Dataset regarding medical expenses for an insurance company.

Model used is Linear Regression and Root Mean Squared error is calculated.

Finally, accuracy of the model after scaling and encoding of data is shown.

```
In [139]:
from urllib.request import urlretrieve
expenses_url = 'https://raw.githubusercontent.com/JovianML/opendatasets/master/data/medical
In [140]:
urlretrieve(expenses_url, 'expenses.csv')
Out[140]:
('expenses.csv', <http.client.HTTPMessage at 0x2ef05f1fc40>)
In [141]:
import pandas as pd
data = pd.read_csv('expenses.csv')
In [142]:
data.shape
Out[142]:
(1338, 7)
In [143]:
data.head()
Out[143]:
          sex
                 bmi children smoker
                                        region
                                                  charges
   age
0
              27.900
                                               16884.92400
        female
                                 ves
                                     southwest
    18
         male
              33.770
                                                1725.55230
                                  no
                                      southeast
```

2

3

28

33

32

male

male

33.000

22.705

male 28.880

3

0

southeast

northwest

northwest

no

4449.46200

21984.47061

3866.85520

```
In [144]:
```

```
data['region'].value_counts()
```

Out[144]:

southeast 364 southwest 325 northwest 325 northeast 324

Name: region, dtype: int64

In [145]:

```
data.describe()
```

Out[145]:

	age	bmi	children	charges	
count	1338.000000	1338.000000	1338.000000	1338.000000	
mean	39.207025	30.663397	1.094918	13270.422265	
std	14.049960	6.098187	1.205493	12110.011237	
min	18.000000	15.960000	0.000000	1121.873900	
25%	27.000000	26.296250	0.000000	4740.287150	
50%	39.000000	30.400000	1.000000	9382.033000	
75%	51.000000	34.693750	2.000000	16639.912515	
max	64.000000	53.130000	5.000000	63770.428010	

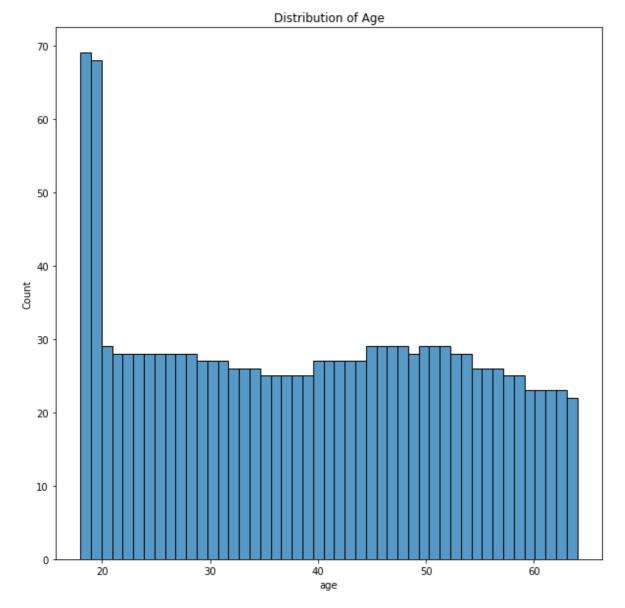
Exploring the data

In [146]:

```
import matplotlib.pyplot as plt
import seaborn as sns
```

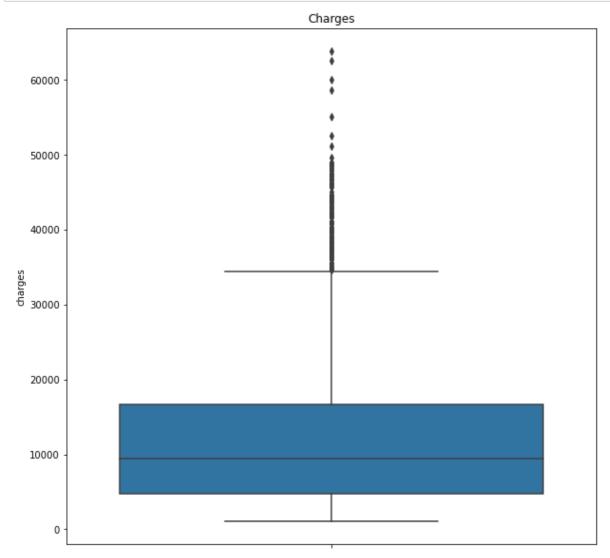
In [147]:

```
plt.figure(figsize=(10,10))
sns.histplot(data=data, x='age', bins=47)
plt.title("Distribution of Age")
plt.show()
```



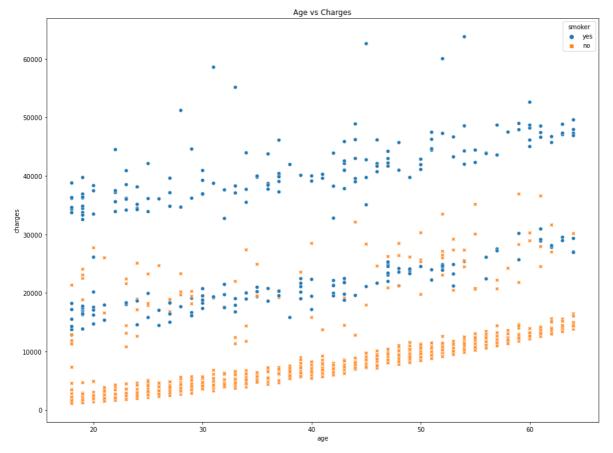
In [148]:

```
plt.figure(figsize=(10,10))
sns.boxplot(data=data, y='charges')
plt.title("Charges")
plt.show()
```



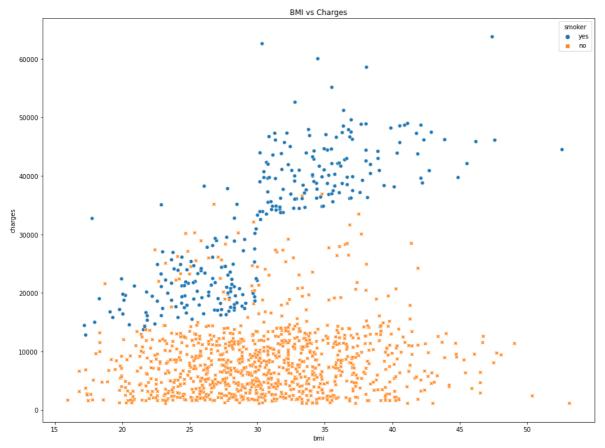
In [149]:

```
plt.figure(figsize=(16,12))
sns.scatterplot(data=data, x='age', y='charges', hue='smoker', style='smoker')
plt.title("Age vs Charges")
plt.show()
```



In [150]:

```
plt.figure(figsize=(16,12))
sns.scatterplot(data=data, x='bmi', y='charges', hue='smoker', style='smoker')
plt.title("BMI vs Charges")
plt.show()
```



In [151]:

data.charges.corr(data.bmi)

Out[151]:

0.19834096883362884

```
In [152]:
```

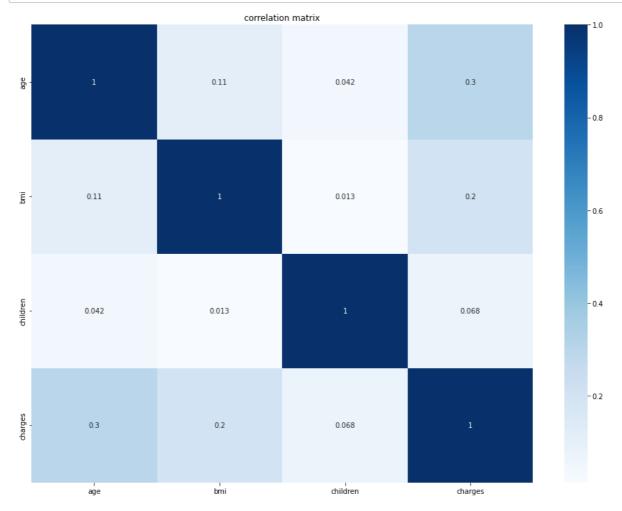
```
data.corr()
```

Out[152]:

	age	bmi	children	charges
age	1.000000	0.109272	0.042469	0.299008
bmi	0.109272	1.000000	0.012759	0.198341
children	0.042469	0.012759	1.000000	0.067998
charges	0.299008	0.198341	0.067998	1.000000

In [153]:

```
plt.figure(figsize=(16,12))
sns.heatmap(data=data.corr(), annot=True, cmap='Blues')
plt.title("correlation matrix")
plt.show()
```



Dataframe with non smokers

```
In [154]:
```

```
non_smoker_data = data[data.smoker=='no']
non_smoker_data
```

Out[154]:

	age	sex	bmi	children	smoker	region	charges
1	18	male	33.770	1	no	southeast	1725.55230
2	28	male	33.000	3	no	southeast	4449.46200
3	33	male	22.705	0	no	northwest	21984.47061
4	32	male	28.880	0	no	northwest	3866.85520
5	31	female	25.740	0	no	southeast	3756.62160
1332	52	female	44.700	3	no	southwest	11411.68500
1333	50	male	30.970	3	no	northwest	10600.54830
1334	18	female	31.920	0	no	northeast	2205.98080
1335	18	female	36.850	0	no	southeast	1629.83350
1336	21	female	25.800	0	no	southwest	2007.94500

1064 rows × 7 columns

Linear regression with one feature

```
In [155]:
```

```
from sklearn.linear_model import LinearRegression
```

```
In [156]:
```

```
model = LinearRegression()
```

```
In [157]:
```

```
inputs = non_smoker_data[['age']]
targets = non_smoker_data['charges']
print(inputs.shape, targets.shape)
```

```
(1064, 1) (1064,)
```

In [158]:

```
model.fit(inputs, targets)
```

Out[158]:

LinearRegression()

Calculating Root Mean Squared Error (RMSE)

```
In [159]:
import numpy as np
def rmse(actual, predict):
    return np.sqrt(np.mean(np.square(actual-predict)))
In [160]:
predictions = model.predict(inputs)
In [161]:
rmse(targets, predictions)
Out[161]:
4662.505766636391
In [162]:
inputs = non_smoker_data[['age', 'bmi']]
inputs.shape
Out[162]:
(1064, 2)
In [163]:
model.fit(inputs, targets)
Out[163]:
LinearRegression()
In [164]:
predictions = model.predict(inputs)
In [165]:
rmse(targets, predictions)
Out[165]:
4662.312835461297
In [166]:
print("Accuracy:",model.score(inputs, targets)*100,"%")
Accuracy: 39.43672876561962 %
```

Linear regression with only one feature fails to capture the whole picture, which was expected. We move on to creating a model with all features included.

Encoding categorical values with mapping and one hot encoder

```
In [167]:
```

```
from sklearn import preprocessing
hot_encoder = preprocessing.OneHotEncoder()
```

In [168]:

```
hot_encoder.fit(data[['region']])
hot_encoder.categories_
```

Out[168]:

```
[array(['northeast', 'northwest', 'southeast', 'southwest'], dtype=object)]
```

In [169]:

```
regions = hot_encoder.transform(data[['region']]).toarray()
regions
```

Out[169]:

In [170]:

```
data[['northeast', 'northwest', 'southeast', 'southwest']] = regions
```

In [171]:

```
data.head()
```

Out[171]:

	age	sex	bmi	children	smoker	region	charges	northeast	northwest	southe
0	19	female	27.900	0	yes	southwest	16884.92400	0.0	0.0	
1	18	male	33.770	1	no	southeast	1725.55230	0.0	0.0	
2	28	male	33.000	3	no	southeast	4449.46200	0.0	0.0	
3	33	male	22.705	0	no	northwest	21984.47061	0.0	1.0	
4	32	male	28.880	0	no	northwest	3866.85520	0.0	1.0	
4										•

In [172]:

```
smokers = {'no':0, 'yes':1}
data['smoker_code'] = data.smoker.map(smokers)
data.head()
```

Out[172]:

	age	sex	bmi	children	smoker	region	charges	northeast	northwest	southe
0	19	female	27.900	0	yes	southwest	16884.92400	0.0	0.0	
1	18	male	33.770	1	no	southeast	1725.55230	0.0	0.0	
2	28	male	33.000	3	no	southeast	4449.46200	0.0	0.0	
3	33	male	22.705	0	no	northwest	21984.47061	0.0	1.0	
4	32	male	28.880	0	no	northwest	3866.85520	0.0	1.0	

→

In [173]:

```
gender = {'female':0, 'male':1}
data['sex_code'] = data.sex.map(gender)
data.head()
```

Out[173]:

	age	sex	bmi	children	smoker	region	charges	northeast	northwest	southe
0	19	female	27.900	0	yes	southwest	16884.92400	0.0	0.0	
1	18	male	33.770	1	no	southeast	1725.55230	0.0	0.0	
2	28	male	33.000	3	no	southeast	4449.46200	0.0	0.0	
3	33	male	22.705	0	no	northwest	21984.47061	0.0	1.0	
4	32	male	28.880	0	no	northwest	3866.85520	0.0	1.0	
4										•

In [174]:

```
inputs2, targets = data[['age','bmi','children', 'smoker_code', 'northeast', 'northwest', '
model.fit(inputs2, targets)
```

Out[174]:

LinearRegression()

In [175]:

```
predictions = model.predict(inputs2)
rmse(targets, predictions)
```

Out[175]:

6042.033215394106

```
In [176]:
```

```
data.head()
```

Out[176]:

	age	sex	bmi	children	smoker	region	charges	northeast	northwest	southe
0	19	female	27.900	0	yes	southwest	16884.92400	0.0	0.0	
1	18	male	33.770	1	no	southeast	1725.55230	0.0	0.0	
2	28	male	33.000	3	no	southeast	4449.46200	0.0	0.0	
3	33	male	22.705	0	no	northwest	21984.47061	0.0	1.0	
4	32	male	28.880	0	no	northwest	3866.85520	0.0	1.0	
4										>

Scaling numerical columns

```
In [177]:
```

```
scaler = preprocessing.StandardScaler()
columns_to_scale = ['age', 'bmi', 'children']
scaler.fit(data[columns_to_scale])
```

Out[177]:

StandardScaler()

In [178]:

```
scaled_columns = scaler.transform(data[columns_to_scale])
scaled_columns
```

Out[178]:

Inputs for model after scaling and encoding

```
In [179]:
```

```
encoded_columns = ['sex_code', 'smoker_code', 'northeast', 'northwest', 'southeast', 'south
categorical_data = data[encoded_columns].values
numerical_data = data[columns_to_scale].values
```

In [180]:

```
inputs2 = np.concatenate((categorical_data, numerical_data), axis=1)
targets = data.charges
```

```
In [181]:
model = LinearRegression()

In [182]:
model.fit(inputs2, targets)

Out[182]:
LinearRegression()

In [183]:
predictions = model.predict(inputs2)

In [184]:
rmse(targets, predictions)

Out[184]:
6041.679651174457

In [185]:
print("Accuracy:",model.score(inputs2, targets)*100,"%")

Accuracy: 75.09130345985207 %
```

Due to a high number of outliers in this dataset, we only achieve an accuracy of 75%, which can be improved further with optimisations