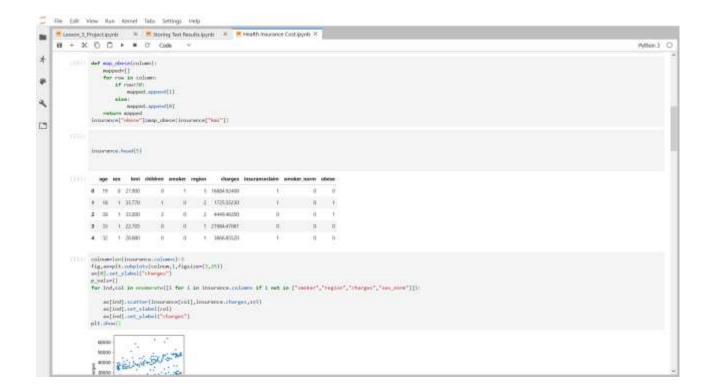
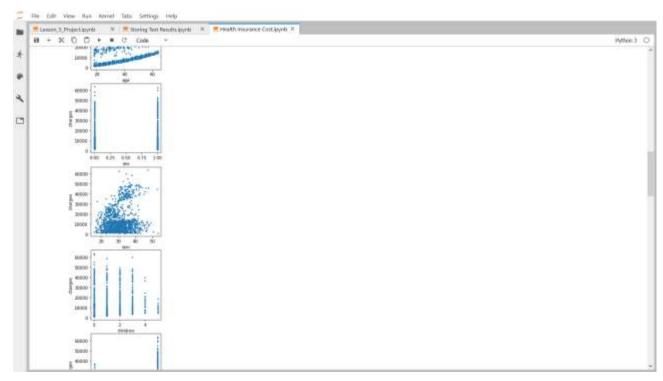
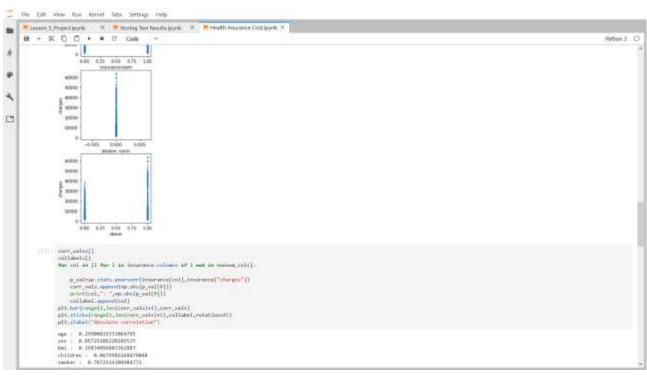
SCREENSHOTS







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    yrique_comPage(*inbusity_comPabesity)*(sembing_comPameking)**reg_intercept
    return y
                       200
```

SOURCE CODE

```
import pandas as pd
import numpy as np
import scipy as sp
import sklearn as sk
```

```
import matplotlib.pyplot as plt
%matplotlib inline
from sklearn.model_selection import cross_val_score, KFold
from sklearn import model_selection
from sklearn import linear_model
from sklearn.metrics import mean_squared_error,mean_absolute_error
```

Loading the data

```
insurance=pd.read csv("insurance2.csv")
insurance.info()
def map smoking(column):
    mapped=[]
    for row in column:
        if row=="yes":
            mapped.append(1)
        else:
            mapped.append(0)
    return mapped
insurance["smoker norm"] = map smoking(insurance["smoker"])
nonnum cols=[col for col in insurance.select dtypes(include=["object"])]
def map obese(column):
    mapped=[]
    for row in column:
        if row>30:
            mapped.append(1)
        else:
            mapped.append(0)
    return mapped
insurance["obese"] = map obese (insurance["bmi"])
insurance.head(5)
```

We now explore the relation between the features given and the insurance costs

```
colnum=len(insurance.columns)-3
fig,ax=plt.subplots(colnum,1,figsize=(3,25))
ax[0].set_ylabel("charges")
p_vals={}
for ind,col in enumerate([i for i in insurance.columns if i not in
["smoker","region","charges","sex_norm"]]):
    ax[ind].scatter(insurance[col],insurance.charges,s=5)
    ax[ind].set_xlabel(col)
    ax[ind].set_ylabel("charges")
plt.show()

corr_vals=[]
collabel=[]
for col in [i for i in insurance.columns if i not in nonnum cols]:
```

```
p val=sp.stats.pearsonr(insurance[col],insurance["charges"])
    corr vals.append(np.abs(p val[0]))
    print(col,": ",np.abs(p val[0]))
    collabel.append(col)
plt.bar(range(1, len(corr vals) +1), corr vals)
plt.xticks(range(1,len(corr vals)+1),collabel,rotation=45)
plt.ylabel("Absolute correlation")
cols not reg3=['age', 'obese', 'smoker norm']
seed=2
results = []
names = []
for name, model in models:
    kfold = KFold(10, random state=None)
    cv results = cross val score (model, insurance[cols not reg3],
insurance["charges"], cv=kfold)
    results.append(cv results)
    names.append(name)
    msg = "%s: %f (%f)" % (name, np.mean(cv results), np.std(cv results))
    print(msg)
kf=KFold(n splits=10, random state=1, shuffle=True)
intercepts=[]
mses=[]
coefs=[]
for train index, test index in kf.split(insurance[cols not reg3]):
    lr=linear model.LinearRegression()
lr.fit(insurance[cols not reg3].iloc[train index],insurance["charges"].iloc[trai
n index])
    lr predictions=lr.predict(insurance[cols not reg3].iloc[test index])
lr mse=mean squared error(insurance["charges"].iloc[test index],lr predictions)
    intercepts.append(lr.intercept )
    coefs.append(lr.coef)
    mses.append(lr mse)
rmses=[x**.5 for x in mses]
avg rmse=np.mean(rmses)
avg intercept=np.mean(intercepts)
age coefs=[]
obesity coefs=[]
smoking_coefs=[]
for vals in coefs:
    #print vals[0]
    age coefs.append(vals[0])
    obesity coefs.append(vals[1])
    smoking coefs.append(vals[2])
age coef=np.mean(age coefs)
```

```
obesity_coef=np.mean(obesity_coefs)
smoking_coef=np.mean(smoking_coefs)
print("a: ",age_coef," b: ",obesity_coef," c: ",smoking_coef," intercept:
",avg_intercept)
```