

```
import pandas as pd
import numpy as np
from scipy.stats import zscore
import matplotlib.pyplot as plt
```

```
from sklearn.datasets import load_wine
```

```
# Load the Wine dataset
wine = load_wine()
X = pd.DataFrame(wine.data, columns=wine.feature_names)
y = pd.Series(wine.target, name='target')
```

```
# Combining X and y into a single DataFrame for the analysis
df = X.copy()
df['target'] = y
```

```
# 1. Display the shape of the dataset
shape_original = X.shape
print(shape_original)
```

```
(178, 13)
```

```
# 2. Print the header and the last few rows
header = df.head()
tail = df.tail()
print(header)
print(tail)
```

	alcohol	malic_acid	ash	alcalinity_of_ash	magnesium	total_phenols	\
0	14.23	1.71	2.43	15.6	127.0	2.80	
1	13.20	1.78	2.14	11.2	100.0	2.65	
2	13.16	2.36	2.67	18.6	101.0	2.80	
3	14.37	1.95	2.50	16.8	113.0	3.85	
4	13.24	2.59	2.87	21.0	118.0	2.80	

	flavanoids	nonflavanoid_phenols	proanthocyanins	color_intensity	hue	\
0	3.06	0.28	2.29	5.64	1.04	
1	2.76	0.26	1.28	4.38	1.05	
2	3.24	0.30	2.81	5.68	1.03	
3	3.49	0.24	2.18	7.80	0.86	
4	2.69	0.39	1.82	4.32	1.04	

	od280/od315_of_diluted_wines	proline	target
0	3.92	1065.0	0
1	3.40	1050.0	0
2	3.17	1185.0	0
3	3.45	1480.0	0
4	2.93	735.0	0

	alcohol	malic_acid	ash	alcalinity_of_ash	magnesium	total_phenols	\
173	13.71	5.65	2.45	20.5	95.0	1.68	
174	13.40	3.91	2.48	23.0	102.0	1.80	
175	13.27	4.28	2.26	20.0	120.0	1.59	
176	13.17	2.59	2.37	20.0	120.0	1.65	
177	14.13	4.10	2.74	24.5	96.0	2.05	

	flavanoids	nonflavanoid_phenols	proanthocyanins	color_intensity	hue	\
173	0.61	0.52	1.06	7.7	0.64	
174	0.75	0.43	1.41	7.3	0.70	
175	0.69	0.43	1.35	10.2	0.59	
176	0.68	0.53	1.46	9.3	0.60	
177	0.76	0.56	1.35	9.2	0.61	

	od280/od315_of_diluted_wines	proline	target
173	1.74	740.0	2
174	1.56	750.0	2
175	1.56	835.0	2
176	1.62	840.0	2
177	1.60	560.0	2

```
# 3. Print a summary of the dataset's statistical details
summary_statistics = df.describe()
print(summary_statistics)
```

	alcohol	malic_acid	ash	alcalinity_of_ash	magnesium	\
count	178.000000	178.000000	178.000000	178.000000	178.000000	
mean	13.000618	2.336348	2.366517	19.494944	99.741573	

std	0.811827	1.117146	0.274344	3.339564	14.282484
min	11.030000	0.740000	1.360000	10.600000	70.000000
25%	12.362500	1.602500	2.210000	17.200000	88.000000
50%	13.050000	1.865000	2.360000	19.500000	98.000000
75%	13.677500	3.082500	2.557500	21.500000	107.000000
max	14.830000	5.800000	3.230000	30.000000	162.000000

	total_phenols	flavanoids	nonflavanoid_phenols	proanthocyanins	\
count	178.000000	178.000000	178.000000	178.000000	
mean	2.295112	2.029270	0.361854	1.590899	
std	0.625851	0.998859	0.124453	0.572359	
min	0.980000	0.340000	0.130000	0.410000	
25%	1.742500	1.205000	0.270000	1.250000	
50%	2.355000	2.135000	0.340000	1.555000	
75%	2.800000	2.875000	0.437500	1.950000	
max	3.880000	5.080000	0.660000	3.580000	

	color_intensity	hue	od280/od315_of_diluted_wines	proline	\
count	178.000000	178.000000	178.000000	178.000000	
mean	5.058090	0.957449	2.611685	746.893258	
std	2.318286	0.228572	0.709990	314.907474	
min	1.280000	0.480000	1.270000	278.000000	
25%	3.220000	0.782500	1.937500	500.500000	
50%	4.690000	0.965000	2.780000	673.500000	
75%	6.200000	1.120000	3.170000	985.000000	
max	13.000000	1.710000	4.000000	1680.000000	

	target
count	178.000000
mean	0.938202
std	0.775035
min	0.000000
25%	0.000000
50%	1.000000
75%	2.000000
max	2.000000

```
# 4. Display a concise summary of the dataframe
info_summary = df.info()
print(info_summary)
```

```
<class 'pandas.core.frame.DataFrame'>
RangeIndex: 178 entries, 0 to 177
Data columns (total 14 columns):
#   Column                                Non-Null Count  Dtype
---  -
0   alcohol                               178 non-null    float64
1   malic_acid                           178 non-null    float64
2   ash                                   178 non-null    float64
3   alcalinity_of_ash                    178 non-null    float64
4   magnesium                             178 non-null    float64
5   total_phenols                        178 non-null    float64
6   flavanoids                           178 non-null    float64
7   nonflavanoid_phenols                 178 non-null    float64
8   proanthocyanins                      178 non-null    float64
9   color_intensity                      178 non-null    float64
10  hue                                   178 non-null    float64
11  od280/od315_of_diluted_wines         178 non-null    float64
12  proline                              178 non-null    float64
13  target                               178 non-null    int64
dtypes: float64(13), int64(1)
memory usage: 19.6 KB
None
```

```
# 5. Add an index column and display its new shape
df.reset_index(inplace=True)
shape_with_index = df.shape
print(shape_with_index)
```

```
(178, 15)
```

```
# 6. Choose a numerical field and print its unique values (using 'alcohol' as an example)
unique_values_alcohol = df['alcohol'].unique()
print(unique_values_alcohol)
```

```
[14.23 13.2  13.16 14.37 13.24 14.2  14.39 14.06 14.83 13.86 14.1  14.12
 13.75 14.75 14.38 13.63 14.3  13.83 14.19 13.64 12.93 13.71 12.85 13.5
 13.05 13.39 13.3  13.87 14.02 13.73 13.58 13.68 13.76 13.51 13.48 13.28
 13.07 14.22 13.56 13.41 13.88 14.21 13.9  13.94 13.82 13.77 13.74 13.29
 13.72 12.37 12.33 12.64 13.67 12.17 13.11 13.34 12.21 12.29 13.49 12.99
 11.96 11.66 13.03 11.84 12.7  12.  12.72 12.08 12.67 12.16 11.65 11.64]
```

```

12.69 11.62 12.47 11.81 12.6 12.34 11.82 12.51 12.42 12.25 12.22 11.61
11.46 12.52 11.76 11.41 11.03 12.77 11.45 11.56 11.87 12.07 12.43 11.79
12.04 12.86 12.88 12.81 12.53 12.84 13.36 13.52 13.62 12.87 13.32 13.08
12.79 13.23 12.58 13.17 13.84 12.45 14.34 12.36 13.69 12.96 13.78 13.45
12.82 13.4 12.2 14.16 13.27 14.13]

```

```

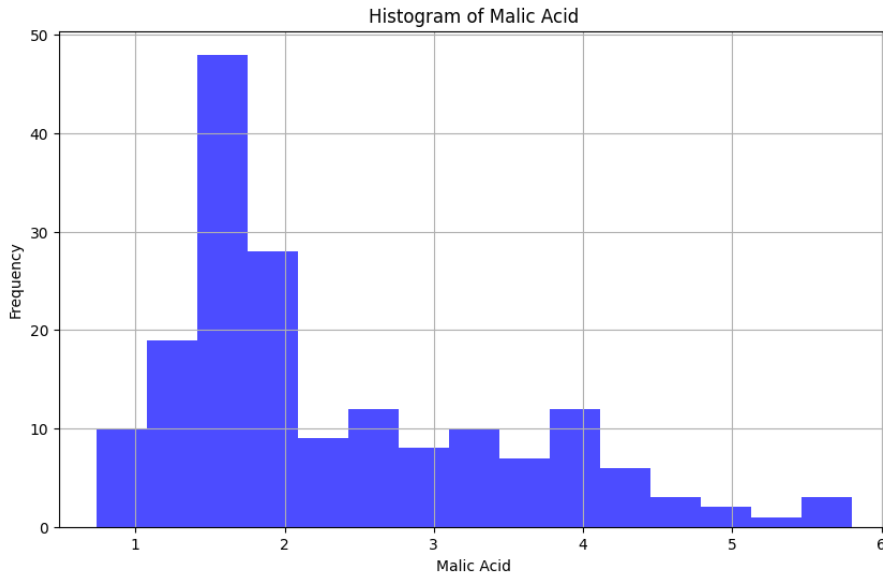
# 7. Replace an extreme set of values in the dataset with NaN (using max value in 'alcohol' as example)
df.loc[df['alcohol'] == df['alcohol'].max(), 'alcohol'] = np.NaN

```

```

# 8. Plot a histogram of a numerical variable (using 'malic_acid')
plt.figure(figsize=(10, 6))
plt.hist(df['malic_acid'], bins=15, color='blue', alpha=0.7)
plt.title('Histogram of Malic Acid')
plt.xlabel('Malic Acid')
plt.ylabel('Frequency')
plt.grid(True)
plt.show()

```



```

# 9. Represent 'target' as a numerical field
tail_with_new_column = df.tail()
print(tail_with_new_column)

```

	index	alcohol	malic_acid	ash	alcalinity_of_ash	magnesium	\
	173	13.71	5.65	2.45	20.5	95.0	
	174	13.40	3.91	2.48	23.0	102.0	
	175	13.27	4.28	2.26	20.0	120.0	
	176	13.17	2.59	2.37	20.0	120.0	
	177	14.13	4.10	2.74	24.5	96.0	

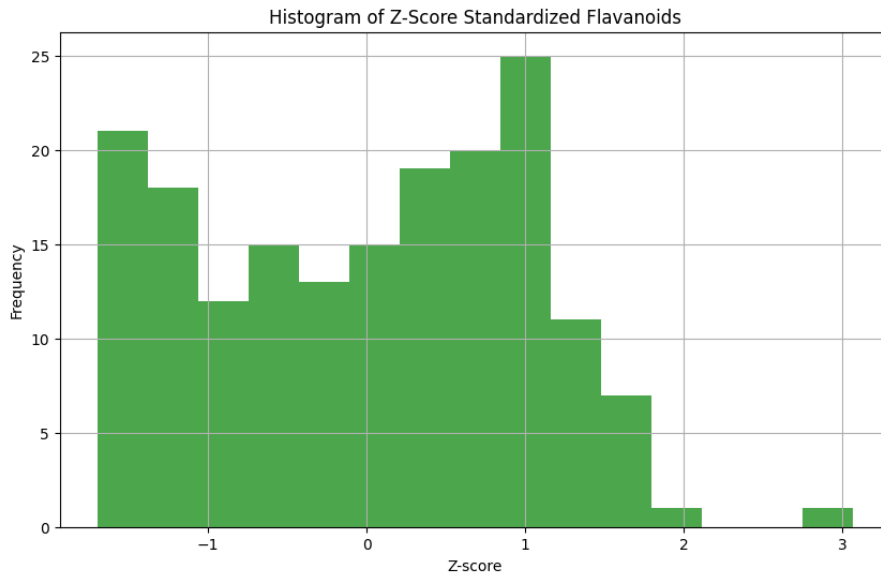
	total_phenols	flavanoids	nonflavanoid_phenols	proanthocyanins	\
173	1.68	0.61	0.52	1.06	
174	1.80	0.75	0.43	1.41	
175	1.59	0.69	0.43	1.35	
176	1.65	0.68	0.53	1.46	
177	2.05	0.76	0.56	1.35	

	color_intensity	hue	od280/od315_of_diluted_wines	proline	target	\
173	7.7	0.64	1.74	740.0	2	
174	7.3	0.70	1.56	750.0	2	
175	10.2	0.59	1.56	835.0	2	
176	9.3	0.60	1.62	840.0	2	
177	9.2	0.61	1.60	560.0	2	

	zscore_flavanoids
173	-1.424900
174	-1.284344
175	-1.344582

```
176 -1.354622
177 -1.274305
```

```
# 10. Standardize a numerical field ('flavanoids') and show its histogram
df['zscore_flavanoids'] = zscore(df['flavanoids'])
plt.figure(figsize=(10, 6))
plt.hist(df['zscore_flavanoids'], bins=15, color='green', alpha=0.7)
plt.title('Histogram of Z-Score Standardized Flavanoids')
plt.xlabel('Z-score')
plt.ylabel('Frequency')
plt.grid(True)
plt.show()
```



```
# 11. Identify outliers (using z-score > 3 or < -3 as criterion for 'flavanoids')
outliers = df[np.abs(zscore(df['flavanoids'])) > 3]
head_outliers = outliers.head()
print(head_outliers)
```

```
   index  alcohol  malic_acid  ash  alkalinity_of_ash  magnesium \
121    121    11.56      2.05  3.23             28.5      119.0

   total_phenols  flavanoids  nonflavanoid_phenols  proanthocyanins \
121           3.18      5.08              0.47             1.87

   color_intensity  hue  od280/od315_of_diluted_wines  proline  target \
121             6.0  0.93              3.69      465.0         1

   zscore_flavanoids
121           3.062832
```

```
# 12. Sort the dataset by 'alcohol' and display 15 interesting fields
df_sorted = df.sort_values(by='alcohol')
interesting_fields = df_sorted.iloc[:, :15].head(15)
print(interesting_fields)
```

```
   index  alcohol  malic_acid  ash  alkalinity_of_ash  magnesium \
115    115    11.03      1.51  2.20             21.5      85.0
113    113    11.41      0.74  2.50             21.0      88.0
120    120    11.45      2.40  2.42             20.0      96.0
110    110    11.46      3.74  1.82             19.5     107.0
121    121    11.56      2.05  3.23             28.5     119.0
109    109    11.61      1.35  2.70             20.0      94.0
 94     94    11.62      1.99  2.28             18.0      98.0
 88     88    11.64      2.06  2.46             21.6      84.0
 87     87    11.65      1.67  2.62             26.0      88.0
 75     75    11.66      1.88  1.92             16.0      97.0
112    112    11.76      2.68  2.92             20.0     103.0
127    127    11.79      2.13  2.78             28.5      92.0
```

96	96	11.81	2.12	2.74	21.5	134.0
103	103	11.82	1.72	1.88	19.5	86.0
116	116	11.82	1.47	1.99	20.8	86.0
	total_phenols	flavanoids	nonflavanoid_phenols	proanthocyanins	\	
115	2.46	2.17	0.52	2.01		
113	2.48	2.01	0.42	1.44		
120	2.90	2.79	0.32	1.83		
110	3.18	2.58	0.24	3.58		
121	3.18	5.08	0.47	1.87		
109	2.74	2.92	0.29	2.49		
94	3.02	2.26	0.17	1.35		
88	1.95	1.69	0.48	1.35		
87	1.92	1.61	0.40	1.34		
75	1.61	1.57	0.34	1.15		
112	1.75	2.03	0.60	1.05		
127	2.13	2.24	0.58	1.76		
96	1.60	0.99	0.14	1.56		
103	2.50	1.64	0.37	1.42		
116	1.98	1.60	0.30	1.53		
	color_intensity	hue	od280/od315_of_diluted_wines	proline	target	
115	1.90	1.71	2.87	407.0	1	
113	3.08	1.10	2.31	434.0	1	
120	3.25	0.80	3.39	625.0	1	
110	2.90	0.75	2.81	562.0	1	
121	6.00	0.93	3.69	465.0	1	
109	2.65	0.96	3.26	680.0	1	
94	3.25	1.16	2.96	345.0	1	
88	2.80	1.00	2.75	680.0	1	
87	2.60	1.36	3.21	562.0	1	
75	3.80	1.23	2.14	428.0	1	
112	3.80	1.23	2.50	607.0	1	
127	3.00	0.97	2.44	466.0	1	
96	2.50	0.95	2.26	625.0	1	
103	2.06	0.94	2.44	415.0	1	
116	1.95	0.95	3.33	495.0	1	