Machine Learning for Diabetes





Photo credit: Pixabay

About one in seven U.S. adults has diabetes now, according to the Centers for Disease Control and Prevention. But by 2050, that rate could skyrocket to as many as one in three. With this in mind, this is what we are going to do today: Learning how to use Machine Learning to help us predict Diabetes. Let's get started!

The Data

The diabetes data set was originated from UCI Machine Learning Repository and can be downloaded from here.

```
import pandas as pd
import numpy as np
import matplotlib.pyplot as plt
%matplotlib inline

diabetes = pd.read_csv('diabetes.csv')
print(diabetes.columns)
```

Index(['Pregnancies', 'Glucose', 'BloodPressure', 'SkinThickness', 'Insulin', 'BMI', 'DiabetesPedigreeFunction', 'Age', 'Outcome'], dtype='object')

```
diabetes.head()
```

	Pregnancies	Glucose	BloodPressure	SkinThickness	Insulin	BMI	DiabetesPedigreeFunction	Age	Outcome
0	6	148	72	35	0	33.6	0.627	50	1
1	1	85	66	29	0	26.6	0.351	31	0
2	8	183	64	0	0	23.3	0.672	32	1
3	1	89	66	23	94	28.1	0.167	21	0
4	0	137	40	35	168	43.1	2.288	33	1

Figure 1

The diabetes data set consists of 768 data points, with 9 features each:

```
print("dimension of diabetes data: {}".format(diabetes.shape))
```

dimension of diabetes data: (768, 9)

"Outcome" is the feature we are going to predict, 0 means No diabetes, 1 means diabetes. Of these 768 data points, 500 are labeled as 0 and 268 as 1:

```
print(diabetes.groupby('Outcome').size())
```

Outcome 0 500 1 268 dtype: int64

Figure 2

import seaborn as sns

sns.countplot(diabetes['Outcome'],label="Count")

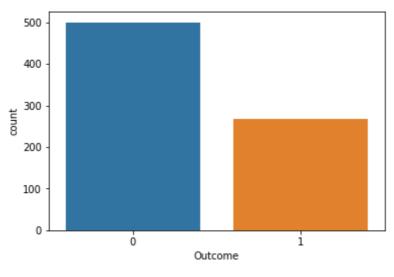


Figure 3

diabetes.info()

```
RangeIndex: 768 entries, 0 to 767
Data columns (total 9 columns):
Pregnancies
                            768 non-null int64
Glucose
                            768 non-null int64
BloodPressure
                            768 non-null int64
SkinThickness
                            768 non-null int64
Insulin
                            768 non-null int64
                            768 non-null float64
DiabetesPedigreeFunction
                            768 non-null float64
                            768 non-null int64
Age
                            768 non-null int64
Outcome
dtypes: float64(2), int64(7)
memory usage: 54.1 KB
```

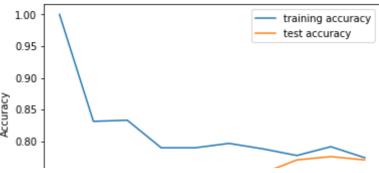
Figure 4

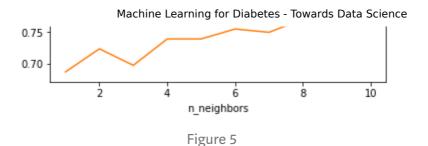
k-Nearest Neighbors

The k-NN algorithm is arguably the simplest machine learning algorithm. Building the model consists only of storing the training data set. To make a prediction for a new data point, the algorithm finds the closest data points in the training data set — its "nearest neighbors."

First, Let's investigate whether we can confirm the connection between model complexity and accuracy:

```
from sklearn.model selection import train test split
X_train, X_test, y_train, y_test = train_test_split(diabetes.loc[:,
diabetes.columns != 'Outcome'], diabetes['Outcome'],
stratify=diabetes['Outcome'], random_state=66)
from sklearn.neighbors import KNeighborsClassifier
training accuracy = []
test accuracy = []
# try n neighbors from 1 to 10
neighbors_settings = range(1, 11)
for n neighbors in neighbors settings:
    # build the model
    knn = KNeighborsClassifier(n_neighbors=n_neighbors)
    knn.fit(X_train, y_train)
    # record training set accuracy
    training_accuracy.append(knn.score(X_train, y_train))
    # record test set accuracy
    test accuracy.append(knn.score(X test, y test))
plt.plot(neighbors_settings, training_accuracy, label="training"
accuracy")
plt.plot(neighbors_settings, test_accuracy, label="test accuracy")
plt.ylabel("Accuracy")
plt.xlabel("n_neighbors")
plt.legend()
plt.savefig('knn_compare_model')
```





The above plot shows the training and test set accuracy on the y-axis against the setting of n_neighbors on the x-axis. Considering if we choose one single nearest neighbor, the prediction on the training set is perfect. But when more neighbors are considered, the training accuracy drops, indicating that using the single nearest neighbor leads to a model that is too complex. The best performance is somewhere around 9 neighbors.

The plot suggests that we should choose n_neighbors=9. Here we are:

```
knn = KNeighborsClassifier(n_neighbors=9)
knn.fit(X_train, y_train)

print('Accuracy of K-NN classifier on training set:
{:.2f}'.format(knn.score(X_train, y_train)))
print('Accuracy of K-NN classifier on test set:
{:.2f}'.format(knn.score(X_test, y_test)))
```

Accuracy of K-NN classifier on training set: 0.79

Accuracy of K-NN classifier on test set: 0.78

Logistic regression

Logistic Regression is one of the most common classification algorithms.

```
from sklearn.linear_model import LogisticRegression

logreg = LogisticRegression().fit(X_train, y_train)
print("Training set score: {:.3f}".format(logreg.score(X_train, y_train)))
print("Test set score: {:.3f}".format(logreg.score(X_test, y_test)))
```

Training set accuracy: 0.781

Test set accuracy: 0.771

The default value of C=1 provides with 78% accuracy on the training and 77% accuracy on the test set.

```
logreg001 = LogisticRegression(C=0.01).fit(X_train, y_train)
print("Training set accuracy:
{:.3f}".format(logreg001.score(X_train, y_train)))
print("Test set accuracy: {:.3f}".format(logreg001.score(X_test, y_test)))
```

Training set accuracy: 0.700

Test set accuracy: 0.703

Using C=0.01 results in lower accuracy on both the training and the test sets.

```
logreg100 = LogisticRegression(C=100).fit(X_train, y_train)
print("Training set accuracy:
{:.3f}".format(logreg100.score(X_train, y_train)))
print("Test set accuracy: {:.3f}".format(logreg100.score(X_test, y_test)))
```

Training set accuracy: 0.785

Test set accuracy: 0.766

Using C=100 results in a little bit higher accuracy on the training set and little bit lower accuracy on the test set, confirming that less regularization and a more complex model may not generalize better than default setting.

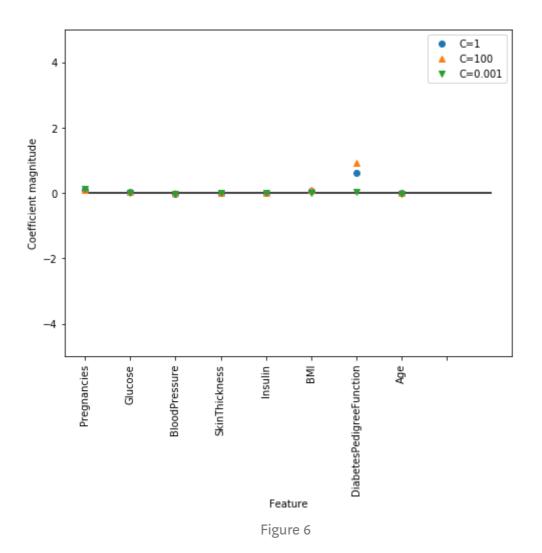
Therefore, we should choose default value C=1.

Let's visualize the coefficients learned by the models with the three different settings of the regularization parameter C.

Stronger regularization (C=0.001) pushes coefficients more and more toward zero. Inspecting the plot more closely, we can also see that feature "DiabetesPedigreeFunction", for C=100, C=1 and C=0.001, the coefficient is positive. This indicates that high "DiabetesPedigreeFunction" feature is related to a sample being "diabetes", regardless which model we look at.

```
diabetes_features = [x for i,x in enumerate(diabetes.columns) if
i!=8]

plt.figure(figsize=(8,6))
plt.plot(logreg.coef_.T, 'o', label="C=1")
plt.plot(logreg100.coef_.T, '^', label="C=100")
plt.plot(logreg001.coef_.T, 'v', label="C=0.001")
plt.xticks(range(diabetes.shape[1]), diabetes_features, rotation=90)
plt.hlines(0, 0, diabetes.shape[1])
plt.ylim(-5, 5)
plt.xlabel("Feature")
plt.ylabel("Coefficient magnitude")
plt.legend()
plt.savefig('log_coef')
```



Decision Tree

from sklearn.tree import DecisionTreeClassifier

```
tree = DecisionTreeClassifier(random_state=0)
tree.fit(X_train, y_train)
print("Accuracy on training set: {:.3f}".format(tree.score(X_train, y_train)))
print("Accuracy on test set: {:.3f}".format(tree.score(X_test, y_test)))
```

Accuracy on training set: 1.000

Accuracy on test set: 0.714

The accuracy on the training set is 100%, while the test set accuracy is much worse. This is an indicative that the tree is overfitting and not generalizing well to new data. Therefore, we need to apply pre-pruning to the tree.

We set max_depth=3, limiting the depth of the tree decreases overfitting. This leads to a lower accuracy on the training set, but an improvement on the test set.

```
tree = DecisionTreeClassifier(max_depth=3, random_state=0)
tree.fit(X_train, y_train)

print("Accuracy on training set: {:.3f}".format(tree.score(X_train, y_train)))
print("Accuracy on test set: {:.3f}".format(tree.score(X_test, y_test)))
```

Accuracy on training set: 0.773

Accuracy on test set: 0.740

Feature Importance in Decision Trees

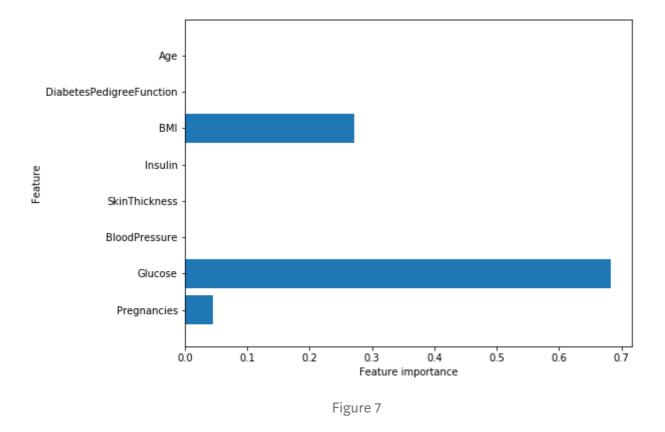
Feature importance rates how important each feature is for the decision a tree makes. It is a number between 0 and 1 for each feature, where 0 means "not used at all" and 1 means "perfectly predicts the target". The feature importances always sum to 1:

```
print("Feature importances:\n{}".format(tree.feature_importances_))
```

Feature importances: [0.04554275 0.6830362 0. 0. 0. 0.27142106 0. 0.]

Then we can visualize the feature importances:

```
def plot_feature_importances_diabetes(model):
    plt.figure(figsize=(8,6))
    n_features = 8
    plt.barh(range(n_features), model.feature_importances_,
align='center')
    plt.yticks(np.arange(n_features), diabetes_features)
    plt.xlabel("Feature importance")
    plt.ylabel("Feature")
    plt.ylim(-1, n_features)
plot_feature_importances_diabetes(tree)
plt.savefig('feature_importance')
```



Feature "Glucose" is by far the most important feature.

Random Forest

Let's apply a random forest consisting of 100 trees on the diabetes data set:

```
from sklearn.ensemble import RandomForestClassifier

rf = RandomForestClassifier(n_estimators=100, random_state=0)
rf.fit(X_train, y_train)
```

```
print("Accuracy on training set: {:.3f}".format(rf.score(X_train, y_train)))
print("Accuracy on test set: {:.3f}".format(rf.score(X_test, y_test)))
```

Accuracy on training set: 1.000

Accuracy on test set: 0.786

The random forest gives us an accuracy of 78.6%, better than the logistic regression model or a single decision tree, without tuning any parameters. However, we can adjust the max_features setting, to see whether the result can be improved.

```
rf1 = RandomForestClassifier(max_depth=3, n_estimators=100,
random_state=0)
rf1.fit(X_train, y_train)
print("Accuracy on training set: {:.3f}".format(rf1.score(X_train,
y_train)))
print("Accuracy on test set: {:.3f}".format(rf1.score(X_test,
y_test)))
```

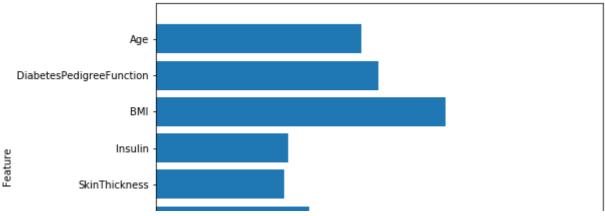
Accuracy on training set: 0.800

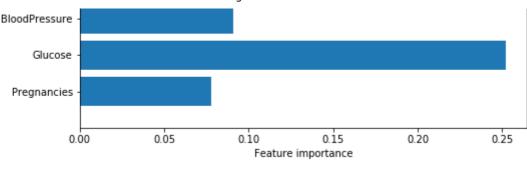
Accuracy on test set: 0.755

It did not, this indicates that the default parameters of the random forest work well.

Feature importance in Random Forest

plot_feature_importances_diabetes(rf)





Similarly to the single decision tree, the random forest also gives a lot of importance to the "Glucose" feature, but it also chooses "BMI" to be the 2nd most informative feature overall. The randomness in building the random forest forces the algorithm to consider many possible explanations, the result being that the random forest captures a much broader picture of the data than a single tree.

Figure 8

Gradient Boosting

```
from sklearn.ensemble import GradientBoostingClassifier

gb = GradientBoostingClassifier(random_state=0)
gb.fit(X_train, y_train)

print("Accuracy on training set: {:.3f}".format(gb.score(X_train, y_train)))
print("Accuracy on test set: {:.3f}".format(gb.score(X_test, y_test)))
```

Accuracy on training set: 0.917

Accuracy on test set: 0.792

We are likely to be overfitting. To reduce overfitting, we could either apply stronger pre-pruning by limiting the maximum depth or lower the learning rate:

```
gb1 = GradientBoostingClassifier(random_state=0, max_depth=1)
gb1.fit(X_train, y_train)

print("Accuracy on training set: {:.3f}".format(gb1.score(X_train, y_train)))
print("Accuracy on test set: {:.3f}".format(gb1.score(X_test, y_test)))
```

Accuracy on training set: 0.804

Accuracy on test set: 0.781

```
gb2 = GradientBoostingClassifier(random_state=0, learning_rate=0.01)
gb2.fit(X_train, y_train)

print("Accuracy on training set: {:.3f}".format(gb2.score(X_train, y_train)))
print("Accuracy on test set: {:.3f}".format(gb2.score(X_test, y_test)))
```

Accuracy on training set: 0.802

Accuracy on test set: 0.776

Both methods of decreasing the model complexity reduced the training set accuracy, as expected. However, in this case, none of these methods increased the generalization performance of the test set.

We can visualize the feature importances to get more insight into our model even though we are not really happy with the model:

plot_feature_importances_diabetes(gb1)

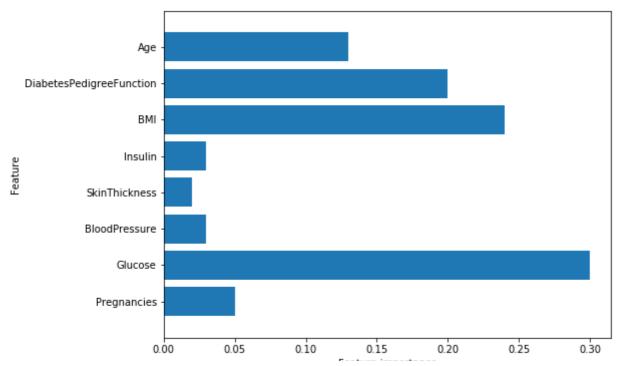


Figure 9

We can see that the feature importances of the gradient boosted trees are somewhat similar to the feature importances of the random forests, it gives weight to all of the features in this case.

Support Vector Machine

```
from sklearn.svm import SVC

svc = SVC()
svc.fit(X_train, y_train)

print("Accuracy on training set: {:.2f}".format(svc.score(X_train, y_train)))
print("Accuracy on test set: {:.2f}".format(svc.score(X_test, y_test)))
```

Accuracy on training set: 1.00

Accuracy on test set: 0.65

The model overfits quite substantially, with a perfect score on the training set and only 65% accuracy on the test set.

SVM requires all the features to vary on a similar scale. We will need to re-scale our data that all the features are approximately on the same scale:

```
from sklearn.preprocessing import MinMaxScaler

scaler = MinMaxScaler()
X_train_scaled = scaler.fit_transform(X_train)
X_test_scaled = scaler.fit_transform(X_test)

svc = SVC()
svc.fit(X_train_scaled, y_train)

print("Accuracy on training set:
{:.2f}".format(svc.score(X_train_scaled, y_train)))
print("Accuracy on test set: {:.2f}".format(svc.score(X_test_scaled, y_test)))
```

Accuracy on training set: 0.77

Accuracy on test set: 0.77

Scaling the data made a huge difference! Now we are actually underfitting, where training and test set performance are quite similar but less close to 100% accuracy. From here, we can try increasing either C or gamma to fit a more complex model.

```
svc = SVC(C=1000)
svc.fit(X_train_scaled, y_train)

print("Accuracy on training set: {:.3f}".format(
        svc.score(X_train_scaled, y_train)))
print("Accuracy on test set: {:.3f}".format(svc.score(X_test_scaled, y_test)))
```

Accuracy on training set: 0.790

Accuracy on test set: 0.797

Here, increasing C allows us to improve the model, resulting in 79.7% test set accuracy.

Deep Learning

```
from sklearn.neural_network import MLPClassifier

mlp = MLPClassifier(random_state=42)
mlp.fit(X_train, y_train)

print("Accuracy on training set: {:.2f}".format(mlp.score(X_train, y_train)))
print("Accuracy on test set: {:.2f}".format(mlp.score(X_test, y_test)))
```

Accuracy on training set: 0.71

Accuracy on test set: 0.67

The accuracy of the Multilayer perceptrons (MLP) is not as good as the other models at all, this is likely due to scaling of the data. deep learning algorithms also expect all

input features to vary in a similar way, and ideally to have a mean of 0, and a variance of 1. We must re-scale our data so that it fulfills these requirements.

Accuracy on training set: 0.823

Accuracy on test set: 0.802

Let's increase the number of iterations:

```
mlp = MLPClassifier(max_iter=1000, random_state=0)
mlp.fit(X_train_scaled, y_train)

print("Accuracy on training set: {:.3f}".format(
        mlp.score(X_train_scaled, y_train)))
print("Accuracy on test set: {:.3f}".format(mlp.score(X_test_scaled, y_test)))
```

Accuracy on training set: 0.877

Accuracy on test set: 0.755

Increasing the number of iterations only increased the training set performance, not the test set performance.

Let's increase the alpha parameter and add stronger regularization of the weights:

```
mlp = MLPClassifier(max_iter=1000, alpha=1, random_state=0)
mlp.fit(X_train_scaled, y_train)
```

```
print("Accuracy on training set: {:.3f}".format(
    mlp.score(X_train_scaled, y_train)))
print("Accuracy on test set: {:.3f}".format(mlp.score(X_test_scaled, y_test)))
```

Accuracy on training set: 0.795

Accuracy on test set: 0.792

The result is good, but we are not able to increase the test accuracy further.

Therefore, our best model so far is default deep learning model after scaling.

Finally, we plot a heat map of the first layer weights in a neural network learned on the diabetes data set.

```
plt.figure(figsize=(20, 5))
plt.imshow(mlp.coefs_[0], interpolation='none', cmap='viridis')
plt.yticks(range(8), diabetes_features)
plt.xlabel("Columns in weight matrix")
plt.ylabel("Input feature")
plt.colorbar()
```

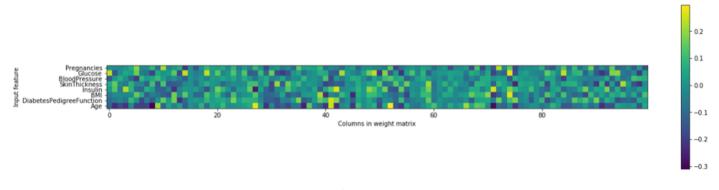


Figure 10

From the heat map, it is not easy to point out quickly that which feature (features) have relatively low weights compared to the other features.

Summary

We practiced a wide array of machine learning models for classification and regression, what their advantages and disadvantages are, and how to control model complexity for

each of them. We saw that for many of the algorithms, setting the right parameters is important for good performance.

We should be able to know how to apply, tune, and analyze the models we practiced above. It's your turn now! Try applying any of these algorithms to the built-in data sets in scikit-learn or any data set at your choice. Happy Machine Learning!

Source code that created this post can be found here. I would be pleased to receive feedback or questions on any of the above.

Reference: Introduction to Machine Learning with Python

Machine Learning Data Science Python Data Visualization Towards Data Science

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