

MedicalCaseStudy

September 7, 2019

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[1]: import numpy as np
from numpy.random import seed
from scipy.stats import friedmanchisquare
import matplotlib.pyplot as plt
# seed the random number generator

[2]: def CreateData(n,p):
    arr = np.zeros(n)
    for i in range(n):
        if(np.random.rand()<p):
            arr[i] = 1
    return arr

[3]: # sample size
seed(1)
n = 50

means = [0.25, 0.23, 0.27, 0.31, 0.28]

data1 = CreateData(n,means[0])
data2 = CreateData(n,means[1])
data3 = CreateData(n,means[2])
data4 = CreateData(n,means[3])
data5 = CreateData(n,means[4])

[4]: # plot
summaryMean = [np.mean(data1), np.mean(data2),np.mean(data3), \
                np.mean(data4),np.mean(data5)]

summaryStd = [np.std(data1), np.std(data2),np.std(data3), \
              np.std(data4),np.std(data5)]

summaryStd = summaryStd/np.sqrt(n)

fig = plt.figure()
ax = fig.add_subplot(111)
```

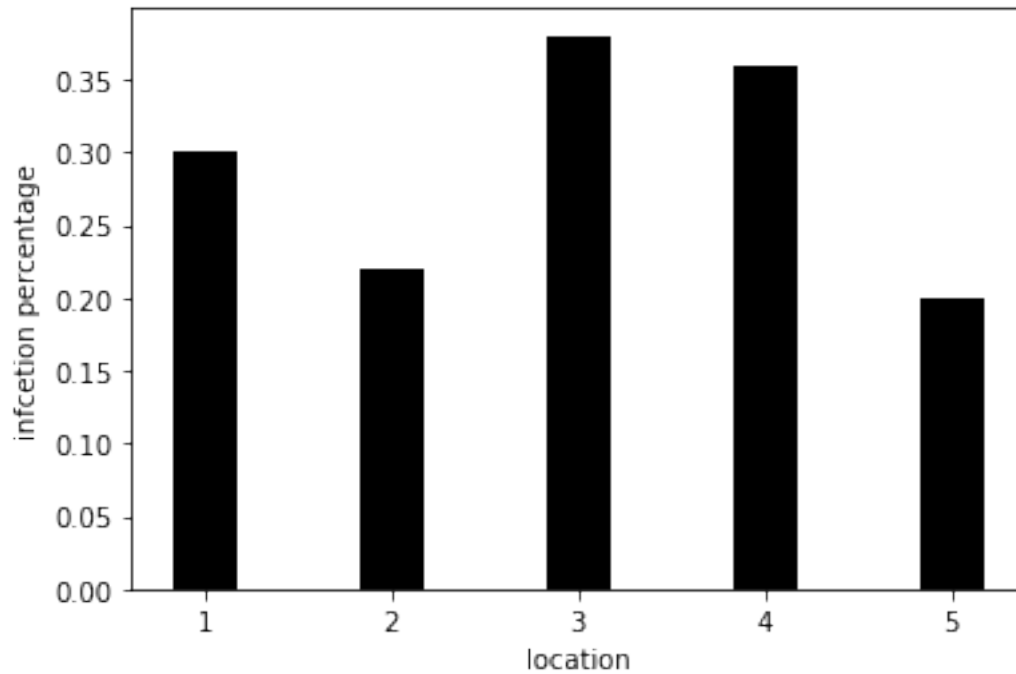
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ax.bar(np.arange(5)+1, summaryMean, 0.35,
       color='black')

plt.xlabel("location")
plt.ylabel("infction percentage")

plt.xlabel("location")
plt.show()

```



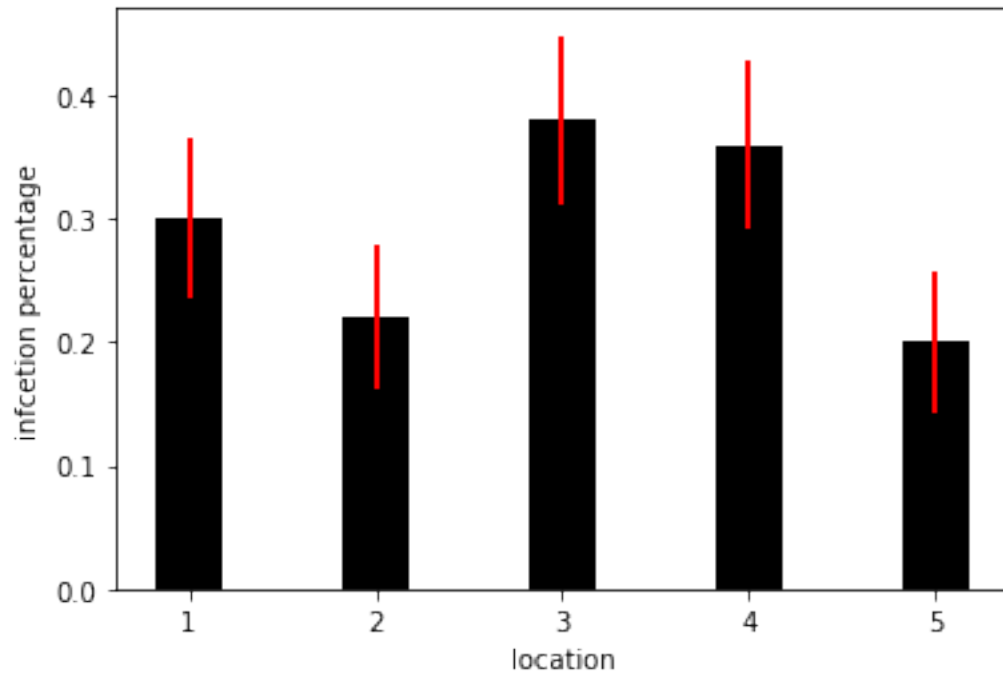
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[5]: fig = plt.figure()
ax = fig.add_subplot(111)

ax.bar(np.arange(5)+1, summaryMean, 0.35,
       color='black',
       yerr=summaryStd,
       error_kw=dict(elinewidth=2,ecolor='red'))

plt.xlabel("location")
plt.ylabel("infction percentage")
plt.show()

```



```
[6]: stat, p = friedmanchisquare(data1, data2, data3, data4, data5)
print('Statistics=%.3f, p=%.3f' % (stat, p))
# interpret
alpha = 0.05
if p > alpha:
    print('Same distributions (fail to reject H0)')
else:
    print('Different distributions (reject H0)')
```

Statistics=6.392, p=0.172
Same distributions (fail to reject H0)