Conductivity 0 Organic_carbon 0 Trihalomethanes 162 Turbidity 0 Potability 0 dtype: int64  data=data.dropna()			
ph 0 Hardness 0 Solids 0 Chloramines 0 Sulfate 0 Conductivity 0 Organic_carbon 0 Trihalomethanes 0 Turbidity 0 Potability 0 dtype: int64			
<pre>plt.figure(figsize=(15,10)) sns.countplot(data.Potability) plt.title("Distribution of unsafe and sa plt.show()  C:\Users\lenovo\anaconda3\lib\site-packag argument will be `data`, and passing othe warnings.warn(</pre>	fe water")  des\seaborn\_decorators.py:36: FutureWarning: Pass the following or arguments without an explicit keyword will result in an error  Distribution of unsafe and safe water	variable as a keyword arg: x. From version 0.12, the only valor misinterpretation.	lid posi
1000 - 800 - 400 - 200 -			
<pre>import plotly.express as px  data = data figure = px.histogram(data, x ="ph",</pre>	Potability  ", fecting water quality: PH")		
Factors affecting water quality: PF			Potak
100 80 60 40			
data = data figure = px.histogram(data, x ="Hardness color = "Potability		10 12 14	
figure.show()  Factors affecting water quality: Ha	fecting water quality: Hardness")		Potak
140 120 100 80 60			•
20 100	150 200 Hardness	250 300	
<pre>data = data figure = px.histogram(data, x ="Solids",</pre>	", fecting water quality: Solids")		<b>-</b> [x] <b>^</b>
100 80 60			Potak
20 0 0 10k	20k 30k Solids	40k 50k	
<pre>data = data figure = px.histogram(data, x = "Chloramic color = "Potability title = "Factors af figure.show()</pre> Factors affecting water quality: Ch	", fecting water quality: Chloramines")		<b>-</b> [X] <b>^</b>
120 100 80			Potak
60 40 20 0 2	4 6 8	10 12	
<pre>data = data figure = px.histogram(data, x ="Sulfate"</pre>	Chloramines  ', ", fecting water quality: Sulfate")		
Factors affecting water quality: Su	Ifate		Potak —
40 20			
<pre>data = data figure = px.histogram(data, x ="Conduction color = "Potability")</pre>		350 400 450	
Factors affecting water quality: Co	nductivity		Potak
40 20			
data = data figure = px.histogram(data, x ="Organic_color = "Potability	400 500  Conductivity  carbon", ", fecting water quality: Organic_carbon")	600 700	
Factors affecting water quality: Or			Potak
100 80 60 40			
data = data figure = px.histogram(data, x ="Trihalom	10 15 Organic_carbon	20 25	
color = "Potability	", fecting water quality: Trihalomethanes")		Potak
100 80 60 40			
20 20	40 60  Trihalomethanes	80 100 120	
ph         1.000000         0.108948         -0.087615           Hardness         0.108948         1.000000         -0.053269           Solids         -0.087615         -0.053269         1.000000	hloramines         Sulfate         Conductivity         Organic_carbon         Trihalomethanes         Turbidity           -0.024768         0.010524         0.014128         0.028375         0.018278         -0.035849           -0.022685         -0.108521         0.011731         0.013224         -0.015400         -0.034831           -0.051789         -0.162769         -0.005198         -0.005484         -0.015668         0.019409	0.014530 -0.001505 0.040674	
Chloramines -0.024768 -0.022685 -0.051789  Sulfate 0.010524 -0.108521 -0.162769  Conductivity 0.014128 0.011731 -0.005198  Organic_carbon 0.028375 0.013224 -0.005484  Trihalomethanes 0.018278 -0.015400 -0.015668  Turbidity -0.035849 -0.034831 0.019409  Potability 0.014530 -0.001505 0.040674	1.000000       0.006254       -0.028277       -0.023808       0.014990       0.013137         0.006254       1.000000       -0.016192       0.026776       -0.023347       -0.009934         -0.028277       -0.016192       1.000000       0.015647       0.004888       0.012495         -0.023808       0.026776       0.015647       1.000000       -0.005667       -0.015428         0.014990       -0.023347       0.004888       -0.005667       1.000000       -0.020497         0.013137       -0.009934       0.012495       -0.015428       -0.020497       1.000000         0.020784       -0.015303       -0.015496       -0.015567       0.009244       0.022682	-0.015303 -0.015496 -0.015567 0.009244 0.022682	
<pre>x = data.drop(["Potability"], axis= 1)  from sklearn.preprocessing import Standa  scaler = StandardScaler() scaler.fit(X)  StandardScaler()</pre>	rdScaler		
from sklearn.model_selection import train from sklearn.metrics import confusion_ma from sklearn.linear_model import Logistic from sklearn.linear_model import LinearRefrom sklearn.svm import SVC from sklearn.naive_bayes import Gaussian from sklearn.neural_network import MLPClafrom sklearn.tree import DecisionTreeClafrom sklearn.ensemble import RandomFores from warnings import filterwarnings filterwarnings ('ignore')	trix, accuracy_score, classification_report, precision_score cRegression egression NB assifier ssifier		
<pre>X_train, X_test, y_train, y_test = train  models =[("LogisticReg", LogisticRegress</pre>	<pre>test_size=0.2,     random_state=42)  ion(max_iter=1000)),("SVC", SVC()), assifier()),("NaiveBayes", GaussianNB()), ssifier()),('GradientBoost',GradientBoostingClassifier()),</pre>		
<pre>results = [] names = [] accuracies = [] finalResults = []  for name, model in models:     model.fit(X_train, y_train)     y_pred = model.predict(X_test)     score = precision_score(y_test, y_predict)     accuracyScore = accuracy_score(y_test)     accuracies.append(accuracyScore)</pre>			
<pre>results.append(score) names.append(name) finalResults.append((name, score, acc)</pre>	ame': names, 'Accuracy-Score': accuracies, 'Precision-Score': re	sults})	
<pre>2 DecisionTree</pre>			
<pre>"max_features": [3, 5, 7, 1]</pre>	0, 11], 180, 200, 300, 500], 3, 15, 16, 18]}  , cv=10, n_jobs=-1, verbose=2)  es, totalling 7500 fits tClassifier(), n_jobs=-1, 2, 14, 15, 17, 19],		
<pre>param_grid={'max_depth': [12</pre>	[3, 5, 7, 10, 11], [1, 13, 15, 16, 18], [100, 150, 180, 200, 300, 500]},		
	<pre>pth = 12, max_features = 7, min_samples_split = 10, n_estimators features=8, min_samples_split=10)</pre>	= 100)	
<pre>RandomForestClassifier(max_depth=10, max_  y_pred = rf_tuned.predict(X_test) accuracy_score(y_test, y_pred)  0.674937965260546</pre>			