In [1]: In [2]:	<pre>import libraries import pandas as pd # Data manipulation import numpy as np # Linear algebra import warnings # Ignore warnings import seaborn as sns #plots import matplotlib.pyplot as plt # plots warnings.filterwarnings("ignore")</pre>
<pre>In [3]: In [4]: Out[4]:</pre>	0 19 female 27.900 0 yes southwest 16884.92400 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
In [5]: Out[5]: In [137	'sex': ['temale', 'male'], 'smoker': ['yes', 'no']}
In [138 In [133	p=s_p=scs : =s()
In [134	northwest 24.29%(325) 24.22%(324) 27.20%(364) 24.29%(325) southwest pie_plots.smoker_pie() Smoker Percent
In [139	79.52%(1064) 20.48%(274) yes
	Sex Percent female 49.48%(662) 50.52%(676) male
In [140 Out[140 In [141 In [142	1 324 2 240 3 157 4 25 5 18 Name: children, dtype: int64 sns.set_style(style="whitegrid")
	<pre>fig ax=plt.subplots(1,1,figsize=(20,1)) ax.set_title(title) ax.hist(df[feature],ec="k",color="MFADAGE",lw=1) ax.axvline(df[feature].mean(),</pre>
In [143	histogram("charges") Charges One of the control o
In [145	The price of insurance is higher for people who smoke? (ggplot(df) + aes(x="smoker",y="charges",fill="smoker") + geom_boxplot() + labs(title="Smoker vs Charges") + facet_wrap("smoker") + theme(legend_position="none") + scale_fill_manual(values=["#90ee90","#ffoccb"])
	Smoker vs Charges 40000 20000 20000 yes smoker
In [146 Out[146	we observe a strong presence of outliers, for the category of non-smokers. df.groupby("smoker")["charges"].mean() smoker no 8434.268298 yes 32050.231832 yame: charges, dtype: float64 The average price of smokers is considerably much higher than non-smokers. Since they usually have poorer physical health and need more treatments. histogram("bmi", "BNI")
	BMI 300 250 150 200 150 200 250 250 2
In [148	Most of the BMI data is within a normal distribution. But even so, it is possible to appreciate outlier values in the upper range. from plotnine.facets import facet_grid from plotnine.geoms import geom_smooth def scatter_plots(feature,title): return((ggplot(df) + aes(x=feature,y="charges",fill="smoker",alpha=0.1) + geom_point() + labs(title=title,x=feature) + facet_wrap("smoker") + theme(legend_position="none") + scale_fill_manual(values=["#90ee90","#ffcccb"])
In [149	People with a high BMI the insurance charge is higher? Scatter_plots "Inst", "BMI vs Charges") BMI vs Charges
Out[149 In [150	eggplot: (8763329574941)> For non-smokers, the data trend remains constant. While for smokers the trend line is linear, that is, one value increases proportionally with another. Does age influence the price of insurance? scatter_plots("age", "Age vs Charges")
	Age vs Charges
Out[150	 For non-smokers, the trend of the data remains linear with respect to age. So we can create a linear regression model to model the data for non-smokers. But with the disadvantage that it would not explain so well for the opposite case. For smokers, the relationship between age seems non-existent. corr_pearson=df.corr(method="pearson") corr_spearman=df.corr(method="spearman") corr_lation_matrix(corr_matrix): plt.figure(figsize=(15, 5)) sns.heatmap(corr_matrix, annot="ruc")
	Correlation Matrix It is used to establish possible relationships between variables Pearson correlation_matrix(corr_pearson)
	1
In [153	Correlation_matrix(corr_spearman)
	96 0057 0016 1 013 013 0 02 013 0 012 0 013 1 0 02 0 02