# Understanding Linear and Polynomial Regression in Python

## Introduction

In this document, we will explore two fundamental regression techniques: **Linear Regression** and **Polynomial Regression**. These are powerful tools used in data science and machine learning to model relationships between variables and make predictions. We'll use the **Diabetes** dataset from sklearn.datasets to demonstrate these models in Python.

## What is Linear Regression?

Linear regression is a statistical method used to model the relationship between a dependent variable and one or more independent variables by fitting a linear equation to observed data. The goal is to find the best-fitting straight line through the data points that can be used for prediction.

In simple linear regression, the model is defined as:

$$y = \beta_0 + \beta_1 x + \epsilon$$

Where:

- y is the dependent variable
- $\cdot x$  is the independent variable
- $\beta_0$  is the intercept
- $\beta_1$  is the slope (coefficient)
- $\epsilon$  is the error term

# What is Polynomial Regression?

Polynomial regression is an extension of linear regression where the relationship between the independent variable and the dependent variable is modeled as an nth-degree polynomial. It is useful when the data shows a curved trend that a straight line cannot capture.

The general form of a polynomial regression equation is:

$$y = \beta_0 + \beta_1 x + \beta_2 x^2 + \ldots + \beta_n x^n + \epsilon$$

This allows the model to fit more complex, non-linear patterns in the data while still using a linear model under the hood through feature transformation.

# 1. Loading the Dataset and Libraries

```
import pandas as pd
import numpy as np
import matplotlib.pyplot as plt
from sklearn.datasets import load_diabetes
from sklearn.model_selection import train_test_split
from sklearn.linear_model import LinearRegression
from sklearn.metrics import r2_score, mean_squared_error, mean_absolute_error
from sklearn.preprocessing import PolynomialFeatures
from sklearn.pipeline import make_pipeline
```

## **Explanation:**

- pandas, numpy, and matplotlib.pyplot are standard libraries for data manipulation and visualization.
- load\_diabetes | provides a sample dataset used for regression tasks.
- train\_test\_split is used to divide our data into training and testing sets.
- LinearRegression creates our linear model.
- PolynomialFeatures and make\_pipeline help us build a polynomial regression model.

## 2. Preparing the Dataset

```
diabetes = load_diabetes()
data = pd.DataFrame(diabetes.data, columns=diabetes.feature_names)
data['target'] = diabetes.target
```

#### **Explanation:**

We load the diabetes dataset into a DataFrame called data and add the target variable (a measure of disease progression) as a new column.

# 3. Selecting a Feature

```
X = data[['bmi']]
y = data['target']
```

#### **Explanation:**

We focus on a single feature: **BMI** (Body Mass Index). X represents the input (independent variable), and y represents the target (dependent variable).

# 4. Splitting the Data

```
X_train, X_test, y_train, y_test = train_test_split(X, y, test_size=0.2)
```

## **Explanation:**

We split the dataset into training (80%) and testing (20%) sets. This ensures that our model is trained on one portion and evaluated on another to avoid overfitting.

# 5. Building a Linear Regression Model

```
lr = LinearRegression()
lr.fit(X_train, y_train)
y_predictions = lr.predict(X_test)
```

## **Explanation:**

We create and train a linear regression model using LinearRegression. After fitting it on training data, we use it to predict the test data.

## 6. Evaluating the Linear Model

```
r2 = r2_score(y_test, y_predictions)
mse = mean_squared_error(y_test, y_predictions)
mae = mean_absolute_error(y_test, y_predictions)
intercept = lr.intercept_

print("The R2 score is:", r2)
print("The Mean Squared Error is:", mse)
print("The Mean Absolute Error is:", mae)
print("The Intercept is:", intercept)
```

## **Explanation:**

- **R2 Score** measures how well the model explains the variation in the data.
- MSE and MAE are error metrics. Lower values indicate better model performance.

# 7. Visualizing Linear Regression

```
plt.scatter(X, y)
plt.xlabel('Persons BMI')
```

```
plt.ylabel('Persons Diabetes level')
plt.show()
```

## **Explanation:**

This scatter plot helps visualize the relationship between BMI and diabetes level.

# 8. Building a Polynomial Regression Model

```
poly_model = make_pipeline(PolynomialFeatures(degree=2), LinearRegression())
poly_model.fit(X_train, y_train)
poly_y_predictions = poly_model.predict(X_test)
```

### **Explanation:**

Polynomial regression models non-linear relationships. PolynomialFeatures transforms the input feature to include powers of the feature (e.g., BMI^2).

# 9. Evaluating the Polynomial Model

```
pol_r2 = r2_score(y_test, poly_y_predictions)
pol_mse = mean_squared_error(y_test, poly_y_predictions)
pol_mae = mean_absolute_error(y_test, poly_y_predictions)

print("The R2 score is:", pol_r2)
print("The Mean Squared Error is:", pol_mse)
print("The Mean Absolute Error is:", pol_mae)
```

### **Explanation:**

We use the same evaluation metrics to assess how well the polynomial model performs in comparison to the linear model.

# **Summary**

- Linear regression fits a straight line to the data.
- **Polynomial regression** allows for a curved line, better capturing non-linear trends.
- Model performance is compared using R2, MSE, and MAE.
- Visualizing and evaluating models helps choose the right approach for your dataset.

Both regression techniques are crucial tools in the data scientist's toolkit, and Python's scikit-learn makes implementing them straightforward and efficient.