In [2]:	Medical Cost Personal Insurance Project import pandas as pd import numpy as np import matplotlib.pyplot as plt %matplotlib inline import seaborn as sns
In [3]: Out[3]:	<pre>import warnings warnings.filterwarnings('ignore') df= pd.read_csv('medical_cost_insurance.csv') df.head() age</pre>
In [4]: Out[4]:	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 df.tail() age sex bmi children smoker region charges 1333 50 male 30.97 3 no northwest 10600.5483
	1334 18 female 31.92 0 no northeast 2205.9808 1335 18 female 36.85 0 no southeast 1629.8335 1336 21 female 25.80 0 no southwest 2007.9450 1337 61 female 29.07 0 yes northwest 29141.3603
	(1338, 7) df.describe() 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
	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% 39.00000 30.400000 1.000000 9382.033000 75% 51.000000 34.693750 2.000000 16639.912515 max 64.000000 53.130000 5.000000 63770.428010
In [7]:	age int64 sex object bmi float64 children int64 smoker object region object charges float64 dtype: object
In [8]: Out[8]:	df.isnull().sum() age
In [9]: Out[9]:	<pre>sns.set(style='whitegrid') f, ax = plt.subplots(1,1, figsize=(12, 8)) ax = sns.distplot(df['charges'], kde = True, color = 'c') plt.title('Distribution of Charges')</pre> Text(0.5, 1.0, 'Distribution of Charges')
	Te-5 Distribution of Charges
	5 Shift 4
	1 0 -10000 0 10000 20000 30000 40000 50000 60000 70000 charges
In [10]:	f, ax = plt.subplots(1, 1, figsize=(12, 8)) ax = sns.distplot(np.log10(df['charges']), kde = True, color = 'r') 1.2
	0.8
	0.6 0.4
	0.0 3.0 3.5 4.0 4.5 5.0
In [11]:	#Now we are calculating total charges by region column #sorting it by ascending value and creating a bar plot to visualize the top few regions with the lowest charges charges = df['charges'].groupby(df.region).sum().sort_values(ascending =True) f, ax = plt.subplots(1, 1, figsize=(8,6)) ax = sns.barplot(x=charges.head(), y=charges.head().index, palette= 'Blues')
	northwest
	northeast
In [12]:	southeast 0 1 2 3 4 5 charges 1e6 #Now I want to compare the charges across different regions with the bars further grouped by sex using hue parameter f, ax = plt.subplots(1, 1, figsize=(12, 8)) ax = sns.barplot(x='region', y='charges', hue='sex', data=df, palette='cool')
	17500 Sex female male
	12500 89 10000
	7500 — 5000 — 6 6 6 6 6 6 6 6 6 6 6 6 6 6 6 6 6
	2500 southwest southeast northwest northeast region
In [13]:	# Now i want to create a bar plot to compare the charges across different regions with the bars grouped by smoker using the hue parameter f, ax = plt.subplots(1,1, figsize=(12,8)) ax = sns.barplot(x = 'region', y = 'charges', hue='smoker', data=df, palette='Reds_r') smoker yes no
	25000
	15000 — 10000
	5000 southwest southeast northwest northeast
In [14]:	
	15000 10000
	5000
In [15]:	southwest southeast northwest northeast region #Now i am creating three separate scatter plots to visualize the relationship between age bmi children with charges ax = sns.lmplot(x = 'age', y = 'charges', data=df, hue='smoker', palette= 'Set1') ax = sns.lmplot(x = 'bmi', y = 'charges', data=df, hue='smoker', palette= 'Set2') ax = sns.lmplot(x = 'children', y = 'charges', data=df, hue='smoker', palette= 'Set3')
	50000
	40000 20000 smoker yes no
	10000 20 30 40 50 60 age 70000
	60000 50000 40000 smoker
	\$\frac{40000}{66} \frac{30000}{20000} \frac{50000}{10000} \frac{5000}{10000} \frac{5000}{10000} \frac{50000}{10000} \frac{50000}{10000} \frac{50000}{10000} \frac{50000}{10000} \qquad \
	20 30 40 50 bmi
	50000 40000 \$\frac{\text{smoker}}{\text{yes}} \text{no}
	20000 10000 0 1 2 3 4 5
In [16]:	#Now I am using violiplot to visualize the distribution of charges on the number of children f, ax = plt.subplots(1,1, figsize=(10, 10)) ax = sns.violinplot(x = 'children', y = 'charges', data=df, orient= 'v', hue='smoker', palette= 'inferno') smoker yes 70000
	60000 50000
	des charges ch
	20000
In [17]: Out[17]:	#Converting objects labels into categorical df[['sex', 'smoker', 'region']] = df[['sex', 'smoker', 'region']].astype('category') df.dtypes age int64 sex category bmi float64 children int64 smoker category
In [18]:	region category charges float64 dtype: object ##Converting category labels into numerical using LabelEncoder from sklearn.preprocessing import LabelEncoder label = LabelEncoder() label.fit(df.sex.drop_duplicates()) df.sex = label.transform(df.sex) label.fit(df.smoker.drop_duplicates())
Out[18]:	<pre>df.smoker = label.transform(df.smoker) label.fit(df.region.drop_duplicates()) df.region = label.transform(df.region) df.dtypes age int64 sex int32 bmi float64 children int64 smoker int32 region int32</pre>
In [19]:	charges float64 dtype: object f, ax = plt.subplots(1, 1, figsize=(10, 10)) ax = sns.heatmap(df.corr(), annot= True, cmap= 'cool') 1 -0.021 0.11 0.042 -0.025 0.0021 0.3
	-0.021 1 0.046 0.017 0.076 0.0046 0.057 -0.8
	[
	1
	89 0.3 0.057 0.2 0.068 0.79 -0.0062 1 -0.00
In [20°	Training model Linear Regression # Importing the necessary libraries from sklearn model selection import train test split as holdout
[∠0]:	<pre>from sklearn.model_selection import train_test_split as holdout from sklearn.linear_model import LinearRegression from sklearn import metrics #Load the dateset x = df.drop(['charges'], axis=1) y = df['charges'] #Split the dataset inot training anf testing Datasets x_train,x_test,y_train,y_test = holdout(x,y, test_size=0.2, random_state=0)</pre>
	#create a Linear Regression model Lin_reg = LinearRegression() Lin_reg.fit(x_train, y_train) #Print the Intercept, Coefficent , and score of the model print(Lin_reg.intercept_) print(Lin_reg.coef_) print(Lin_reg.score(x_test, y_test)) -11661.983908824424 [253.99185244 -24.32455098 328.40261701 443.72929547 23568.87948381 -288.50857254]
In [21]:	23568.87948381 -288.50857254] 0.7998747145449959 Ridge Regression #Importing the required library from sklearn.linear_model import Ridge #Creating the Ridge Regression Model
	Ridge = Ridge(alpha=0.5) #Train the ridge Regression Model Ridge.fit(x_train, y_train) #Print the intercept, coeff, and score of the model print(Ridge.intercept_) print(Ridge.coef_) print(Ridge.score(x_test, y_test)) -11643.440927495836 [2.53893751e+02 -2.15112284e+01 3.28339566e+02 4.44238477e+02
In [22]:	[2.53893751e+02 -2.15112284e+01
	Lasso = Lasso(alpha=0.2, fit_intercept=True, precompute=False, max_iter=1000,
In [23]:	-11661.838929039533 [2.53991436e+02 -2.34569821e+01
	<pre>from sklearn.ensemble import RandomForestRegressor as rfr #load the dataset x= df.drop(['charges'],axis = 1) y= df.charges #Create a Random Forest Regression Rfr = rfr(n_estimators = 100, criterion = 'squared_error',</pre>
	<pre>Rfr.fit(x_train,y_train) #Make predictions of the model x_train_pred= Rfr.predict(x_train) x_test_pred= Rfr.predict(x_test) #print the mean square of the model print('MSE train data: %.3f, MSE test data: %.3f' %</pre>
In [24]:	<pre>Plt.figure(figsize=(8,6)) plt.scatter(x_train_pred, x_train_pred - y_train,</pre>
Out[24]:	<pre>label = 'Test data') plt.xlabel('Predicted values') plt.ylabel('Actual values') plt.legend(loc = 'upper right') plt.hlines(y= 0, xmin =0 , xmax =6000, lw =2, color = 'red') <matplotlib.collections.linecollection 0x240e4574a60="" at=""></matplotlib.collections.linecollection></pre> 20000 Train data
	15000 10000 5000
	-5000 -15000
In [33]:	-20000 0 10000 20000 30000 40000 50000 60000 Predicted values print('Feature importance ranking\n\n') importances = Rfr.feature_importances_ std = np.std([tree.feature_importances_ for tree in Rfr.estimators_], axis =0)
	<pre>std = np.std([tree.feature_importances_ for tree in Rfr.estimators_], axis =0) indices = np.argsort(importances)[::-1] variables = ['age', 'sex', 'bmi', 'children', 'smoker', 'region'] importance_list = [] for f in range(x.shape[1]): variable = variables[indices[f]] importance_list.append(variable) print("%d.%s(%f)"% (f + 1, variable, importances[indices[f]])) #Plot the feature importance of the forest plt.figure() plt.title("Feature importances")</pre>
	<pre>plt.title("Feature importances") plt.bar(importance_list, importances[indices],</pre>
Out[33]:	
	0.4 0.3 0.2
	0.1 smoker bmi age children region sex
In [36]:	#Importing the library from sklearn.preprocessing import PolynomialFeatures #Load the Data set x = df.drop(['charges', 'sex', 'region'], axis =1) y = df.charges #Create polynomial features pol = PolynomialFeatures (degree = 2)
	<pre>pol = PolynomialFeatures (degree = 2) x_pol = pol.fit_transform(x) #Split the dataset into training and testing x_train, x_test, y_train, y_test = holdout(x_pol, y, test_size=0.2, random_state=0) #Create the polynomial regression model Pol_reg = LinearRegression() Pol_reg.fit(x_train, y_train) #make predictions of the model</pre>
	<pre>y_train_pred = Pol_reg.predict(x_train) y_test_pred = Pol_reg.predict(x_test) #print the intercept, coeff, and score of the model print(Pol_reg.intercept_) print(Pol_reg.coef_) print(Pol_reg.score(x_test, y_test)) -5325.881705252405 [0.00000000e+00 -4.01606591e+01 5.23702019e+02 8.52025026e+02 -9.52698471e+03 3.04430186e+00 1.84508369e+00 6.01720286e+00</pre>
In [37]:	#Evaluating the performance of the algorithm print('Mean Absolute Error:', metrics.mean_absolute_error(y_test, y_test_pred)) print('Mean Squared Error:', np.sqrt(metrics.mean_squared_error(y_test, y_test_pred))) Mean Absolute Error: 2824.4950454776426 Mean Squared Error: 18895160.09878039 Root Mean Squared Error: 4346.856346692445
<pre>In [39]: Out[39]:</pre>	
	610 8547.69130 10440.782266 569 45702.02235 48541.022951 1034 12950.07120 14140.067522 198 9644.25250 8636.235727 1084 15019.76005 16712.196281 726 6664.68595 8654.565461 1132 20709.02034 12372.050609
	1132 20709.02034 12372.050609 725 40932.42950 41465.617268 963 9500.57305 10941.780705