotlib.pyplot
- 4. ipython magic functions
- 5. Pandas DOC
- 6. Breast Cancer Wisconsin Dataset
- 7. sklearn.model_selection.train_test_split
- 8. sklearn.neural_network.MLPClassifier
- 9. sklearn.metrics.accuracy_score
- 10. sklearn.metrics.precision_score
- 11. sklearn.metrics.recall_score
- 12. sklearn.metrics.f1_score
- 13. Harmonic Mean
- 14. sklearn.metrics.confusion_matrix
- 15. https://scikit-learn.org/stable/auto_examples/model_selection/ plot_confusion_matrix.html
- 16. Cross-validation