Problem Set 4

Research method Problem Set 4 due Fri 23th Nov, 23:59

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For Problem Set 4, I use the following packages:

```
import matplotlib.pyplot as plt
import numpy as np
import scipy as sp
import scipy.stats as ss
import math
```

Problem 1

Answer:

Follow the procedure of MLE, first define the PDF and then define the likelihood function. For this case, the beta value is the maximum likelihood value in the trials. The code is below:

```
def time (x, b):
    return 1.0/b * math.exp (-x / b)

def likelihood (trials, arrival):
    problist =[]
    for line in trials :
        prob = 1.
```

```
for l in arrival :

prob = prob * time(l, line)

problist.append(prob)

return problist, trials[problist.index(max(problist))]

arrive = [3.2, 2.1, 5.3, 4.2, 1.2, 2.8, 6.4, 1.5, 1.9, 3.0]

expeTrials = np.linspace(0.0001, 10, 10000)

likely, b = likelihood(expeTrials, arrive)

print 'The max likelihood estimate beta is:', b

plt.figure()

plt.plot (expeTrials, likely)

plt.xlabel ('beta lambda')

plt.ylabel ('Likelihood')
```

I set the trial range np.linspace(0.0001, 10, 10000) and meanwhile, obtain the max value in trials[]. The result is shown in Figure 1 and the plot figure is shown in Figure 2.

The max likelihood estimate beta is: 3.160384428442845

Figure 1

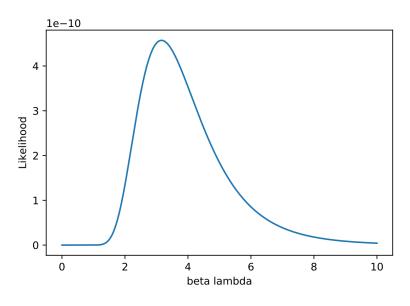


Figure 2

Problem 2

Answer:

1)

I use the equation 1 to calculate the T^2 where X is a vector of sample means, C^{-1} is a vector of the corresponding variance-covariance matrix, and μ is a vector of the corresponding population means. The code is shown below:

$$T^2 = n(ar{oldsymbol{X}} - oldsymbol{\mu})^T oldsymbol{C}^{-1} (ar{oldsymbol{X}} - oldsymbol{\mu})$$

Equation 1

```
def Ex2():

meanMat = np.mat([60, 24, 161])

whole = np.mat([[65, 28, 175], [80, 26, 159], [70, 27, 180], [62, 29, 167], [74, 24, 170]]).T

C = np.cov(whole)
```

```
sample1 = [65, 80, 70, 62,74]

m1 = np.mean(sample1)

sample2 = [28, 26,27, 29, 24]

m2 = np.mean(sample2)

sample3 = [175, 159, 180, 167, 170]

m3 = np.mean(sample3)

meanSample = np.mat([m1, m2, m3])

T_square = 5.0 * (meanSample - meanMat) * np.matrix(C).I * (np.mat([m1, m2, m3]).T-np.mat([60, 24, 161]).T)

F = (5.0 - 3.0) / (3.0*(5-1)) * T_square

Fc = ss.f.ppf(0.99, 3, 5-3)

print T_square, F, Fc

Ex2()
```

The result is:

```
T^2 = 173.22918028,

F = 28.87153005,

Fc = 99.16620137447148

2)
```

At the α = 0.01 significance level, the critical F-value F < Fc therefore there is no sample significantly different from the population means.

Problem 3

1)

The error(within) for DoF is 16 and for SS is 159.89. MSB = SSB/k-1 = SSB/3 = 12.05 and MSW = SSW/n-k= SSW/16 = 9.9931. F = MSB/MSW = 1.2058

Source	DoF	SS	MS	F	P
Factor (between)	3	36.15	12.05	1.2058	0.66
Error (within)	16	159.89	9.9931		
Total	19	196.04			

The code is below:

```
H1 = [98,97,99,96]
H2 = [91,90,93,92]
H3 = [96,95,97,95]
H4 = [95,96,99,98]
def SSW (*arg):
    n = len(arg[0])
     levels = len(arg)
     ssw = 0
     for i in range(levels):
          var =np.var(arg[i], ddof=0)
          ssw = ssw + var
     df =(n*levels)-levels
     return (n* ssw), df
```

```
def SSB (*arg):
    grandmean =0
    levels = len(arg)
    for i in range(len(arg)):
         grandmean = grandmean +np.mean(arg[i])
    grandmean = grandmean /len(arg)
    ssb = 0
    for i in range (len(arg)):
         temp =np.mean(arg[i])
         ssb = ssb + (temp-grandmean) **2
    ssb = len(arg[0])*ssb
    df= levels-1
    return ssb,df
ssw, sswdf = SSW(H1, H2, H3, H4)
ssb, ssbdf = SSB(H1, H2, H3, H4)
sst1 = np.var(H1 + H2 + H3 + H4)* len(H1 + H2 + H3 + H4)
sst2 = ssw + ssb
msw = ssw / sswdf
msb = ssb / ssbdf
F = msb / msw
```

```
Fc=ss.f.ppf(0.95,3,12)

print " SSW =", ssw

print " MSW =", msw

print " SSB =", ssb

print " MSB =", msb

print "F=", F

print "Critical F=", Fc

if F > Fc:

print "F>Fc , Reject H0"

else:

print "F<Fc , Accept H0"
```

The answer is shown in Figure 3:

Figure 3

- 2) SSB = 89.1875
- 3) MSB = 29.72916666666668
- 4) SSE = 22.75
- 6)F = 15.681318681318682 > Fc = 3.490294819497605, so we reject H_0 there is a significant difference between the looms at the 0.05 significance level.