## Problem 1

Remember from last week we discussed that skewness and kurtosis functions in statistical packages are often biased. Is your function biased? Prove or disprove your hypothesis.

### Solution:

The function used are scipy.stats.skew() and scipy.stats.kurtosis(). By definition, these two functions are default biased. As both of the functions have a parameter bias.

#### Notes

The sample skewness is computed as the Fisher-Pearson coefficient of skewness, i.e.

$$g_1 = rac{m_3}{m_2^{3/2}}$$

where

$$m_i = rac{1}{N} \sum_{n=1}^N (x[n] - ar{x})^i$$

is the biased sample ith central moment, and  $\bar{x}$  is the sample mean. If bias is False, the calculations are corrected for bias and the value computed is the adjusted Fisher-Pearson standardized moment coefficient, i.e.

$$G_1 = rac{k_3}{k_2^{3/2}} = rac{\sqrt{N(N-1)}}{N-2} rac{m_3}{m_2^{3/2}}.$$

We solve this problem by doing the hypothesis test.

### Steps

- 1. Sample 10, 100, 1,000, 10,000 standardized random normal values.
- 2. Calculate the skewness and kurtosis by python function skew() and kurtosis().
- 3. Sample the skewness and kurtosis by repeating steps 1 and 2 100 times.
- 4. Calculate the p-value ( $\mu$ 0 = 0).
- 5. If the value is lower than your threshold (typically 5%), then you reject the hypothesis that the kurtosis function is unbiased.

We have the following result:

```
10 samples:
skewness p-value is 0.010022125067440038.
Reject the null hypothesis.

kurtosis p-value is 9.892089260419227e-12.
Reject the null hypothesis.

100 samples:
skewness p-value is 0.835437664512038.
Fail to reject the null hypothesis.

kurtosis p-value is 0.6467761933137416.
Fail to reject the null hypothesis.

1000 samples:
skewness p-value is 0.0375664513602751.
Reject the null hypothesis.

kurtosis p-value is 0.9977733631042587.
Fail to reject the null hypothesis.

10000 samples:
skewness p-value is 0.472536613094932.
Fail to reject the null hypothesis.

kurtosis p-value is 0.472536613094932.
Fail to reject the null hypothesis.
```

These two functions are **biased** as we rejected the null hypothesis at significant level of 0.05 under 10 sample size tested.

As sample size going larger and larger, the functions tend to be unbiased.

## Problem 2

2.1 Fit the data in problem2.csv using OLS and calculate the error vector. Look at its distribution. How well does it fit the assumption of normally distributed errors?

### Solution:

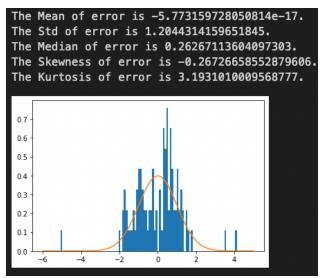
Using OLS regression by using the python sm.OLS() function. The OLS Regression results is shown by ols model.summary() as

	OLS Regression	n Results			
Dep. Variable:			0.195		
Model:			0.186		
Method:	Least Squares		23.68		
Date:			4.34e-06		
Time:					
No. Observations:	100	AIC:	324.0		
Df Residuals:	98	BIC:	329.2		
Df Model:	1				
Covariance Type: nonrobust					
coef st	derr t P	> t  [0.025 0.975]			
ones 0.1198 (	0.121 0.990 0.3	25 -0.120 0.360			
x 0.6052 0	0.124 4.867 0.0	00 0.358 0.852			
Omnibus:	14.146 Durbin-	-Watson: 1.885			
Prob(Omnibus):	0.001 Jarque-B	era (JB): 43.673			
Skew:	-0.267 F	Prob(JB): 3.28e-10			
Kurtosis:	6.193 C	ond. No. 1.03			

The error vector is calculated by ols model.resid. Part of the error vector is shown as following:

0	-0.838485
1	0.835296
2	1.027428
3	1.319711
4	-0.152317
95	-1.590264
95 96	-1.590264 -1.694848
96	-1.694848

The 4 moments of the distribution is calculated. The histogram of distribution is also plotted.



The Skewness of error is -0.26726658552879606. The Kurtosis of error is 3.1931010009568777. The Std of error is 1.2044314159651845. These three moments is far from that of the normal distribution. The histogram of it is not like normal distribution as well. It does not fit the assumption of normally distributed errors.

## 2.2 & 2.3

Fit the data using MLE given the assumption of normality. Then fit the MLE using the assumption of a T distribution of the errors. Which is the best fit?

What are the fitted parameters of each and how do they compare? What does this tell us about the breaking of the normality assumption in regards to expected values in this case?

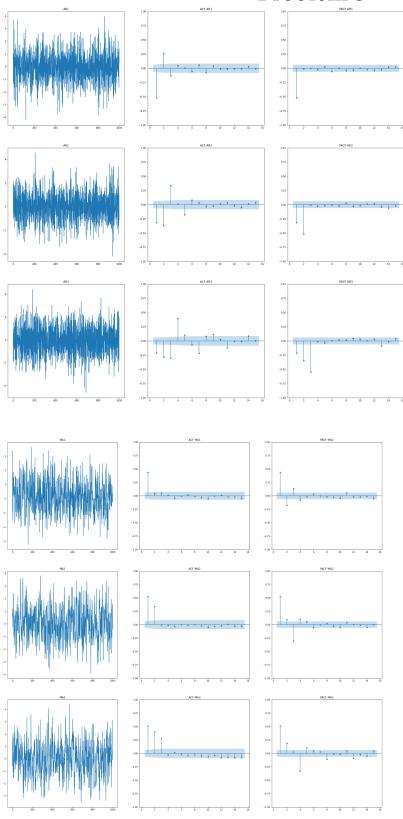
The Goodness of fitness statistic R Square and Information criteria AIC and BIC is used in model selection.

	OLS	MLE_Norm	MLE_T
Const	0.1198	0.1198	0.1426
X	0.6052	0.6052	0.5576
R_Squared	0.195	0.1946	0.1934
AIC	324.0	324.0	315.0
BIC	329.2	329.2	320.2

The OLS and the MLE with normal distribution led to the same result. Then MLE\_Norm and MLE\_T are compared. The first model selection statistic R\_Squared is quite similar. But the MLE T has smaller AIC and BIC. The MLE t is the best fit.

Both OLS and MLE\_Norm based on the assumption of normal distribution. From this result we can see that the normality assumption might be broken. And use the MLE with t distribution might have a better result.

# Problem 3



The simulate process are drawn as labeled in the graph.

	ACF	PACF
AR	Decrease exponentially	Rapid decrease after order
MA	Rapid decrease after order	Decrease exponentially

Decrease refers to absolute value

By the pattern of decrease rapidly we can identify the type and order of each process.