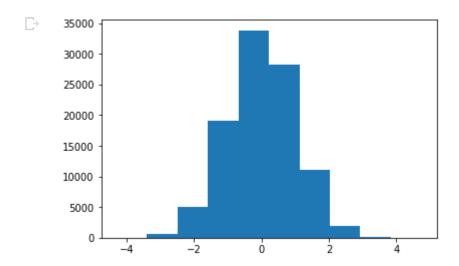
```
import numpy as np
import matplotlib.pyplot as plt
import seaborn as sns
import math
```

Generating Random sample data in Normal Distribustion form

```
sample_data = np.random.normal(size=100000)
plt.hist(sample_data)
plt.show() #to plot histogram without axis
```



sns.distplot(sample_data)
#sns.displot(sample data)

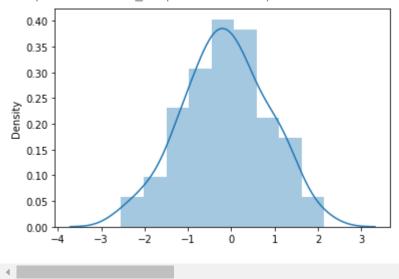
/usr/local/lib/python3.7/dist-packages/seaborn/distributions.py:2619: FutureWarning: `di warnings.warn(msg, FutureWarning)
<matplotlib.axes._subplots.AxesSubplot at 0x7f3a8e68a450>

Normal Dist sample data

sample_data = np.random.normal(size=100)
sns.distplot(sample_data)

/usr/local/lib/python3.7/dist-packages/seaborn/distributions.py:2619: FutureWarnin warnings.warn(msg, FutureWarning)

<matplotlib.axes. subplots.AxesSubplot at 0x7f3a844e6d50>

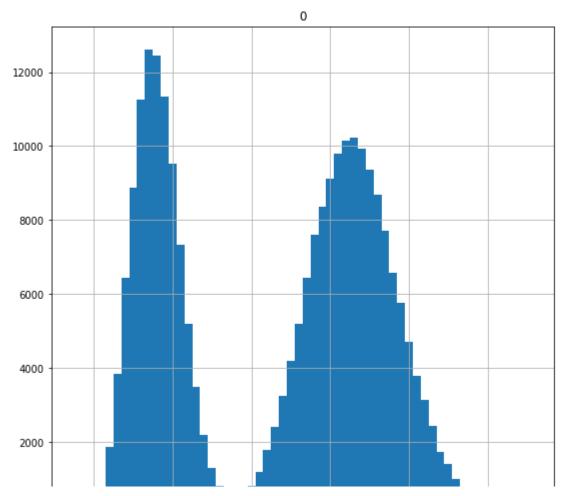


Uniform Dist sample data

sample_data= np.random.uniform(size=10000)
sns.distplot(sample data)

```
/usr/local/lib/python3.7/dist-packages/seaborn/distributions.py:2619: FutureWarnin
       warnings.warn(msg, FutureWarning)
     <matplotlib.axes. subplots.AxesSubplot at 0x7f3a844e3a10>
Point Estimates
from scipy.stats import poisson
      일 0.6 1
np.random.seed(10)
population ages1 =poisson.rvs(loc=18, mu=35, size=150000)
population ages2 =poisson.rvs(loc=18, mu=10, size=100000)
population ages = np.concatenate((population ages1, population ages2)) # concat
population ages.mean() # what is the true mean age of the population of 250000 people?
     43.002372
np.random.seed(6)
sample ages = np.random.choice(a=population ages, size=500) # Sample 500 values
sample_ages.mean() # Show sample mean
                                          point estimate
     42.388
population ages.mean() - sample ages.mean()
     0.6143720000000003
import random
  import pandas as pd
  import scipy.stats as stats
Studying Sampling distribution
  pd.DataFrame(population ages).hist(bins=58,range=(17.5,75.5),figsize=(9,9))
  print( stats.skew(population ages) )
```

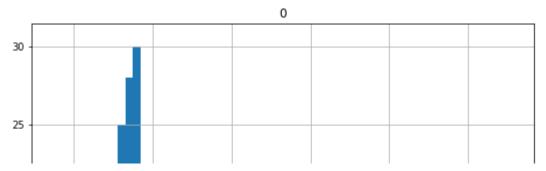
-0.12008483603917186



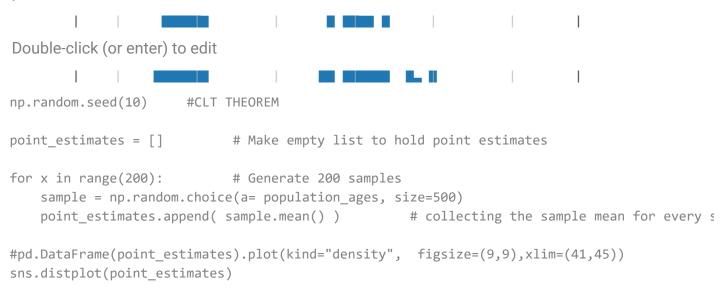
pd.DataFrame(sample_ages).hist(bins=58,range=(17.5,75.5),figsize=(9,9))

print(stats.skew(sample_ages))

-0.056225282585406065

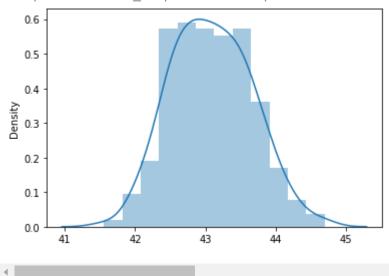


create a sampling distribution by taking 200 samples from our population and then making 200 point estimates of the mean:



/usr/local/lib/python3.7/dist-packages/seaborn/distributions.py:2619: FutureWarning: `di warnings.warn(msg, FutureWarning)

<matplotlib.axes. subplots.AxesSubplot at 0x7f3a83cf9110>



#The sampling distribution appears to be roughly normal, despite the bimodal population dist #In addition, the mean of the sampling distribution approaches the true population mean

Confidence Intervals

```
#S.D is known
sample_size = 1000
sample = np.random.choice(a= population_ages, size = sample_size) #draw some random sample
sample mean = sample.mean() #point estimate
print(sample mean)
z_{critical} = stats.norm.ppf(q = 0.975) # Get the z-critical value* # area under curve
print("z-critical value:")
                                # Check the z-critical value
print(z critical)
pop stdev = population ages.std() # Get the population standard deviation
margin_of_error = z_critical * (pop_stdev/math.sqrt(sample_size)) #MG= Z*SIGMA/SQ(N)
confidence interval = (sample mean - margin of error, sample mean + margin of error)
                                                                                        #X-
print("Confidence interval:")
print(confidence interval)
    43.266
    z-critical value:
    1.959963984540054
    Confidence interval:
    (42.44606406882683, 44.08593593117317)
sample_mean
    43.282
stats.norm.interval(alpha=0.95,
                                 # Confidence level
                   loc=sample mean,
                   scale=pop_stdev/math.sqrt(sample_size)) # Scaling factor
     (42.46206406882683, 44.101935931173166)
```

T-distribution

```
#SD is unknown
sample_size = 25
sample = np.random.choice(a= population ages, size = sample size) #
sample mean = sample.mean()
t critical = stats.t.ppf(q = 0.975, df=24) # Get the t-critical value* df=n-1
print("t-critical value:")
                                          # Check the t-critical value
print(t critical)
sample stdev = sample.std() # Get the sample standard deviation
sigma = sample_stdev/math.sqrt(sample_size) # Standard deviation estimate MG= t*SIGMA/S
margin of error = t critical * sigma
confidence interval = (sample mean - margin of error, sample mean + margin of error)
print("Confidence interval:")
print(confidence interval)
    t-critical value:
    2.0638985616280205
    Confidence interval:
     (37.659096531075285, 47.46090346892472)
# Check the difference between critical values with a sample size of 1000
stats.t.ppf(q=0.975, df=999) - stats.norm.ppf(0.975)
    0.0023774765933946007
                                          # Confidence level
stats.t.interval(alpha = 0.95,
                df= 24,
                                           # Degrees of freedom
                loc = sample_mean,
                                          # Sample mean
                scale = sigma)
                                           # Standard deviation estimate
     (37.659096531075285, 47.46090346892472)
```

Confidence interval using Proportions