

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

```
In [2]: # 1. Generate synthetic wine data using Gaussian distributions
np.random.seed(42)
num_samples = 1000

# Synthetic features based on Gaussian assumptions
data = {
    "fixed_acidity": np.random.normal(7.0, 0.7, num_samples),
    "volatile_acidity": np.random.normal(0.5, 0.1, num_samples),
    "citric_acid": np.random.normal(0.3, 0.1, num_samples),
    "residual_sugar": np.random.normal(6.0, 1.5, num_samples),
    "chlorides": np.random.normal(0.05, 0.01, num_samples),
    "alcohol": np.random.normal(10.0, 1.0, num_samples)
}

# Generate wine quality based on a combination of other features
# Add randomness to simulate real-world data
wine_quality = (
    0.3 * data["alcohol"] -
    1.5 * data["volatile_acidity"] +
    0.8 * data["citric_acid"] +
    np.random.normal(0, 0.5, num_samples)
).round().astype(int)
```

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In [3]: # Clamp wine quality between 3 and 8
wine_quality = np.clip(wine_quality, 3, 8)

# Add wine_quality to the dataset
data["wine_quality"] = wine_quality
```

```
In [4]: # Convert to DataFrame
df = pd.DataFrame(data)

# 2. Fit Gaussian models and 3. Compute statistical measures
print("Statistical Summary:")
print(df.describe())
```

Statistical Summary:

	fixed_acidity	volatile_acidity	citric_acid	residual_sugar	\
count	1000.000000	1000.000000	1000.000000	1000.000000	
mean	7.013532	0.507084	0.300583	5.971921	
std	0.685451	0.099745	0.098345	1.540699	
min	4.731113	0.205961	-0.001951	1.605827	
25%	6.546687	0.439376	0.235200	4.893869	
50%	7.017710	0.506308	0.299975	6.000277	
75%	7.453561	0.572888	0.366092	7.000418	
max	9.696912	0.819311	0.692624	10.864639	

	chlorides	alcohol	wine_quality
count	1000.000000	1000.000000	1000.000000
mean	0.049507	9.953262	3.04600
std	0.009924	1.007389	0.20959
min	0.018233	7.100486	3.00000
25%	0.043174	9.306307	3.00000
50%	0.049818	9.957173	3.00000
75%	0.056391	10.612447	3.00000
max	0.081129	13.098299	4.00000

In [9]: *#Visualizations - Histogram & PDF*

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features = ["alcohol", "volatile_acidity", "citric_acid", "wine_quality"]
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for feature in features:
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    mean = np.mean(df[feature])
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    std = np.std(df[feature])
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    x = np.linspace(df[feature].min(), df[feature].max(), 100)
```

```
    pdf = norm.pdf(x, mean, std)
```

```
plt.figure(figsize=(8, 5))
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sns.histplot(df[feature], bins=30, kde=False, color='skyblue', stat="density")
```

```
plt.plot(x, pdf, 'r--', label=f'Gaussian PDF\(\mu={mean:.2f}, \sigma={std:.2f}\)')
```

```
plt.title(f'Distribution for {feature}')
```

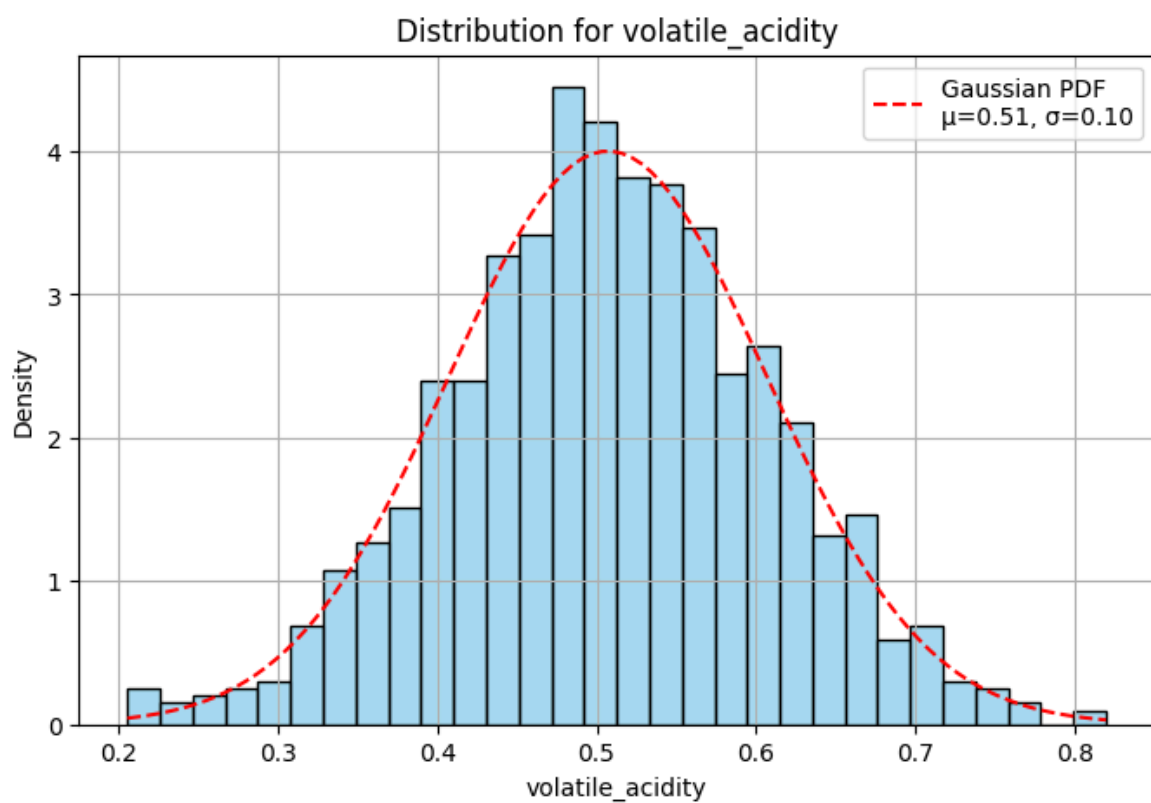
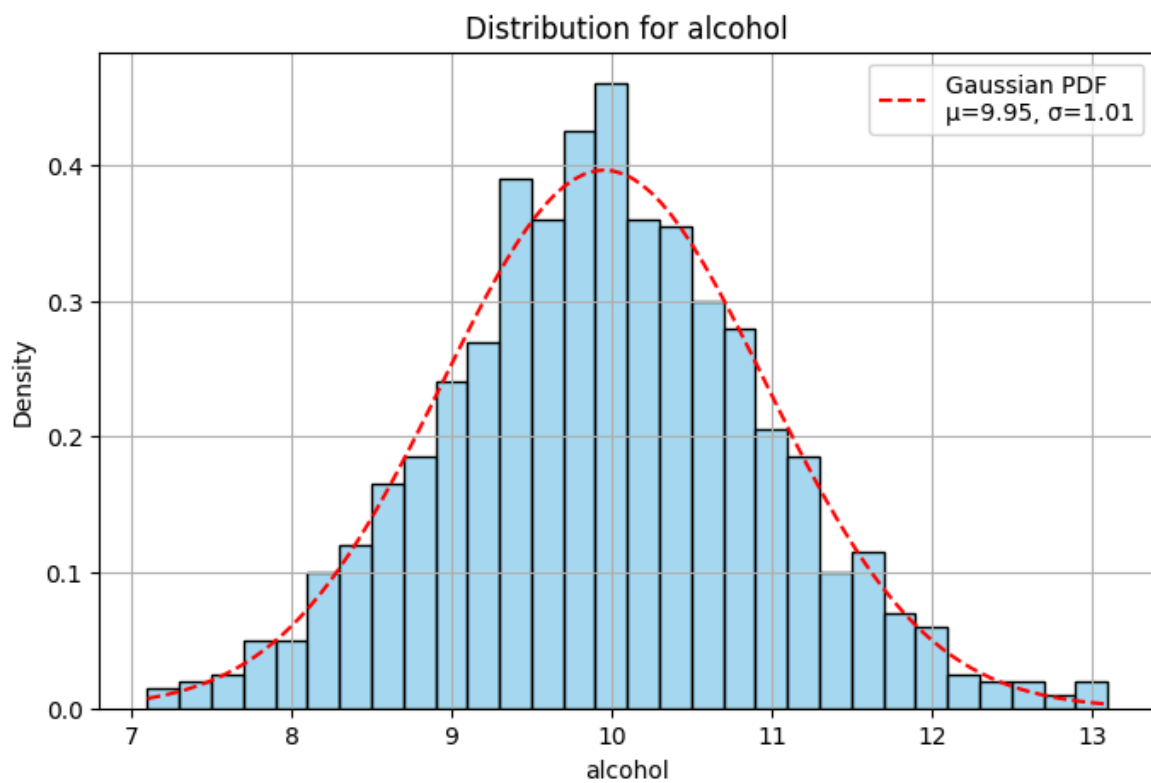
```
plt.xlabel(feature)
```

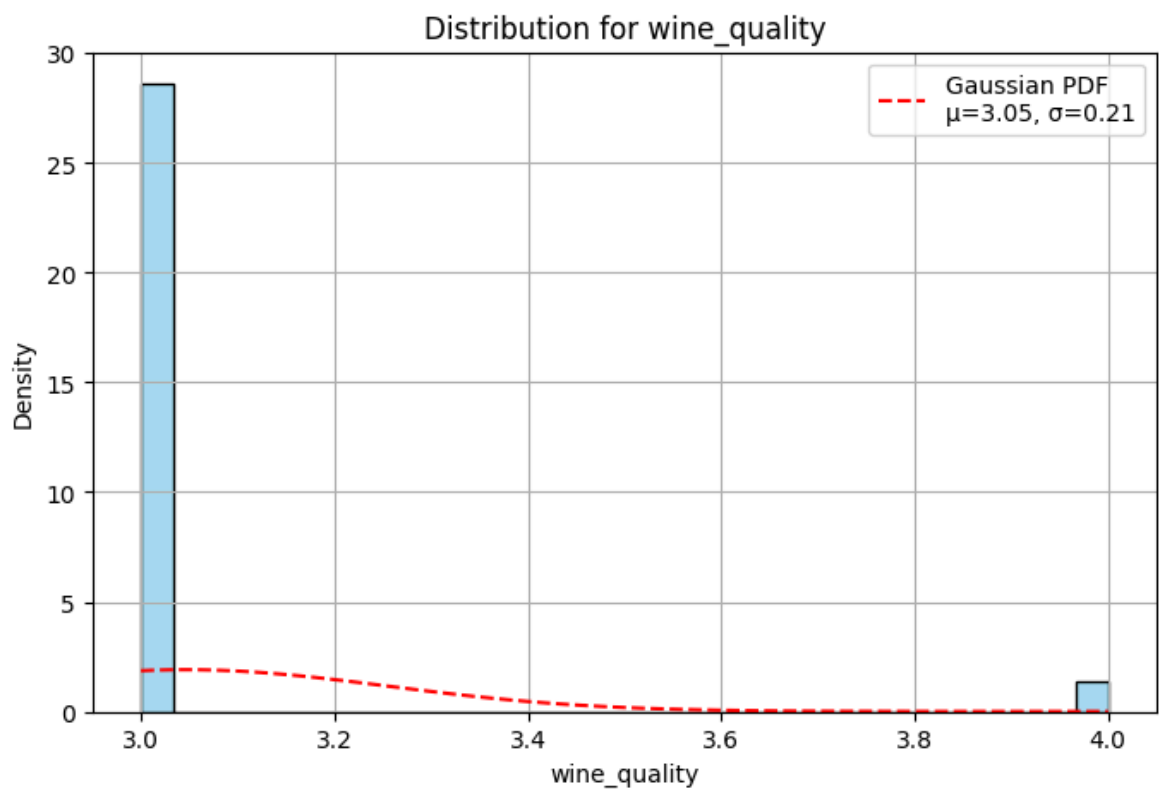
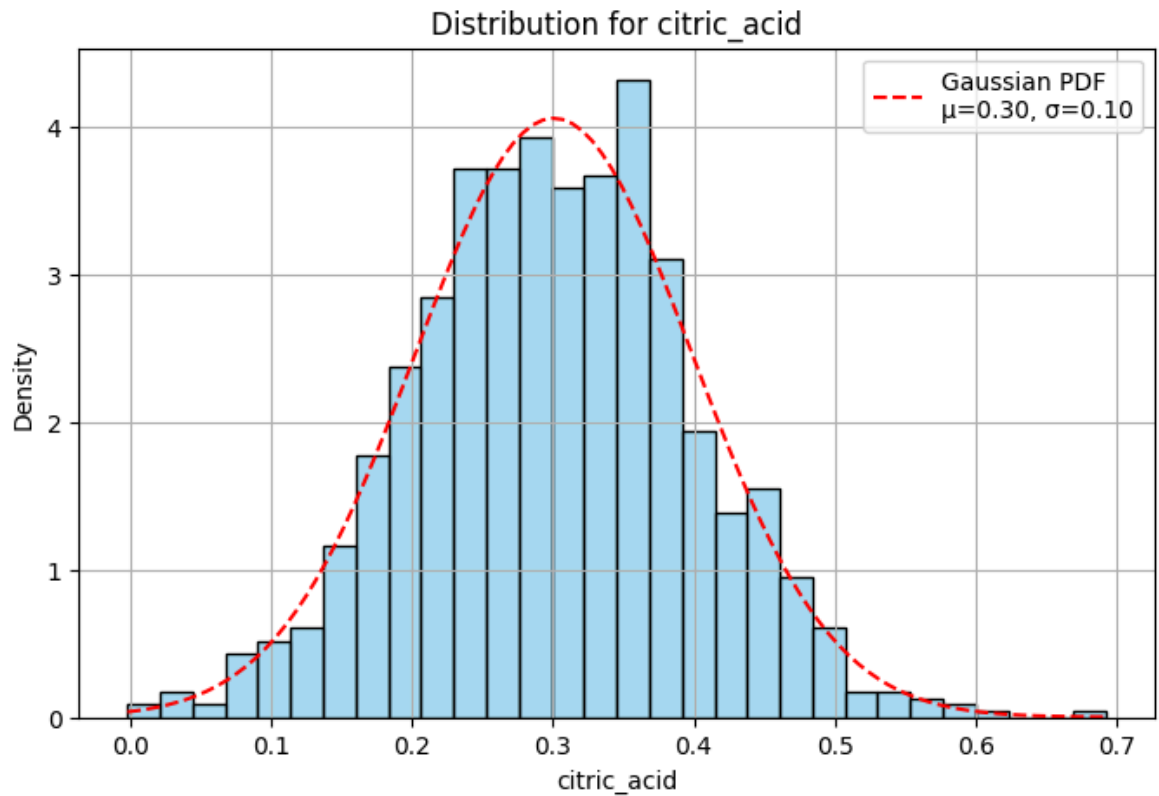
```
plt.ylabel("Density")
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plt.legend()
```

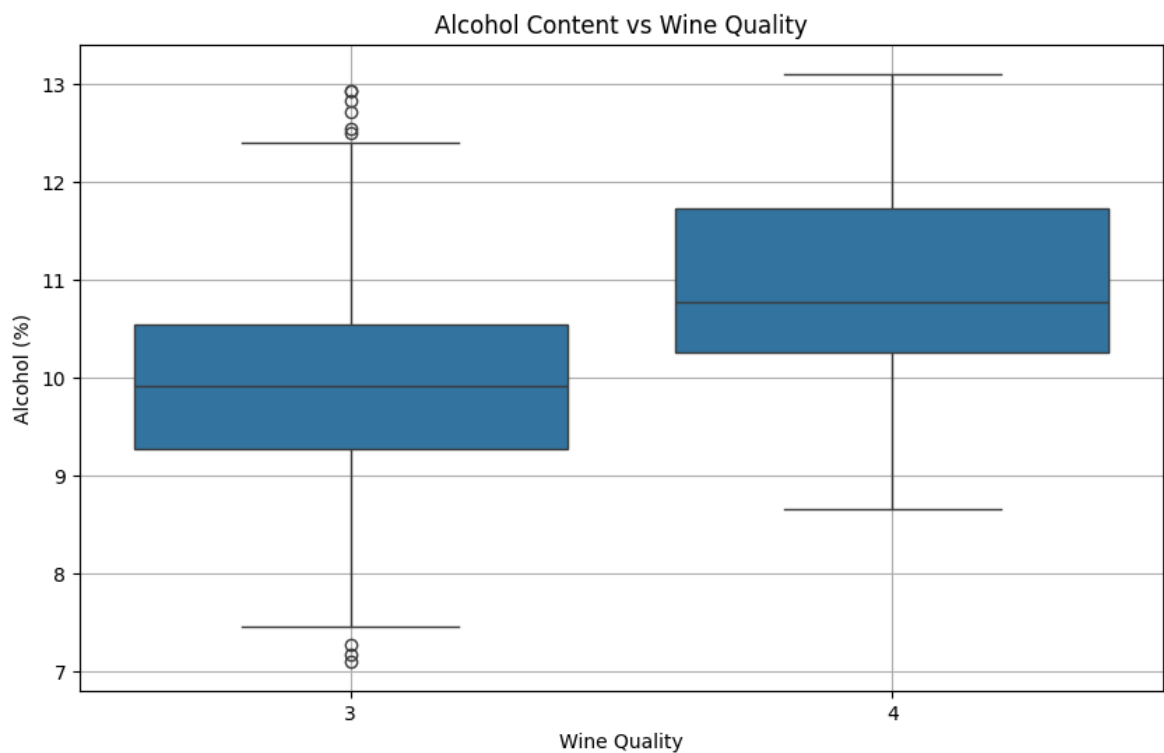
```
plt.grid(True)
```

```
plt.show()
```

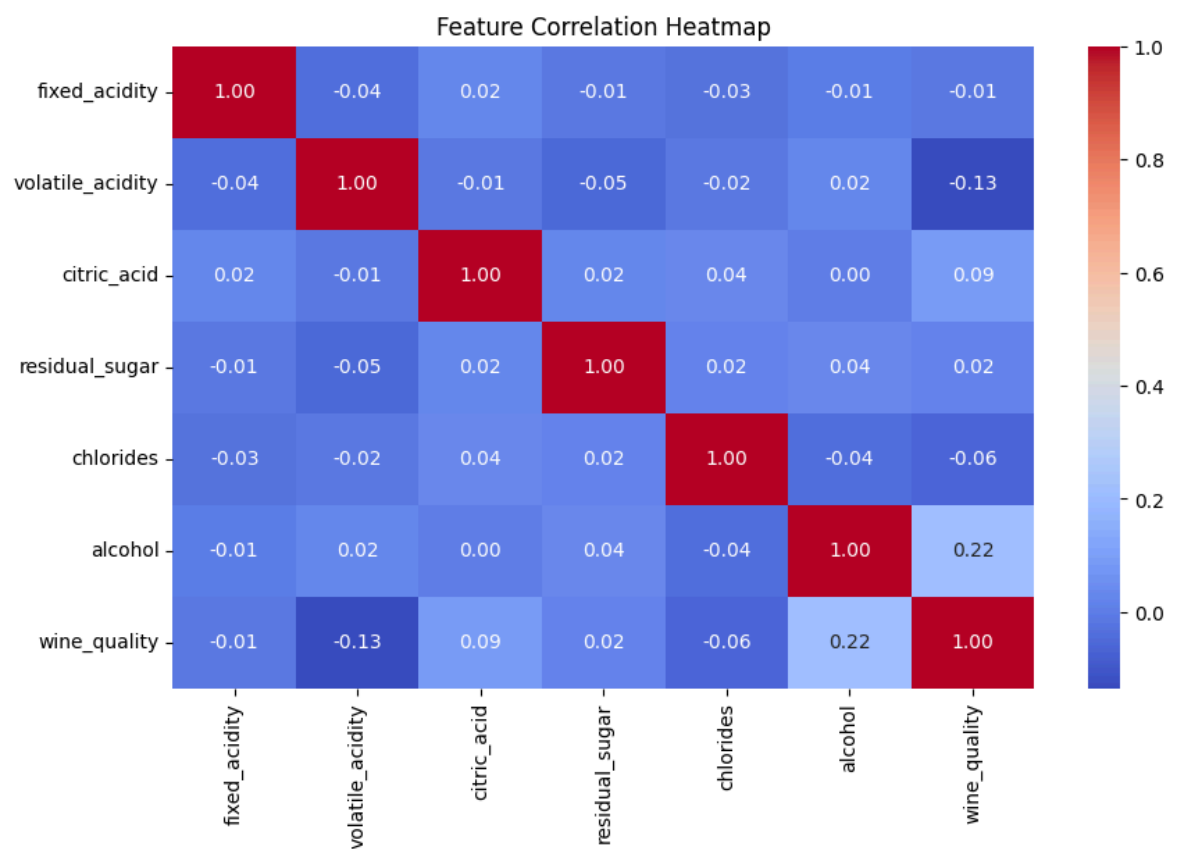




```
In [7]: # Analyze feature vs wine_quality
plt.figure(figsize=(10, 6))
sns.boxplot(x="wine_quality", y="alcohol", data=df)
plt.title("Alcohol Content vs Wine Quality")
plt.xlabel("Wine Quality")
plt.ylabel("Alcohol (%)")
plt.grid(True)
plt.show()
```



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In [8]: # Correlation heatmap
plt.figure(figsize=(10, 6))
sns.heatmap(df.corr(), annot=True, cmap="coolwarm", fmt=".2f")
plt.title("Feature Correlation Heatmap")
plt.show()
```



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In [ ]:
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