Problem 1

Step 1: After having a look with df.info() and df.describe(), the output shows as follows:

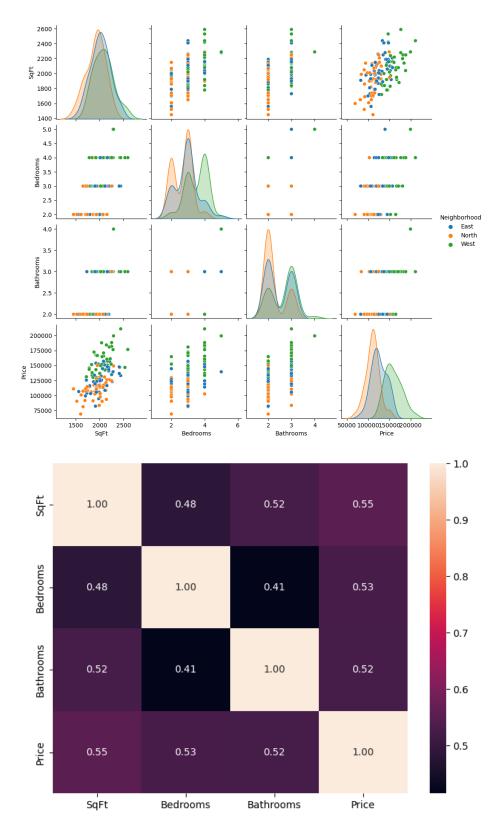
	SqFt	Bedrooms	Bathrooms	Price
count	128.000000	128.000000	128.000000	128.000000
mean	2000.937500	3.023438	2.445312	130427.343750
std	211.572431	0.725951	0.514492	26868.770371
min	1450.000000	2.000000	2.000000	69100.000000
25%	1880.000000	3.000000	2.000000	111325.000000
50%	2000.000000	3.000000	2.000000	125950.000000
75%	2140.000000	3.000000	3.000000	148250.000000
max	2590.000000	5.000000	4.000000	211200.000000

<class 'pandas.core.frame.DataFrame'>
RangeIndex: 128 entries, 0 to 127
Data columns (total 5 columns):

#	Column	Non-Null Count	Dtype			
0	SqFt	128 non-null	int64			
1	Bedrooms	128 non-null	int64			
2	Bathrooms	128 non-null	int64			
3	Neighborhood	128 non-null	category			
4	Price	128 non-null	int64			
dtypes: category(1), int64(4)						
memory usage: 4.4 KB						
None						

The dataset contains 128 entries and 5 attributes: SqFt, Bedrooms, Bathrooms, Neighborhood, and Price. The Neighborhood feature is **categorical**, and the rest are numerical. Houses range from 1450 to 2590 square feet, with 2 to 5 bedrooms and 2 to 4 bathrooms. Prices vary significantly, from 69,100 to 211,200. There are no missing values like NaN or null, and the dataset occupies 4.4 KB of memory.

Step 2: After some visualization, the output shows as:



The pairplot indicates that **larger square footage**, **more bedrooms**, **and more bathrooms** are all moderately associated with higher house prices. The distribution of points suggests that **the neighborhood** also influences house prices, with some neighborhoods appearing to have consistently higher or lower prices.

The heatmap confirms **moderate positive correlations** between house size, number of bedrooms, number of bathrooms, and price, with the correlation coefficients all above 0.5 but not so high.

Step 3: After performing the transformer using ColumnTransformer, we transfer the previous data set into the following df_encoded, with the first three column as one-hot encoding representing *Neighbourhood*, followed by the rest as *SqFt*, *Bedrooms* and *Bathrooms*. Here is part of the array:

```
array([[1.000e+00, 0.000e+00, 0.000e+00, 1.790e+03, 2.000e+00, 2.000e+00, 1.143e+05], [1.000e+00, 0.000e+00, 0.000e+00, 2.030e+03, 4.000e+00, 2.000e+00, 1.142e+05], [1.000e+00, 0.000e+00, 0.000e+00, 1.740e+03, 3.000e+00, 2.000e+00, 1.148e+05],
```

After that, we randomly split the data with train test split into two parts for later analysis.

Step 4: By executing the linear regression using model.fit and model.predict, we report the **training RMSE** to be 1.4408543108972288e-12 and the **testing RMSE** to be 0, which seems to be a valid and accurate match.

Problem 2

Step 1: As instructed, we load the diabetes dataset. To better coordinate with the matrix operations, I add to [X] an additional first column of all ones, functioning as the intercept. Also, [y] is converted into an 2D array.

With reference, learning_rate = 0.1 and iterations = 1000 are chosen for the problem. m indicates the rows of the design matrix with intercept x_b , and theta is initialized to be $\theta_i \sim N(0,1)$.

For the step of gradient descent, the gradient $rac{2}{m}X_b^T(heta^TX_b-y)$ is implemented as

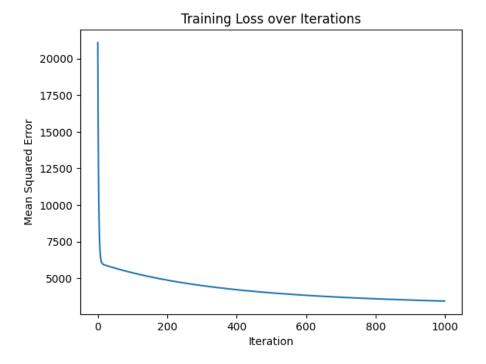
```
gradients = 2/m * X_b.T.dot(X_b.dot(theta) - y)
```

and update as $\theta := \theta - \alpha \nabla f(\mathbf{x})$.

Step 2: After splitting the whole dataset into two, I run the linear regression again on the test set. This time a list train_loss is initialized, and for each iteration the loss is appended:

```
loss = mean_squared_error(y_train, X_train.dot(theta))
train_loss.append(loss)
```

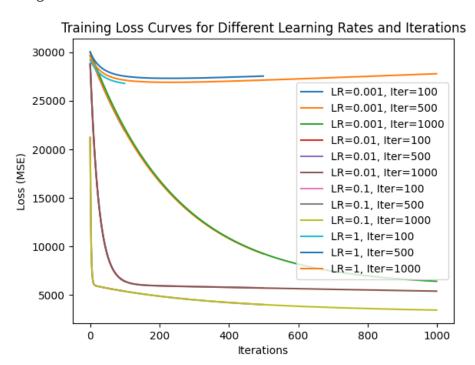
After the training, we generate both the prediction of training set and that on the test set, <code>y_train_pred</code> and <code>y_test_pred</code>, after which the RMSE is calculated respectively: The **training RMSE** is roughly 58.672, and the **testing RMSE** is 55.583. The plot looks like:



Step 3: In this part, I tuned different hyper-parameter of gradient descents, which are **learning rate** LR and iterations Iter. The only difference lies in encapsulating the steps into the <code>gradient_descent</code> function, and add two <code>for</code> iterations for every experiment. The result tells that:

```
LR: 0.001, Iter: 100, Training RMSE: 148.08575669916843, Testing RMSE: 138.59615078130028 LR: 0.001, Iter: 500, Training RMSE: 96.1373205583258, Testing RMSE: 87.37643581536264 LR: 0.001, Iter: 1000, Training RMSE: 80.07649571882452, Testing RMSE: 73.31543992980345 LR: 0.01, Iter: 100, Training RMSE: 80.09388107011452, Testing RMSE: 73.38531514979655 LR: 0.01, Iter: 500, Training RMSE: 75.69320096292356, Testing RMSE: 70.96342208236184 LR: 0.01, Iter: 1000, Training RMSE: 73.46176222414701, Testing RMSE: 68.75493975871984 LR: 0.1, Iter: 100, Training RMSE: 73.39366161382566, Testing RMSE: 68.68437119492309 LR: 0.1, Iter: 500, Training RMSE: 63.28627286684141, Testing RMSE: 59.17672788209183 LR: 0.1, Iter: 1000, Training RMSE: 58.68920034516669, Testing RMSE: 55.604002757840455 LR: 1, Iter: 100, Training RMSE: 163.6610006570304, Testing RMSE: 161.99054529715568 LR: 1, Iter: 500, Training RMSE: 165.95113689077763, Testing RMSE: 169.33777373700747 LR: 1, Iter: 1000, Training RMSE: 166.67603473647068, Testing RMSE: 170.46821567504807
```

with the corresponding curves as:



From the plot the influnces of these hyper-parameters could be analysed:

- **Iterations:** A sufficent number of iterations often yields a complete gradient descent, but it will be a waste of resources if this number is too large. An iteration of 600 could have been enough.
- **Learning rate:** If the learning rate is too large, the step might be so large that it steps over the optima all the time. If the learning rate is too small, it might take such a huge number of iterations to descend to the optima. A learning rate of 0.1 seems good.