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# **Applied Machine Learning: Module 3 (Evaluation)**

## 03-01: Evaluation for Classification

## **Preamble**

In [2]:

```
%matplotlib notebook
import numpy as np
import pandas as pd
import seaborn as sns
import matplotlib.pyplot as plt
from sklearn.model_selection import train_test_split
from sklearn.datasets import load_digits

dataset = load_digits() # Loads the digits dataset
X, y = dataset.data, dataset.target

for class_name, class_count in zip(dataset.target_names, np.bincount(dataset.target)):
    print(class_name,class_count) # count the number of instances of each digit.
```

- 0 178
- 1 182
- 2 177
- 3 183
- 4 181
- 5 182
- 6 181
- 7 179
- 8 174
- 9 180

```
In [3]:
# Creating a dataset with imbalanced binary classes:
# Negative class (0) is 'not digit 1'
# Positive class (1) is 'digit 1'
y binary imbalanced = y.copy()
y binary imbalanced[y binary imbalanced != 1] = 0
print('Original labels:\t', y[1:30])
print('New binary labels:\t', y_binary_imbalanced[1:30])
Original labels:
                         [1 2 3 4 5 6 7 8 9 0 1 2 3 4 5 6 7 8 9 0 1
2 3 4 5 6 7 8 91
New binary labels:
                         [1 0 0 0 0 0 0 0 0 1 0 0 0 0 0 0 0 1
0 0 0 0 0 0 0 0 0 1
In [4]:
np.bincount(y binary imbalanced)
                                    # Negative class (0) is the most frequent cl
ass
Out[4]:
array([1615, 182])
In [5]:
X train, X test, y train, y test = train test split(X, y binary imbalanced, rand
om state=0)
# Accuracy of Support Vector Machine classifier
from sklearn.svm import SVC
svm = SVC(kernel='rbf', C=1).fit(X train, y train)
svm.score(X test, y test)
Out[5]:
```

# **Dummy Classifiers**

0.9088888888888886

DummyClassifier is a classifier that makes predictions using simple rules, which can be useful as a baseline for comparison against actual classifiers, especially with imbalanced classes.

```
In [6]:
from sklearn.dummy import DummyClassifier
# Negative class (0) is most frequent
dummy_majority = DummyClassifier(strategy = 'most_frequent').fit(X_train, y_trai
n)
# Therefore the dummy 'most frequent' classifier always predicts class 0
# Most frequent parameter is used to predict the most frequent class
y dummy predictions = dummy majority.predict(X test)
y dummy predictions
Out[6]:
0, 0, 0,
  0, 0, 0,
  0, 0, 0,
  0, 0, 0,
  0, 0, 0,
  0, 0, 0,
  0, 0, 0,
  0, 0, 0,
  0, 0, 0,
  0, 0, 0,
  0, 0, 0,
  0, 0, 0,
  0, 0, 0,
  0, 0, 0,
```

0, 0, 0, 0, 0, 0, 0, 0, 0, 0, 0, 0, 0]

0, 0, 0,

0, 0, 0,

0, 0, 0,

0, 0, 0,

```
In [7]:
```

```
dummy_majority.score(X_test, y_test)
Out[7]:
```

## **Change the Type of Kernel**

0.90444444444445

```
In [8]:
```

```
svm = SVC(kernel='linear', C=1).fit(X_train, y_train)
svm.score(X_test, y_test)
```

```
Out[8]:
0.9777777777777775
```

## **Confusion matrices**

Binary (two-class) confusion matrix on Dummy Classifier with strategy = 'most frequent'

In [9]:

```
from sklearn.metrics import confusion_matrix

# Negative class (0) is most frequent
dummy_majority = DummyClassifier(strategy = 'most_frequent').fit(X_train, y_train)
y_majority_predicted = dummy_majority.predict(X_test)
# To get the confusion matrix, we simply pass the y-test set of predicted labels,
# and the y-predicted labels into the confusion matrix
# and then print the output.

confusion = confusion_matrix(y_test, y_majority_predicted)
print('Most frequent class (dummy classifier)\n', confusion)
```

```
Most frequent class (dummy classifier) [[407 0] [ 43 0]]
```

The order of the cells of the little matrix output here is the same as the one in the slides

```
[[TN FP] FN TP]]
```

In particular, the successful predictions of the classifier are **on the diagonal**, where the true class matches the predicted class. Elements on the **subdiagonal** represents errors of different types.

Here we compute the confusion matrices for different choices of classifier in the problem so we can see how they shift slightly with different choice of model.

## Binary (Two Class) confusion matrix on Dummy Classifier with strategy = 'stratified'

```
In [10]:
```

```
# produces random predictions w/ same class proportion as training set
dummy_classprop = DummyClassifier(strategy='stratified').fit(X_train, y_train)
y_classprop_predicted = dummy_classprop.predict(X_test)
confusion = confusion_matrix(y_test, y_classprop_predicted)
print('Random class-proportional prediction (dummy classifier)\n', confusion)
```

```
Random class-proportional prediction (dummy classifier) [[368 39] [39 4]]
```

#### Binary (Two Class) Confusion matrix on Support Linear Vector Machine with Linear Kernel

```
In [11]:
```

```
svm = SVC(kernel='linear', C=1).fit(X_train, y_train)
svm_predicted = svm.predict(X_test)
confusion = confusion_matrix(y_test, svm_predicted)

print('Support vector machine classifier (linear kernel, C=1)\n', confusion)

Support vector machine classifier (linear kernel, C=1)
[[402 5]
[ 5 38]]
```

## Binary (Two Class) Confusion matrix on Logistic Regression model

```
In [12]:
```

```
from sklearn.linear_model import LogisticRegression

lr = LogisticRegression().fit(X_train, y_train)
lr_predicted = lr.predict(X_test)
confusion = confusion_matrix(y_test, lr_predicted)

print('Logistic regression classifier (default settings)\n', confusion)
```

```
Logistic regression classifier (default settings) [[401 6] [ 6 37]]
```

#### Binary (Two Class) Confusion matrix on Decision Tree Classifier of max\_depth 2

In [13]:

```
from sklearn.tree import DecisionTreeClassifier

dt = DecisionTreeClassifier(max_depth=2).fit(X_train, y_train)
tree_predicted = dt.predict(X_test)
confusion = confusion_matrix(y_test, tree_predicted)

print('Decision tree classifier (max_depth = 2)\n', confusion)

Decision tree classifier (max_depth = 2)
[[400 7]
[ 17 26]]
```

Note that the decision tree makes **twice as more** false negative errors as false positive errors.

# 03-02: Evaluation metrics for binary classification

**Evaluation Metrics for Binary Classification - Decision Tree Classifier** 

In [14]:

```
from sklearn.metrics import accuracy_score, precision_score, recall_score, f1_sc
ore
# Accuracy = TP + TN / (TP + TN + FP + FN)
# Precision = TP / (TP + FP)
# Recall = TP / (TP + FN) Also known as sensitivity, or True Positive Rate
# F1 = 2 * Precision * Recall / (Precision + Recall)
print('Accuracy: {:.2f}'.format(accuracy_score(y_test, tree_predicted)))
print('Precision: {:.2f}'.format(precision_score(y_test, tree_predicted)))
print('Recall: {:.2f}'.format(recall_score(y_test, tree_predicted)))
```

Accuracy: 0.95 Precision: 0.79 Recall: 0.60 F1: 0.68

Using classification report

```
# Combined report with all above metrics
from sklearn.metrics import classification_report

print(classification_report(y_test, tree_predicted, target_names=['not 1', '1'
])) # Look at the SKL documentation
# to see what other output options are available to you.
```

support	f1-score	recall	precision	
407 43	0.97 0.68	0.98 0.60	0.96 0.79	not 1 1
450	0.94	0.95	0.94	avg / total

## **Classification Report on Dummy Classifier**

## In [16]:

Random class-	-proportional precision		f1-score	support
not 1	0.90	0.90	0.90	407
1	0.09	0.09	0.09	43
avg / total	0.83	0.83	0.83	450

## **Classification Report on SVM Classifier**

#### In [17]:

SVM

	precision	recall	f1-score	support
not 1	0.99	0.99	0.99	407
1	0.88	0.88	0.88	43
avg / total	0.98	0.98	0.98	450

## **Classification Report on Logistic Regression Classifier**

## In [18]:

Logistic regression

	precision	recall	f1-score	support
not 1	0.99	0.99	0.99	407
1	0.86	0.86	0.86	43
avg / total	0.97	0.97	0.97	450

## **Classification Report on Decision Tree Classifier**

## In [19]:

Decision tree

	precision	recall	f1-score	support
not 1	0.96	0.98	0.97	407
1	0.79	0.60	0.68	43
avg / total	0.94	0.95	0.94	450

# 03-03: Classifier Decision functions

Using decision\_function() method

```
In [20]:
```

```
X_train, X_test, y_train, y_test = train_test_split(X, y_binary_imbalanced, rand
om_state=0)
#lr was actually a logistic regression classifier from the previous lecture.
# lr = LogisticRegression().fit(X_train, y_train)
y_scores_lr = lr.fit(X_train, y_train).decision_function(X_test)
y_score_list = list(zip(y_test[0:20], y_scores_lr[0:20]))
# show the decision_function scores for first 20 instances
y_score_list
```

## Out[20]:

```
[(0, -23.172292973469546),
 (0, -13.542576515500063),
(0, -21.717588760007867),
(0, -18.903065133316439),
 (0, -19.733169947138638),
 (0, -9.7463217496747667),
 (1, 5.2327155658831135),
 (0, -19.308012306288916),
 (0, -25.099330209728528),
 (0, -21.824312362996),
 (0, -24.14378275072049),
 (0, -19.578811099762508),
 (0, -22.568371393280199),
 (0, -10.822590225240777),
 (0, -11.907918741521932),
 (0, -10.977026853802803),
 (1, 11.206811164226373),
(0, -27.64415761980748),
(0, -12.857692102545409),
 (0, -25.848149140240199)
```

Using the predict proba() method

```
In [21]:
```

```
X_train, X_test, y_train, y_test = train_test_split(X, y_binary_imbalanced, rand
om_state=0)
y_proba_lr = lr.fit(X_train, y_train).predict_proba(X_test)
y_proba_list = list(zip(y_test[0:20], y_proba_lr[0:20,1]))
# show the probability of positive class for first 20 instances
y_proba_list
```

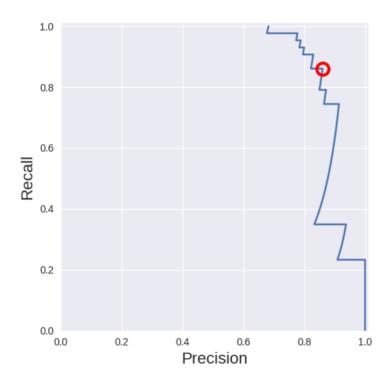
## Out[21]:

```
[(0, 8.6377579220606777e-11),
 (0, 1.3138118599563783e-06),
 (0, 3.6997386039099529e-10),
 (0, 6.1730972504865465e-09),
 (0, 2.6914925394345074e-09),
 (0, 5.8506057771143608e-05),
 (1, 0.99468934644404694),
 (0, 4.1175302368500096e-09),
 (0, 1.2574750894253029e-11),
 (0, 3.3252290754668869e-10),
 (0, 3.2695529799373086e-11),
 (0, 3.1407283576084884e-09),
 (0, 1.5800864117150149e-10),
 (0, 1.9943442430612578e-05),
 (0, 6.7368003023860014e-06),
 (0, 1.7089540581641637e-05),
 (1, 0.9999864188091131),
 (0, 9.8694940340195476e-13),
 (0, 2.6059983600823893e-06),
 (0, 5.9469113009063784e-12)]
```

## 03-04: Precision-recall and ROC Curves

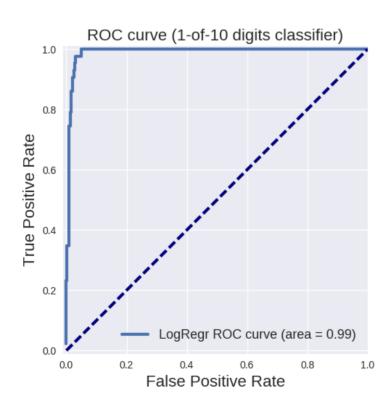
**Precision Recall Curves** 

```
from sklearn.metrics import precision recall curve
precision, recall, thresholds = precision recall curve(y test, y scores lr) ## c
all its constructor
# this returns an array of tuples in the above stated order, where all indices a
re shared across the 3 parameters.
closest zero = np.argmin(np.abs(thresholds)) # this gives the index of the point
with a threshold closest to 0
closest zero p = precision[closest zero] # Find the precision of the closest zer
o threshold element
closest zero r = recall[closest zero] #Find the recall of the closest zero thres
hold element.
# Plot a graph of recall against precision
plt.figure()
plt.xlim([0.0, 1.01])
plt.ylim([0.0, 1.01])
plt.plot(precision, recall, label='Precision-Recall Curve')
plt.plot(closest_zero_p, closest_zero_r, 'o', markersize = 12, fillstyle = 'non
e', c='r', mew=3)
plt.xlabel('Precision', fontsize=16)
plt.ylabel('Recall', fontsize=16)
plt.axes().set aspect('equal')
plt.show()
```

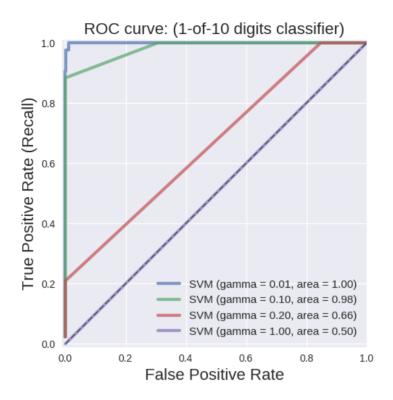


## **ROC curves, Area-Under-Curve (AUC)**

```
from sklearn.metrics import roc curve, auc
X train, X test, y train, y test = train test split(X, y binary imbalanced, rand
om state=0)
# Lr represents the logistic regression classifier - fit the LRC with data and r
un the decision function on it
y_score_lr = lr.fit(X_train, y_train).decision_function(X_test)
#False positive rate, true positive rate and one extra parameter which we don't
need - so we just pass it off as
fpr lr, tpr lr, = roc curve(y test, y score lr)
#Area under the ROC curve.
roc auc lr = auc(fpr lr, tpr lr)
# Plot the ROC curve
plt.figure()
plt.xlim([-0.01, 1.00])
plt.ylim([-0.01, 1.01])
plt.plot(fpr lr, tpr lr, lw=3, label='LogRegr ROC curve (area = {:0.2f})'.format
(roc auc lr))
plt.xlabel('False Positive Rate', fontsize=16)
plt.ylabel('True Positive Rate', fontsize=16)
plt.title('ROC curve (1-of-10 digits classifier)', fontsize=16)
plt.legend(loc='lower right', fontsize=13)
plt.plot([0, 1], [0, 1], color='navy', lw=3, linestyle='--') # random classifie
plt.axes().set aspect('equal')
plt.show()
```



```
from matplotlib import cm #colour map
X train, X test, y train, y test = train test split(X, y binary imbalanced, rand
om state=0)
# Plot a figure of the TPR with respect to the FPR for varying gamma
plt.figure()
plt.xlim([-0.01, 1.00])
plt.ylim([-0.01, 1.01])
# Range of values of gamma
for g in [0.01, 0.1, 0.20, 1]:
    # Create a SVM classifier with RBF kernel and fit the training data to the m
odel
    svm = SVC(gamma=g).fit(X_train, y_train)
    # Find the decision function
    y score svm = svm.decision function(X test)
    # Find the false positive rate and true positive rate for this SVM
    fpr_svm, tpr_svm, _ = roc_curve(y_test, y_score_svm)
    # Find the area under the curve as an evaluation metric
    roc_auc_svm = auc(fpr_svm, tpr_svm)
    # Calculate its accuracy on the test data
    accuracy svm = svm.score(X test, y test)
    print("gamma = {:.2f} accuracy = {:.2f} AUC = {:.2f}".format(g, accuracy
svm,
                                                                     roc_auc_svm
))
    # Plot the ROC curve for this value of gamma
    plt.plot(fpr svm, tpr svm, lw=3, alpha=0.7,
             label='SVM (gamma = {:0.2f}, area = {:0.2f})'.format(g, roc auc svm
))
    # Repeat for other values of gamma
# Label the rest of your axes
plt.xlabel('False Positive Rate', fontsize=16)
plt.ylabel('True Positive Rate (Recall)', fontsize=16)
plt.plot([0, 1], [0, 1], color='k', lw=0.5, linestyle='--')
plt.legend(loc="lower right", fontsize=11)
plt.title('ROC curve: (1-of-10 digits classifier)', fontsize=16)
plt.axes().set aspect('equal')
plt.show()
```



```
      gamma = 0.01
      accuracy = 0.91
      AUC = 1.00

      gamma = 0.10
      accuracy = 0.90
      AUC = 0.98

      gamma = 0.20
      accuracy = 0.90
      AUC = 0.66

      gamma = 1.00
      accuracy = 0.90
      AUC = 0.50
```

## 03-05: Evaluation measures for multi-class classification

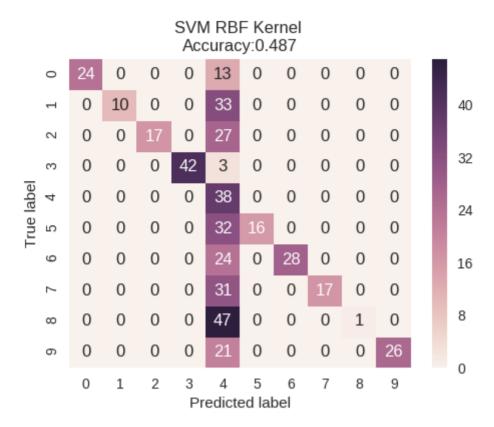
**Multi-class confusion matrix** 

```
dataset = load digits()
X, y = dataset.data, dataset.target
X train mc, X test mc, y train mc, y test mc = train test split(X, y, random sta
te=0)
# Create an SVC with linear kernel
svm = SVC(kernel = 'linear').fit(X train mc, y train mc)
# Get a list of predictions
svm predicted mc = svm.predict(X test mc)
# Create a multiclass confusion matrix of name confusion mc
confusion mc = confusion matrix(y test mc, svm predicted mc)
# Create a dataframe to store the values where each row is the true value and ea
ch column is the predicted value
df cm = pd.DataFrame(confusion mc,
                     index = [i for i in range(0,10)], columns = [i for i in range(0,10)]
ge(0,10)])
# Plot the confusion matrix
plt.figure(figsize=(5.5,4))
# HEY HELLO SEABORN
sns.heatmap(df_cm, annot=True)
# Other MPL support
plt.title('SVM Linear Kernel \nAccuracy: {0:.3f}'.format(accuracy score(y test mc
                                                                         svm predi
cted mc)))
plt.ylabel('True label')
plt.xlabel('Predicted label')
# Here is the 2nd classifier for comparison that uses an SVC rbf kernel - this c
lassifier is meant to perform
# much worse than the prior.
svm = SVC(kernel = 'rbf').fit(X train mc, y train mc)
svm predicted mc = svm.predict(X test mc)
confusion_mc = confusion_matrix(y_test_mc, svm_predicted_mc)
df cm = pd.DataFrame(confusion mc, index = [i for i in range(0,10)],
                  columns = [i \text{ for } i \text{ in } range(0,10)])
plt.figure(figsize = (5.5,4))
sns.heatmap(df cm, annot=True)
plt.title('SVM RBF Kernel \nAccuracy:{0:.3f}'.format(accuracy_score(y_test_mc,
                                                                      svm predicte
dmc)))
plt.ylabel('True label')
plt.xlabel('Predicted label');
```

Accuracy:0.971 ↵ က  $\infty$ 

Predicted label

SVM Linear Kernel



As an aside, it sometimes is useful to display a confusion matrix as a **heat map** in order to highlight the relative frequencies of different kinds of errors.

For comparison, I've included the RBF kernel SVM which did substantially worse on the dataset.

Although you can see from the accuracy score, the RBF kernel has an accuracy of 0.487 compared to the linear kernel, but this accuracy metric alone doesn't give much insight into why this is the case.

For some reason, the SVM RBF kernel **keeps predicting the digit 4.** For instance, out of 44 instances of the digit 2 the classifier only correctly classified 17 of them, and the rest of them was predicted as the digit 4. This would mean an error in the **feature preprocessing** which has caused too many numbers to map to the digit 4. It is **the confusion matrix** that gives you insight into these issues.

## Multi-class classification report

```
print(classification_report(y_test_mc, svm_predicted_mc))
                          recall f1-score
             precision
                                                support
                                        0.79
          0
                   1.00
                             0.65
                                                     37
          1
                   1.00
                              0.23
                                        0.38
                                                     43
          2
                             0.39
                                        0.56
                   1.00
                                                     44
          3
                             0.93
                   1.00
                                        0.97
                                                     45
          4
                                                     38
                   0.14
                             1.00
                                        0.25
          5
                   1.00
                             0.33
                                        0.50
                                                     48
          6
                   1.00
                             0.54
                                        0.70
                                                     52
          7
                   1.00
                             0.35
                                        0.52
                                                     48
          8
                   1.00
                             0.02
                                        0.04
                                                     48
          9
                   1.00
                             0.55
                                        0.71
                                                     47
                                                    450
avg / total
                   0.93
                             0.49
                                        0.54
```

## Micro- vs. macro-averaged metrics using the average = parameter

.format(precision score(y test mc, svm predicted mc, average = 'macro')))

```
Micro-averaged precision = 0.49 (treat instances equally)
Macro-averaged precision = 0.91 (treat classes equally)
```

#### In [28]:

```
Micro-averaged f1 = 0.49 (treat instances equally)
Macro-averaged f1 = 0.54 (treat classes equally)
```

# 03-06: Regression evaluation metrics

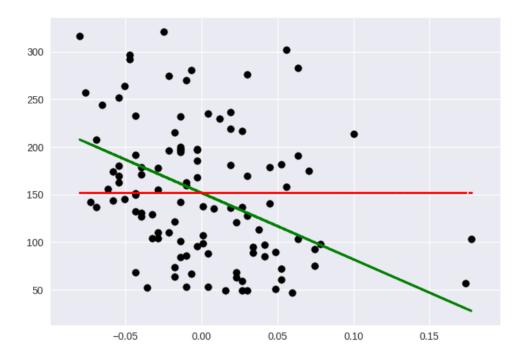
```
%matplotlib notebook
import matplotlib.pyplot as plt
import numpy as np
from sklearn.model selection import train test split
from sklearn import datasets
from sklearn.linear model import LinearRegression
from sklearn.metrics import mean squared error, r2 score
from sklearn.dummy import DummyRegressor
diabetes = datasets.load diabetes()
X = diabetes.data[:, None, 6]
y = diabetes.target
X train, X test, y train, y test = train test split(X, y, random state=0)
# Least Squares Regression
lm = LinearRegression().fit(X train, y train)
# Dummy Regressor
lm dummy mean = DummyRegressor(strategy = 'mean').fit(X train, y train)
# Get both predicted values for both models.
y predict = lm.predict(X test)
y predict dummy mean = lm dummy mean.predict(X test)
# Output the coefficients to the linear model using the coef attribute
print('Linear model, coefficients: ', lm.coef )
# Calculate the mean Squared error for the dummy and for the linear model
print("Mean squared error (dummy): {:.2f}".format(mean squared error(y test,
                                                                     y_predict_d
ummy mean)))
print("Mean squared error (linear model): {:.2f}".format(mean squared error(y te
st, y predict)))
# Calculate the r2 score for the dummy and linear model
print("r2_score (dummy): {:.2f}".format(r2_score(y_test, y_predict_dummy_mean)))
print("r2 score (linear model): {:.2f}".format(r2 score(y test, y predict)))
# Plot outputs
plt.scatter(X test, y test, color='black')
plt.plot(X test, y predict, color='green', linewidth=2)
plt.plot(X_test, y_predict_dummy_mean, color='red', linestyle = 'dashed',
         linewidth=2, label = 'dummy')
plt.show()
```

Linear model, coefficients: [-698.80206267] Mean squared error (dummy): 4965.13

Mean squared error (linear model): 4646.74

r2\_score (dummy): -0.00

r2\_score (linear model): 0.06



We can see that, as expected, the dummy regressor achieves an  $\mathbb{R}^2$  score of 0, since it always makes a constant prediction without looking at the output. In this instance, the linear model provides only **slightly** better fit than the dummy regressor according to both the **mean squared error** and the  $\mathbb{R}^2$  score.

# 03-07: Model selection using evaluation metrics

### **Cross-validation example**

```
In [31]:
```

```
from sklearn.model selection import cross val score
from sklearn.svm import SVC
dataset = load digits()
# again, making this a binary problem with 'digit 1' as positive class
# and 'not 1' as negative class
X, y = dataset.data, dataset.target == 1 # appreciate how neat this syntax look
# Create an SVC linear kernel with default value of C
clf = SVC(kernel='linear', C=1)
# accuracy is the default scoring metric
print('Cross-validation (accuracy)', cross val score(clf, X, y, cv=5))
# use AUC as scoring metric
print('Cross-validation (AUC)', cross val score(clf, X, y, cv=5, scoring = 'roc
auc'))
# use recall as scoring metric
print('Cross-validation (recall)', cross val score(clf, X, y, cv=5, scoring = 'r
ecall'))
Cross-validation (accuracy) [ 0.91944444 0.98611111 0.97214485
97493036 0.969359331
002 0.986756111
Cross-validation (recall) [ 0.81081081  0.89189189  0.83333333  0.83
```

## Grid search example

333333 0.83333333]

```
from sklearn.svm import SVC
from sklearn.model selection import GridSearchCV
from sklearn.metrics import roc auc score
dataset = load digits()
X, y = dataset.data, dataset.target == 1
X_train, X_test, y_train, y_test = train_test_split(X, y, random_state=0)
# Create a classifier with an rbf kernel
clf = SVC(kernel='rbf')
# Parameter values of the variable gamma
grid values = {'gamma': [0.001, 0.01, 0.05, 0.1, 1, 10, 100]}
# ----- default metric to optimize over grid parameters: accuracy----
# Call the GridSearchCV constructor and assign it the grid clf acc variable
grid clf acc = GridSearchCV(clf, param grid = grid values)
# Fit the model to the data
grid clf acc.fit(X train, y train)
# Run the decision function on it
y_decision_fn_scores_acc = grid_clf_acc.decision_function(X_test)
# Use the best params method from the GridSearchCV object to obtain the best pa
rameter for gamma
print('Grid best parameter (max. accuracy): ', grid_clf_acc.best_params_)
# And the score for this best gamma
print('Grid best score (accuracy): ', grid_clf_acc.best_score_)
# ----- Do the same thing for the alternative metric to optimize ove
r grid parameters: AUC -----
grid clf auc = GridSearchCV(clf, param grid = grid values, scoring = 'roc auc')
grid clf auc.fit(X train, y train)
y decision fn scores auc = grid clf auc.decision function(X test)
print('Test set AUC: ', roc_auc_score(y_test, y_decision_fn_scores_auc))
print('Grid best parameter (max. AUC): ', grid clf auc.best params )
print('Grid best score (AUC): ', grid clf auc.best score )
Grid best parameter (max. accuracy): {'gamma': 0.001}
Grid best score (accuracy): 0.996288047513
Test set AUC: 0.999828581224
Grid best parameter (max. AUC): {'gamma': 0.001}
Grid best score (AUC): 0.99987412783
```

In this case, they are the same. But you'll see for some cases that this value of  $\gamma$  is different.

#### **Evaluation metrics supported for model selection**

# from sklearn.metrics.scorer import SCORERS print(sorted(list(SCORERS.keys())))

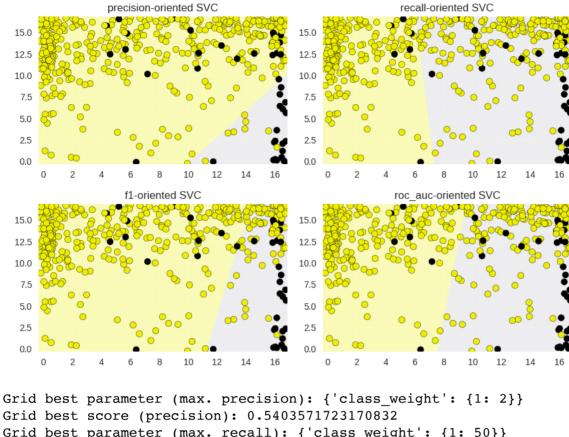
```
['accuracy', 'adjusted_rand_score', 'average_precision', 'f1', 'f1_m acro', 'f1_micro', 'f1_samples', 'f1_weighted', 'log_loss', 'mean_ab solute_error', 'mean_squared_error', 'median_absolute_error', 'neg_l og_loss', 'neg_mean_absolute_error', 'neg_mean_squared_error', 'neg_median_absolute_error', 'precision', 'precision_macro', 'precision_m icro', 'precision_samples', 'precision_weighted', 'r2', 'recall', 'r ecall_macro', 'recall_micro', 'recall_samples', 'recall_weighted', 'roc_auc']
```

You can see stuff like **precision\_micro** that represents micro-averaged-precision and also the **r2** metric for  $R^2$  regression loss.

## Two-feature classification example using the digits dataset

Optimizing a classifier using different evaluation metrics

```
from sklearn.datasets import load digits
from sklearn.model selection import train test split
from adspy shared utilities import plot class regions for classifier subplot
from sklearn.svm import SVC
from sklearn.model selection import GridSearchCV
dataset = load digits()
X, y = dataset.data, dataset.target == 1
X train, X test, y train, y test = train test split(X, y, random state=0)
# Create a two-feature input vector matching the example plot above
# We jitter the points (add a small amount of random noise) in case there are ar
eas
# in feature space where many instances have the same features.
jitter delta = 0.25
X twovar train = X train[:,[20,59]]+ np.random.rand(X train.shape[0], 2) - jitte
r delta
X_twovar_test = X_test[:,[20,59]] + np.random.rand(X_test.shape[0], 2) - jitter
delta
# Create an SVC linear kernel and fit the training data to it
clf = SVC(kernel = 'linear').fit(X twovar train, y train)
# Wait what is this??
grid values = {'class weight':['balanced', {1:2},{1:3},{1:4},{1:5},{1:10},{1:20}
},{1:50}]}
# Plot the figure of 4 graphs - a precision oriented SVC, recall oriented SVC, f
1 oriented SVC and a roc auc SVC
plt.figure(figsize=(9,6))
for i, eval metric in enumerate(('precision', 'recall', 'f1', 'roc auc')):
    # Call the constructor with the classifier used, grid values we want to opti
mise and the scoring metric we want
    # to optimise by.
   grid_clf_custom = GridSearchCV(clf, param_grid=grid_values, scoring=eval_met
ric)
   # Fit the parameters to this model
    grid clf custom.fit(X twovar train, y train)
   # Find the best parameter.
   print('Grid best parameter (max. {0}): {1}'
          .format(eval_metric, grid_clf_custom.best_params_))
   print('Grid best score ({0}): {1}'
          .format(eval metric, grid clf custom.best score ))
   # some MPL support to plot the SVC
   plt.subplots adjust(wspace=0.3, hspace=0.3)
   plot_class_regions_for_classifier_subplot(grid_clf_custom, X_twovar_test, y_
test, None,
                                             None, None, plt.subplot(2, 2, i+1
))
   plt.title(eval metric+'-oriented SVC')
plt.tight_layout()
plt.show()
```



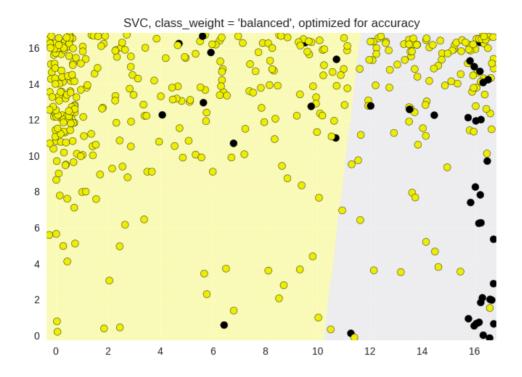
```
Grid best parameter (max. precision): {'class_weight': {1: 2}}
Grid best score (precision): 0.5403571723170832
Grid best parameter (max. recall): {'class_weight': {1: 50}}
Grid best score (recall): 0.9284310837047003
Grid best parameter (max. f1): {'class_weight': {1: 4}}
Grid best score (f1): 0.500943108983028
Grid best parameter (max. roc_auc): {'class_weight': {1: 20}}
Grid best score (roc_auc): 0.8874264392933449
```

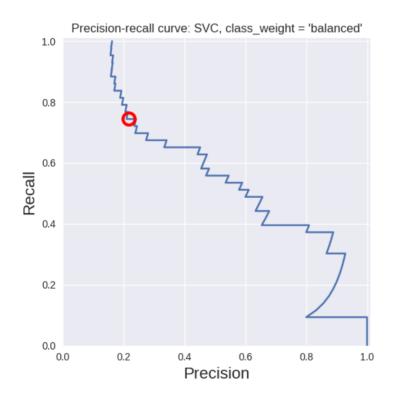
Note that the **F1-oriented** SVC is kind of in the middle of the precision and recall SVCs. This makes sense, because F1 is the **harmonic mean of precision and recall**.

The AUC oriented classifier with an optimal class weight of 5 has a similar decision boundary to the F1-oriented classifier, but shifted slightly in favour of higher recall.

Precision-recall curve for the default SVC classifier (with balanced class weights)

```
from sklearn.model selection import train test split
from sklearn.metrics import precision recall curve
from adspy shared utilities import plot class regions for classifier
from sklearn.svm import SVC
dataset = load digits()
X, y = dataset.data, dataset.target == 1
X_train, X_test, y_train, y_test = train_test_split(X, y, random_state=0)
# create a two-feature input vector matching the example plot above
jitter delta = 0.25
X twovar train = X train[:,[20,59]]+ np.random.rand(X train.shape[0], 2) - jitte
r delta
X twovar test = X test[:,[20,59]] + np.random.rand(X test.shape[0], 2) - jitter
_delta
clf = SVC(kernel='linear', class weight='balanced').fit(X twovar train, y train)
y scores = clf.decision function(X twovar test)
precision, recall, thresholds = precision recall curve(y test, y scores)
closest zero = np.argmin(np.abs(thresholds))
closest zero p = precision[closest zero]
closest_zero_r = recall[closest_zero]
plot class regions for classifier(clf, X twovar test, y test)
plt.title("SVC, class weight = 'balanced', optimized for accuracy")
plt.show()
plt.figure()
plt.xlim([0.0, 1.01])
plt.ylim([0.0, 1.01])
plt.title ("Precision-recall curve: SVC, class weight = 'balanced'")
plt.plot(precision, recall, label = 'Precision-Recall Curve')
plt.plot(closest_zero_p, closest_zero_r, 'o', markersize=12, fillstyle='none', c
='r', mew=3)
plt.xlabel('Precision', fontsize=16)
plt.ylabel('Recall', fontsize=16)
plt.axes().set aspect('equal')
plt.show()
print('At zero threshold, precision: {:.2f}, recall: {:.2f}'
      .format(closest zero p, closest zero r))
```





Take a moment to imagine how the extreme lower right part of the curve represents a decision boundary that is highly precision oriented. As the decision threshold is shifted to become less and less conservative, tracing the curve up and into the left, the classifier becomes more and more like the **recall-oriented** SVC example. Again the red circle represents the precision-recall trade-off achieved at the **zero score mark**, which is the actual decision boundary chosen for the trained classifier.

In [ ]:			