

# hw4

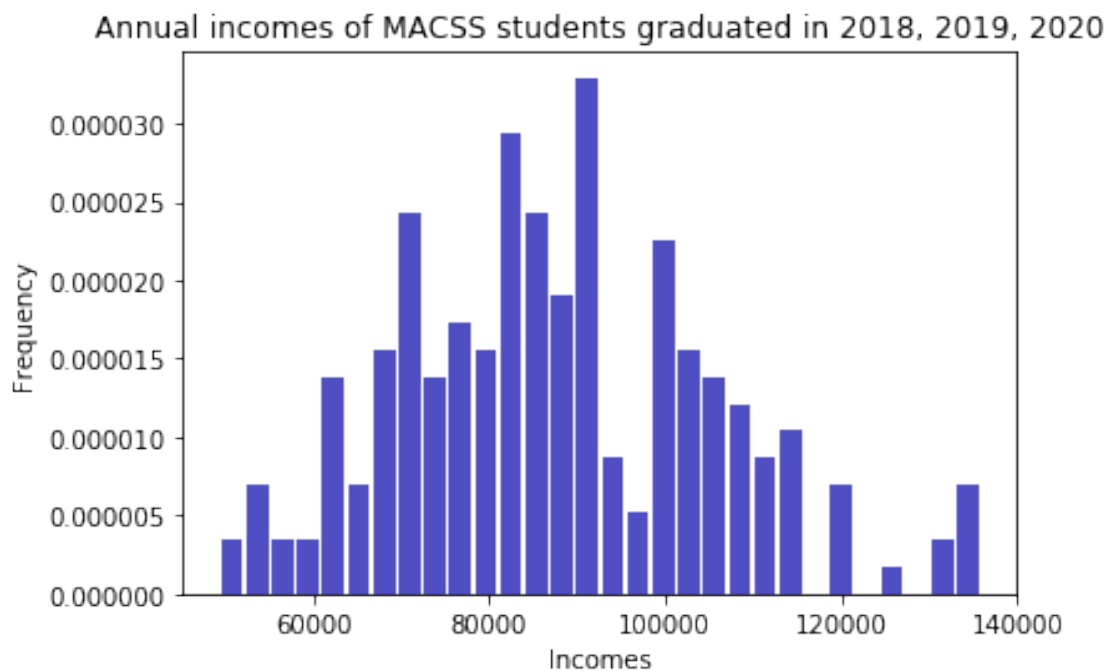
February 10, 2019

```
In [1]: import numpy as np
import pandas as pd
import matplotlib.pyplot as plt
import scipy.stats as sts
from scipy.integrate import quad
```

0.1 1

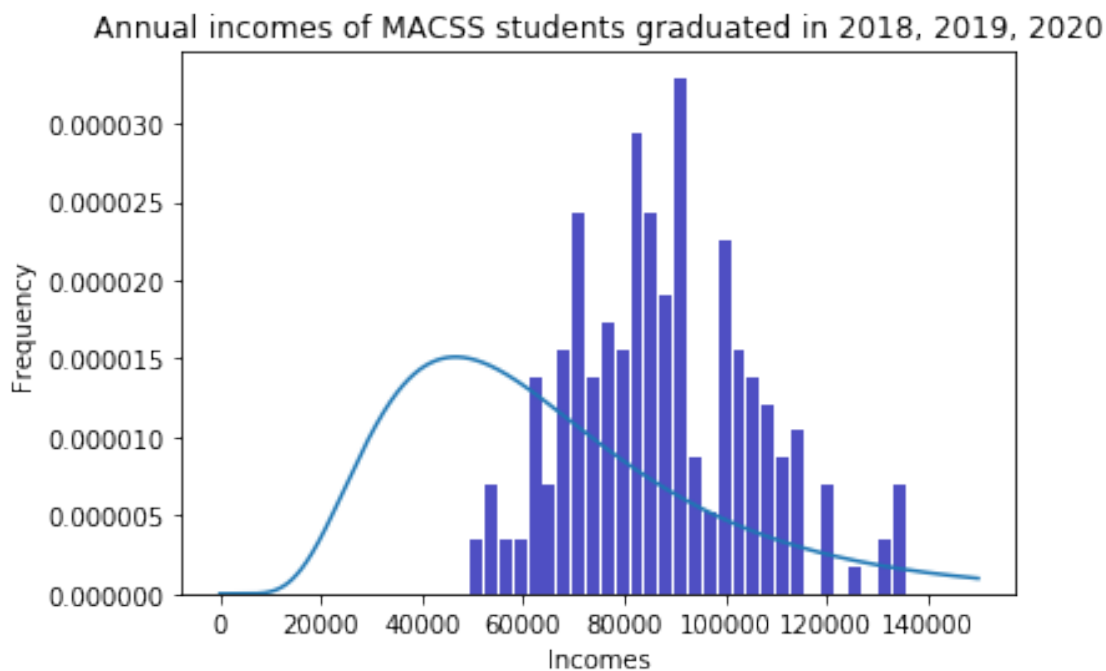
0.1.1 a

```
In [2]: Incomes = np.loadtxt('data/incomes.txt')
fig,ax = plt.subplots()
ax.set_xlabel('Incomes')
ax.set_ylabel('Frequency')
ax.set_title('Annual incomes of MACSS students graduated in 2018, 2019, 2020')
ax.hist(x=Incomes, bins=30, color='#0504aa', alpha=0.7, rwidth=0.85, normed = True)
plt.show()
```



## 0.1.2 b

```
In [3]: def lognorm(x, mu=11, sigma=0.5):
        return sts.lognorm.pdf(x, s = abs(sigma), scale = np.exp(mu))
def log_lik_truncnorm(x, mu=11, sigma=0.5):
    pdf_vals = lognorm(x, mu, sigma)
    polishedpdf = pdf_vals
    ln_pdf_vals = np.log(polishedpdf)
    log_lik_val = ln_pdf_vals.sum()
    return log_lik_val
X = np.linspace(0,150000, 150001)
fig,ax = plt.subplots()
ax.set_xlabel('Incomes')
ax.set_ylabel('Frequency')
ax.set_title('Annual incomes of MACSS students graduated in 2018, 2019, 2020')
ax.plot(X, lognorm(X))
ax.hist(x=Incomes, bins=30, color='#0504aa', alpha=0.7, rwidth=0.85, normed = True)
plt.show()
LLV = log_lik_truncnorm(Incomes)
print('The value of the log likelihood value is', LLV)
```



The value of the log likelihood value is -2385.856997808558

### 0.1.3 c

```
In [4]: def crit(params):
        mu, sigma = params
        log_lik_val = log_lik_truncnorm(Incomes, mu, sigma)
        neg_log_lik_val = -log_lik_val
        return neg_log_lik_val

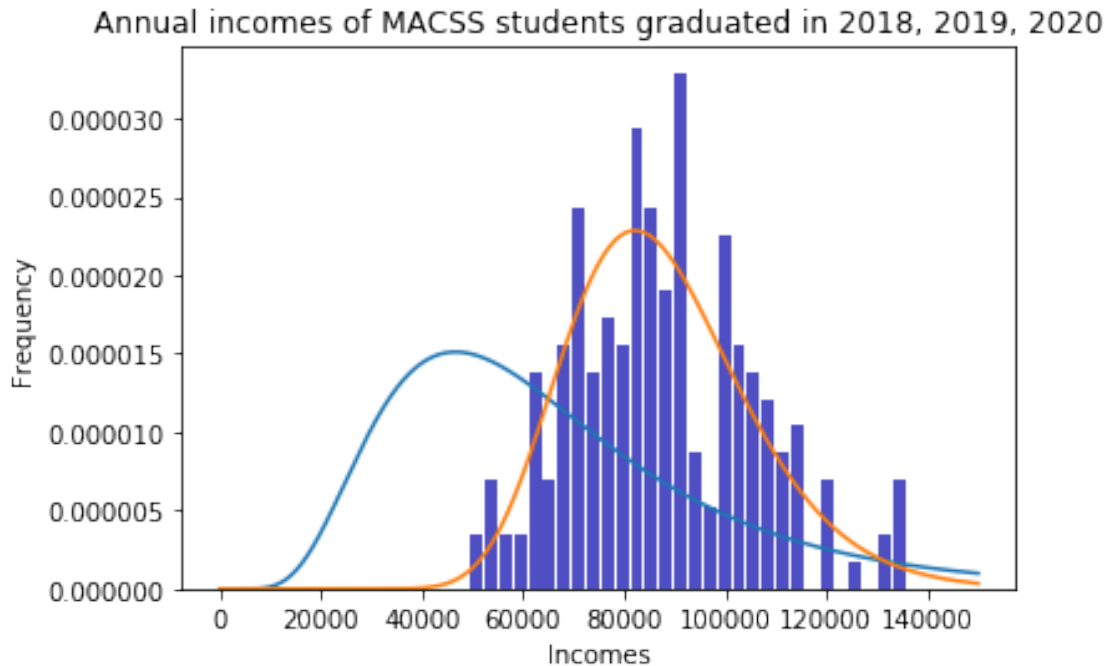
In [5]: import scipy.optimize as opt
        mu_init = 11
        sig_init = 0.5
        params_init = np.array([mu_init, sig_init])

        results = opt.minimize(crit, params_init, tol=1e-14, method='L-BFGS-B')
        mu_MLE, sig_MLE = results.x
        LLV_MLE = log_lik_truncnorm(Incomes, mu_MLE, sig_MLE)
        VCV_MLE = results.hess_inv.todense()

        print('mu_MLE=', mu_MLE, ' sig_MLE=', sig_MLE)
        print('The value of the log likelihood value is', LLV_MLE)
        print('The variance-covariance matrix is', VCV_MLE)

mu_MLE= 11.359022999045031  sig_MLE= 0.20817732039955567
The value of the log likelihood value is -2241.7193013573583
The variance-covariance matrix is [[2.40426054e-04 4.06249275e-06]
 [4.06249275e-06 1.09631836e-04]]

In [6]: X = np.linspace(0,150000, 150001)
        fig,ax = plt.subplots()
        ax.set_xlabel('Incomes')
        ax.set_ylabel('Frequency')
        ax.set_title('Annual incomes of MACSS students graduated in 2018, 2019, 2020')
        ax.plot(X, lognorm(X), label = 'initial guess')
        ax.plot(X, lognorm(X, mu=mu_MLE,sigma=sig_MLE), label = 'MLE result')
        ax.hist(x=Incomes, bins=30, color='#0504aa', alpha=0.7, rwidth=0.85, normed = True)
        plt.show()
```



#### 0.1.4 d

```
In [7]: mu_new, sig_new = np.array([11, 0.5])
log_lik_h0 = log_lik_truncnorm(Incomes, mu_new, sig_new)
print('hypothesis value log likelihood', log_lik_h0)
log_lik_mle = log_lik_truncnorm(Incomes, mu_MLE, sig_MLE)
print('MLE log likelihood', log_lik_mle)
LR_val = 2 * (log_lik_mle - log_lik_h0)
print('likelihood ratio value', LR_val)
pval_h0 = 1.0 - sts.chi2.cdf(LR_val, 2)
print('chi squared of H0 with 2 degrees of freedom p-value = ', pval_h0)
```

hypothesis value log likelihood -2385.856997808558

MLE log likelihood -2241.7193013573583

likelihood ratio value 288.2753929023993

chi squared of H0 with 2 degrees of freedom p-value = 0.0

#### 0.1.5 e

```
In [8]: print("The probability that you will earn more than $100,000 is",
quad(lambda x: lognorm(x, mu=mu_MLE, sigma=sig_MLE), 100000, np.inf)[0])
print("The probability that you will earn less than $75,000 is",
quad(lambda x: lognorm(x, mu=mu_MLE, sigma=sig_MLE), 0, 75000)[0])
```

The probability that you will earn more than \$100,000 is 0.22986683846424502  
The probability that you will earn less than \$75,000 is 0.2602342648237652

## 0.2 2

### 0.2.1 a

```
In [9]: def norm_pdf(x, sigma):
        sigma = np.abs(sigma)
        pdf_vals = (1/(sigma * np.sqrt(2 * np.pi))) *
                    np.exp( - x**2 / (2 * sigma**2)))
        return pdf_vals
def log_lik_norm(x, beta0, beta1, beta2, beta3, sigma):
    error = x[:,0] - beta0 - beta1*x[:,1] - beta2*x[:,2] - beta3*x[:,3]
    pdf_vals = norm_pdf(error, sigma)
    ln_pdf_vals = np.log(pdf_vals)
    log_lik_val = ln_pdf_vals.sum()
    return log_lik_val
def new_crit(params, *args):
    beta0, beta1, beta2, beta3, sigma = params
    x = args[0]
    log_lik_val = log_lik_norm(x,beta0, beta1, beta2, beta3, sigma)
    neg_log_lik_val = -log_lik_val
    return neg_log_lik_val
Data = []
with open('data/sick.txt') as f:
    for line in f:
        line = line.strip()
        data = line.split(',')
        Data.append(data)
Data.pop(0)
for i in range(len(Data)):
    Data[i] = [float(x) for x in Data[i]]
Data = np.array(Data)
new_Data = np.ones_like(Data)
new_Data[:,1:] = Data[:,1:]
beta0_init, beta1_init, beta2_init, beta3_init = np.linalg.lstsq(new_Data,Data[:,0])[0]
sigma_init = 0.01
parameters_init = np.array([beta0_init, beta1_init, beta2_init, beta3_init, sigma_init])
results2 = opt.minimize(new_crit, parameters_init, args=(Data),tol=1e-14, method='L-BFGS-B')
beta0_MLE, beta1_MLE, beta2_MLE, beta3_MLE, sigma_MLE = results2.x
new_LLVMLE = log_lik_norm(Data, beta0_MLE, beta1_MLE, beta2_MLE, beta3_MLE, sigma_MLE)
new_VCMLE = results2.hess_inv.todense()
print('beta0_MLE =',beta0_MLE,
      'beta1_MLE =',beta1_MLE,
      'beta2_MLE =',beta2_MLE,
      'beta3_MLE =',beta3_MLE,
```

```

        'sigma_MLE = ',sigma_MLE)
print('The value of the log likelihood value is', new_LLVMLE)
print('The variance-covariance matrix is', new_VCV_MLE)
Error = Data[:,0] - beta0_MLE - beta1_MLE*Data[:,1] - beta2_MLE*Data[:,2] - beta3_MLE*
X2 = np.linspace(-0.01,0.01,10001)
fig,ax = plt.subplots()
ax.set_xlabel('Errors')
ax.set_ylabel('Frequency')
ax.set_title('Estimate Errors')
ax.plot(X2, norm_pdf(X2,sigma_MLE))
ax.hist(x=Error, bins='auto', color='#0504aa', alpha=0.7, rwidth=0.85, normed = True)
plt.show()

```

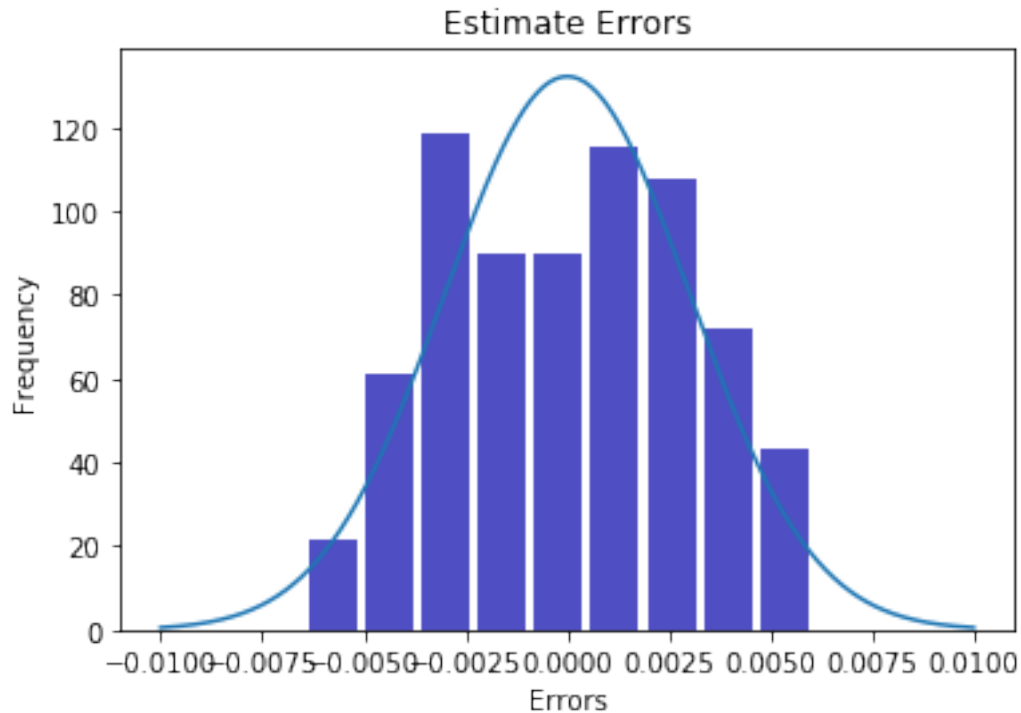
```

beta0_MLE = 0.25164168313073476 beta1_MLE = 0.01293361306625627 beta2_MLE = 0.4004998342959863
The value of the log likelihood value is 876.8650774281919
The variance-covariance matrix is [[ 1.00526159e+00  1.91017126e-03  6.33297010e-03  9.30956564e-03
-7.36909408e-01]
[ 1.91017126e-03  9.45409236e-02 -7.77969811e-02 -1.16709317e-02
-2.02835103e+00]
[ 6.33297010e-03 -7.77969811e-02  9.95819379e-01  7.43380313e-02
-9.53637969e-01]
[ 9.30956564e-03 -1.16709317e-02  7.43380313e-02  1.36436593e-01
-2.54565331e+00]
[-7.36909408e-01 -2.02835103e+00 -9.53637969e-01 -2.54565331e+00
 1.03496822e+02]]

```

C:\ProgramData\Anaconda3\lib\site-packages\ipykernel\_launcher.py:30: FutureWarning: `rcond` parameter will be deprecated, use `rcond=1e-16` instead, to keep using the old default.

To use the future default and silence this warning we advise to pass `rcond=None`, to keep using the old default.



### 0.2.2 b

```
In [10]: parameters_new = np.array([1,0,0,0,0.1])
log_lik_h0 = log_lik_norm(Data, *parameters_new)
print('hypothesis value log likelihood', log_lik_h0)
log_lik_mle = log_lik_norm(Data, *results2.x)
print('MLE log likelihood', log_lik_mle)
LR_val = 2 * (log_lik_mle - log_lik_h0)
print('likelihood ratio value', LR_val)
pval_h0 = 1.0 - sts.chi2.cdf(LR_val, 5)
print('chi squared of H0 with 5 degrees of freedom p-value = ', pval_h0)
```

```
hypothesis value log likelihood -2253.700688042125
MLE log likelihood 876.8650774281919
likelihood ratio value 6261.131530940634
chi squared of H0 with 5 degrees of freedom p-value = 0.0
```

The results show that it's not likely that age, number of children, and average winter temperature have no effect on the number of sick days.