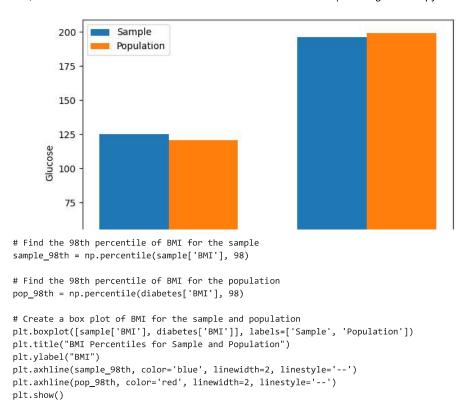
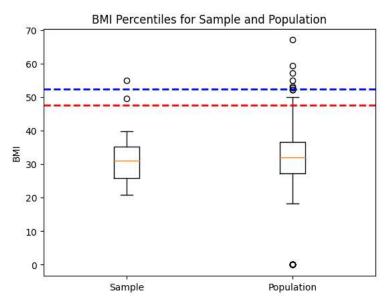
diabetes.head()

ightharpoons	ı	Pregnancies	Glucose	BloodPressure	SkinThickness	Insulin	BMI	DiabetesPedigreeFunction	Age	Outcome	1
	0	6	148	72	35	0	33.6	0.627	50	1	
	1	1	85	66	29	0	26.6	0.351	31	0	
	2	8	183	64	0	0	23.3	0.672	32	1	
	3	1	89	66	23	94	28.1	0.167	21	0	
	4	0	137	40	35	168	43.1	2.288	33	1	

```
# Set random seed
np.random.seed(99)
# Generate a random sample of 25 observations
sample = diabetes.sample(n=25)
# Calculate mean and max of sample
mean_sample = np.mean(sample['Glucose'])
max_sample = np.max(sample['Glucose'])
# Calculate mean and max of population
mean_pop = diabetes['Glucose'].mean()
max_pop = diabetes['Glucose'].max()
# Create a bar chart to compare mean and max values
labels = ['Mean Glucose', 'Max Glucose']
sample_values = [mean_sample, max_sample]
pop_values = [mean_pop, max_pop]
x = np.arange(len(labels))
width = 0.35
fig, ax = plt.subplots()
rects1 = ax.bar(x - width/2, sample_values, width, label='Sample')
rects2 = ax.bar(x + width/2, pop_values, width, label='Population')
ax.set_ylabel('Glucose')
ax.set_xticks(x)
ax.set_xticklabels(labels)
ax.legend()
fig.tight_layout()
plt.show()
```





```
# Set random seed
np.random.seed(123)
# Define function to calculate mean, standard deviation, and percentile of BloodPressure
def calc_stats(data):
    return np.mean(data), np.std(data), np.percentile(data, 98)
# Generate 500 bootstrap samples of 150 observations each
n_samples = 500
sample_size = 150
bootstrap_means = []
bootstrap_stds = []
bootstrap_percentiles = []
for i in range(n_samples):
    sample = np.random.choice(diabetes['BloodPressure'], size=sample_size, replace=True)
    mean, std, percentile = calc_stats(sample)
   bootstrap_means.append(mean)
   bootstrap_stds.append(std)
   bootstrap_percentiles.append(percentile)
```

```
# Calculate population statistics for BloodPressure
pop_mean = np.mean(diabetes['BloodPressure'])
pop_std = np.std(diabetes['BloodPressure'])
pop_percentile = np.percentile(diabetes['BloodPressure'], 98)
# Create histograms to compare bootstrap means, standard deviations, and percentiles to population statistics
fig, axs = plt.subplots(3, 1, figsize=(8, 12))
axs[0].hist(bootstrap_means, bins=30)
axs[0].axvline(pop_mean, color='r', linestyle='--')
axs[0].set_xlabel('BloodPressure Mean')
axs[0].set_ylabel('Frequency')
axs[1].hist(bootstrap_stds, bins=30)
axs[1].axvline(pop_std, color='r', linestyle='--')
axs[1].set_xlabel('BloodPressure Standard Deviation')
axs[1].set_ylabel('Frequency')
axs[2].hist(bootstrap_percentiles, bins=30)
axs[2].axvline(pop_percentile, color='r', linestyle='--')
axs[2].set_xlabel('BloodPressure 98th Percentile')
axs[2].set_ylabel('Frequency')
plt.tight_layout()
plt.show()
```

