

```
In [1]: import numpy as np
import pandas as pd
import matplotlib.pyplot as plt
import seaborn as sns
```

```
In [3]: data = pd.read_csv("C:/Users/iafri/Documents/ds practice/water_potability.csv")
```

```
In [4]: data.head()
```

	ph	Hardness	Solids	Chloramines	Sulfate	Conductivity	Organic_carbon	Trihalomethanes	Turbidity	Potability
0	NaN	204.890455	20791.318981	7.300212	368.516441	564.308654	10.379783	86.990970	2.963135	0
1	3.716080	129.422921	18630.057858	6.635246	NaN	592.885359	15.180013	56.329076	4.500656	0
2	8.099124	224.236259	19909.541732	9.275884	NaN	418.606213	16.868637	66.420093	3.055934	0
3	8.316766	214.373394	22018.417441	8.059332	356.886136	363.266516	18.436524	100.341674	4.628771	0
4	9.092223	181.101509	17978.986339	6.546600	310.135738	398.410813	11.558279	31.997993	4.075075	0

```
In [5]: data.info()

<class 'pandas.core.frame.DataFrame'>
RangeIndex: 3276 entries, 0 to 3275
Data columns (total 10 columns):
#   Column              Non-Null Count  Dtype  
---  -
0   ph                   2785 non-null   float64
1   Hardness             3276 non-null   float64
2   Solids               3276 non-null   float64
3   Chloramines          3276 non-null   float64
4   Sulfate              2495 non-null   float64
5   Conductivity         3276 non-null   float64
6   Organic_carbon       3276 non-null   float64
7   Trihalomethanes      3114 non-null   float64
8   Turbidity            3276 non-null   float64
9   Potability           3276 non-null   int64   
dtypes: float64(9), int64(1)
memory usage: 256.1 KB
```

```
In [6]: data.isnull().sum()
```

ph	491
Hardness	0
Solids	0
Chloramines	0
Sulfate	781
Conductivity	0
Organic_carbon	0
Trihalomethanes	162
Turbidity	0
Potability	0
dtype:	int64

```
In [7]: data.describe()
```

	ph	Hardness	Solids	Chloramines	Sulfate	Conductivity	Organic_carbon	Trihalomethanes	Turbidity	Potability
count	2785.000000	3276.000000	3276.000000	3276.000000	2495.000000	3276.000000	3276.000000	3114.000000	3276.000000	3276
mean	7.080795	196.369496	22014.092526	7.122277	333.775777	426.205111	14.284970	66.396293	3.966786	0
std	1.594320	32.879761	8768.570828	1.583085	41.416840	80.824064	3.308162	16.175008	0.780382	0
min	0.000000	47.432000	320.942611	0.352000	129.000000	181.483754	2.200000	0.738000	1.450000	0
25%	6.093092	176.850538	15666.690297	6.127421	307.699498	365.734414	12.065801	55.844536	3.439711	0
50%	7.036752	196.967627	20927.833607	7.130299	333.073546	421.884968	14.218338	66.622485	3.955028	0
75%	8.062066	216.667456	27332.762127	8.114887	359.950170	481.792304	16.557652	77.337473	4.500320	1
max	14.000000	323.124000	61227.196008	13.127000	481.030642	753.342620	28.300000	124.000000	6.739000	1

```
In [8]: for col in ["ph", "Sulfate", "Trihalomethanes"]:
data[col].fillna(value=data[col].mean(), inplace=True)
```

```
In [9]: # Reassessing Data Quality
data.info()
```

```
<class 'pandas.core.frame.DataFrame'>
RangeIndex: 3276 entries, 0 to 3275
Data columns (total 10 columns):
#   Column                Non-Null Count  Dtype
---  -
0   ph                     3276 non-null   float64
1   Hardness               3276 non-null   float64
2   Solids                 3276 non-null   float64
3   Chloramines            3276 non-null   float64
4   Sulfate                3276 non-null   float64
5   Conductivity           3276 non-null   float64
6   Organic_carbon         3276 non-null   float64
7   Trihalomethanes        3276 non-null   float64
8   Turbidity              3276 non-null   float64
9   Potability             3276 non-null   int64
dtypes: float64(9), int64(1)
memory usage: 256.1 KB
```

```
In [10]: data.isnull().sum()
```

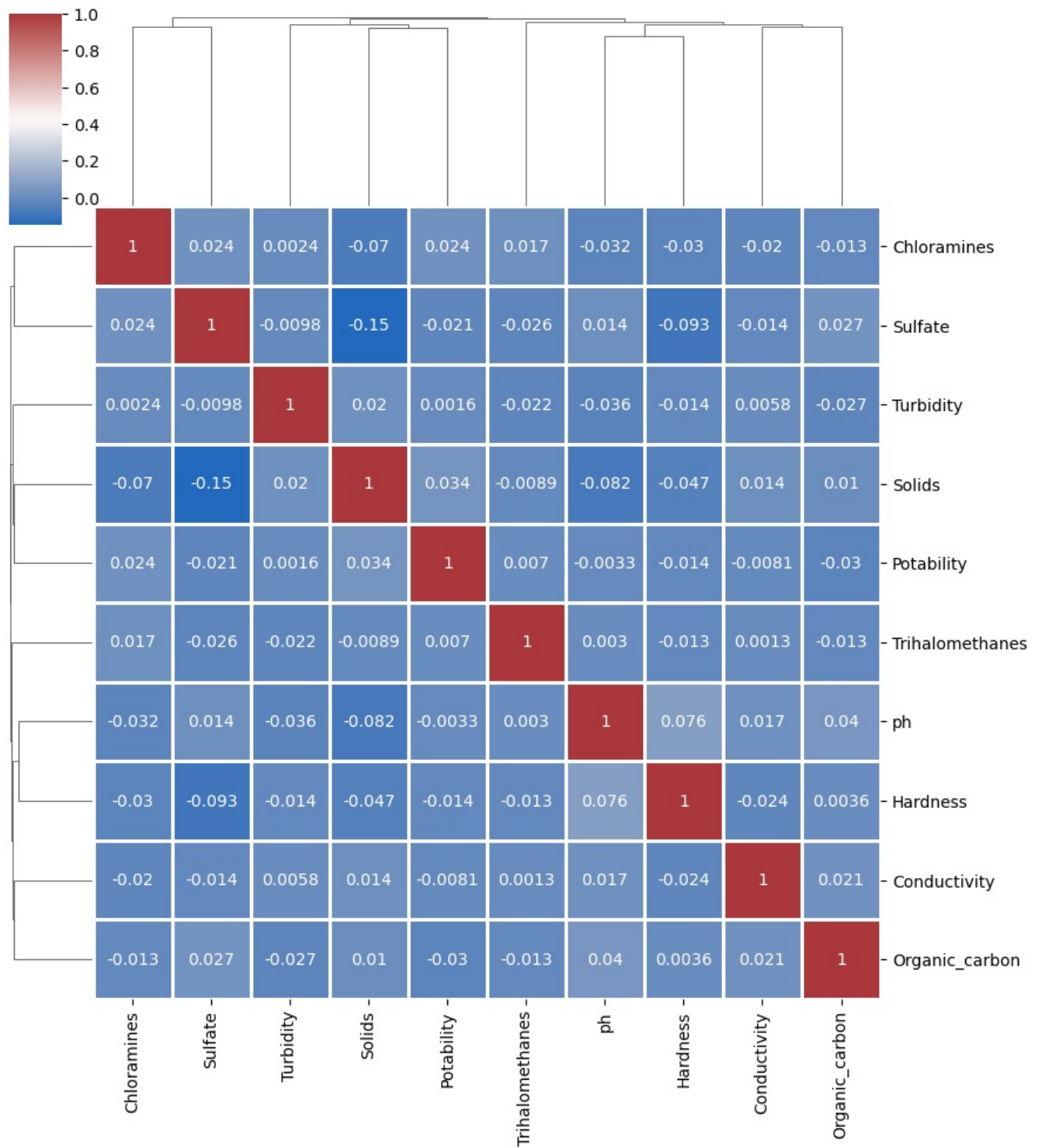
```
Out[10]: ph                0
Hardness              0
Solids                0
Chloramines           0
Sulfate               0
Conductivity          0
Organic_carbon        0
Trihalomethanes       0
Turbidity             0
Potability            0
dtype: int64
```

```
In [11]: data["Potability"].value_counts()
```

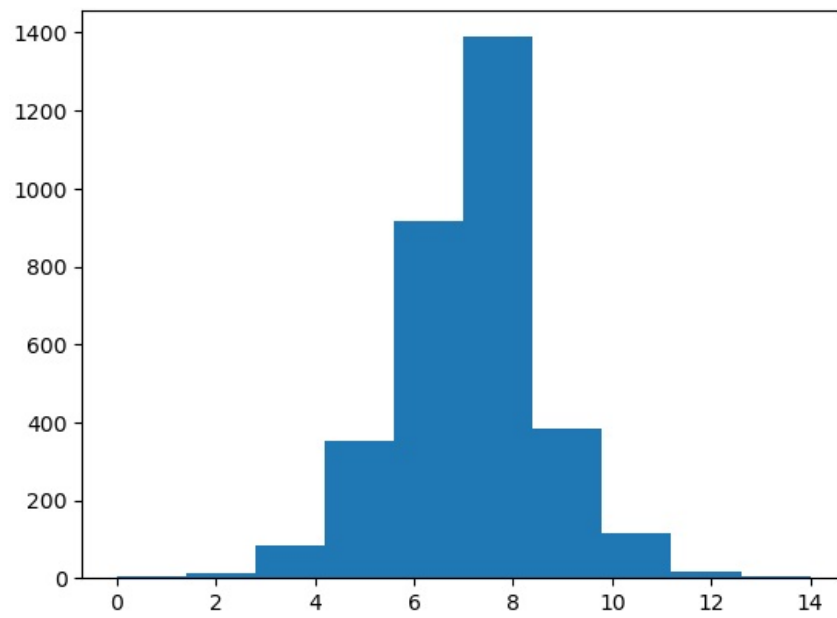
```
Out[11]: Potability
0      1998
1      1278
Name: count, dtype: int64
```

```
In [12]: #Showing the correlation of the features (with missing values)
sns.clustermap(data.corr(), cmap="vlag", dendrogram_ratio=(0.1, 0.2), annot=True, linewidths=.8, figsize=(9, 10))
```

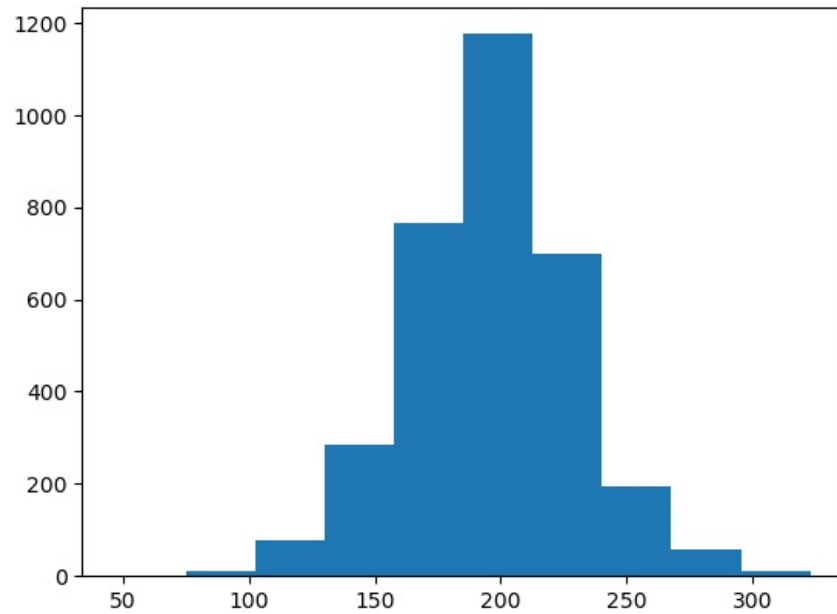
```
Out[12]: <seaborn.matrix.ClusterGrid at 0x214eaf14510>
```



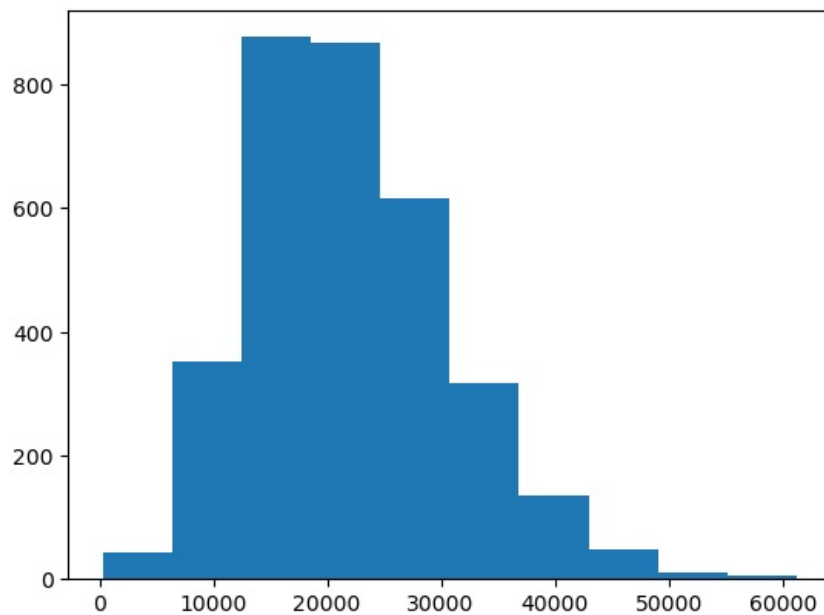
```
In [13]: plt.figure()
plt.hist(data['ph'])
plt.show()
```



```
In [14]: plt.figure()
plt.hist(data['Hardness'])
plt.show()
```

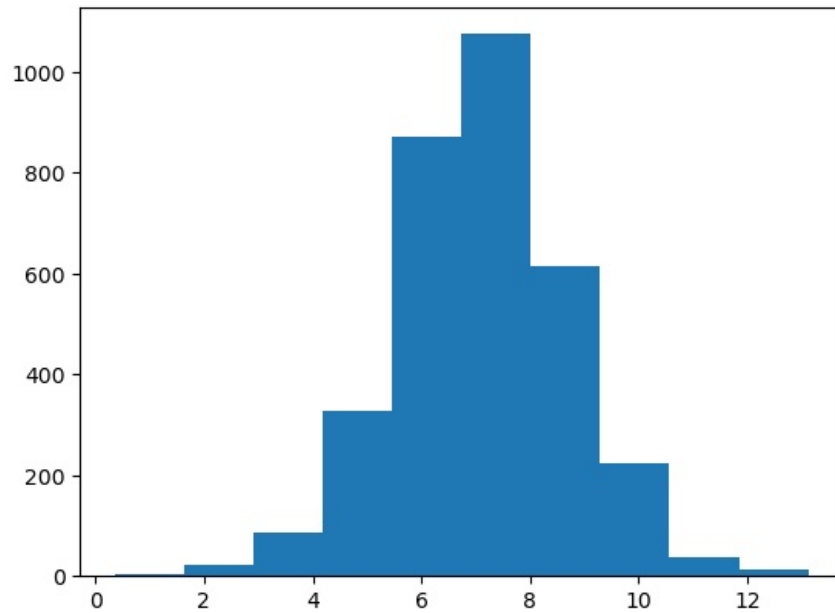


```
In [15]: plt.figure()
plt.hist(data['Solids'])
plt.show()
```

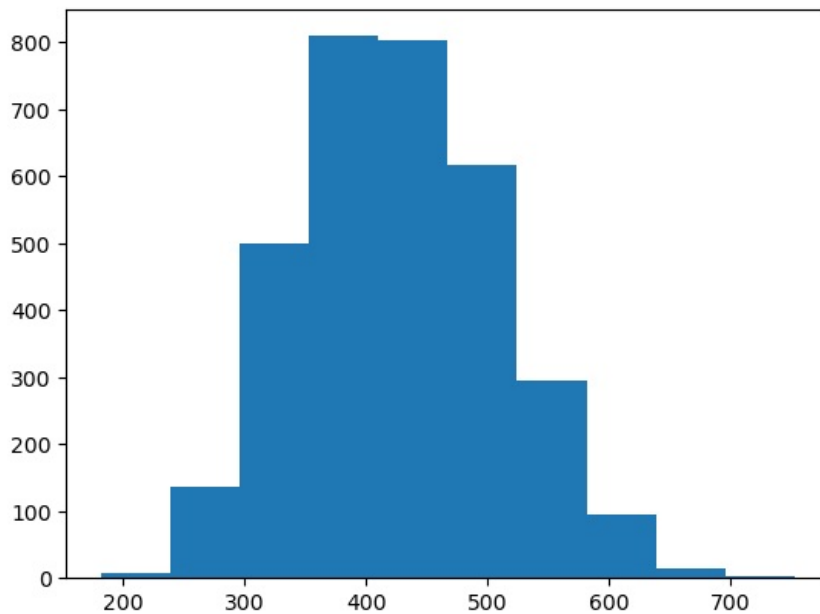


```
In [16]: plt.figure()
```

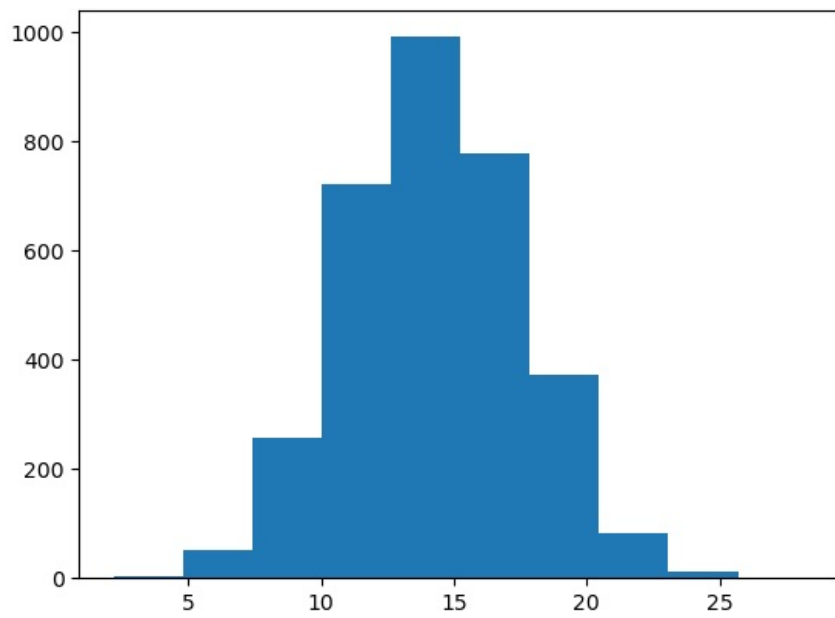
```
In [16]: plt.figure()  
plt.hist(data['Chloramines'])  
plt.show()
```



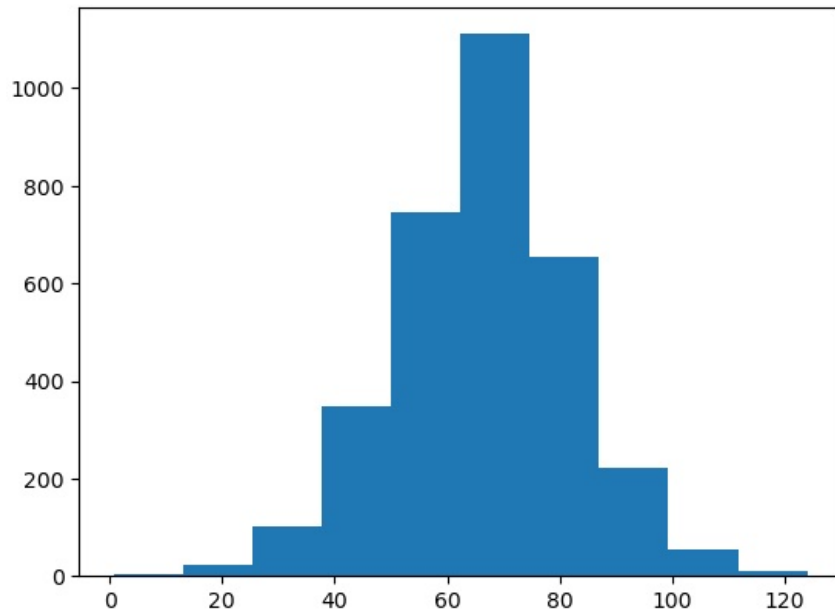
```
In [17]: plt.figure()  
plt.hist(data['Conductivity'])  
plt.show()
```



```
In [18]: plt.figure()  
plt.hist(data['Organic_carbon'])  
plt.show()
```



```
In [19]: plt.figure()
plt.hist(data['Trihalomethanes'])
plt.show()
```



```
In [20]: plt.figure()
plt.hist(data['Turbidity'])
plt.show()
```

