BI-PST/24-25 Homework

Oleksandr Slyvka
slyvkole@cvut.cz

Illia Lyhin
lyhinill@cvut.cz

Maksym Khavil
khavimak@cvut.cz

December 2, 2024

Abstract

With this document we present homework for BI-PST. Oleksandr Slyvka was chosen as a represntative for our group. He was born 14.04.2006, so we get $K = 14$ and $L = 6$, then $M = 4$. Such value of M corresponds to case0202 Sleuth dataset, volume of hippocampus with respect to schizofrenia. We will analyse this dataset using numpy, pandas, matplotlib and scipy.stats Python modules.

1 Data and estimating moments

Firstly, we will load data and display it as a table. There are $n = 15$ samples in both categories, values seem to be greater than 1cm^3 and peak around 2cm^3, each row repesents a pair of twins, one of whom was affected by schizophrenia and other was not.

"Unaffect"	"Affected"
1.94000005722046	1.26999998092651
1.44000005722046	1.62999999523163
1.55999994277954	1.47000002861023
1.58000004291534	1.38999998569489
2.05999994277954	1.92999994754791
1.6599999666214	1.25999999046326
1.75	1.71000003814697
1.76999998092651	1.66999995708466
1.77999997138977	1.27999997138977
1.91999995708466	1.85000002384186
1.25	1.01999998092651
1.92999994754791	1.3400000333786
2.03999996185303	2.01999998092651
1.62000000476837	1.5900000333786
2.07999992370605	1.97000002861023

1.1 Estimations

Let X_{unaff} and X_{aff} denote random variables of hippocampues volume of those who are unaffected and affected by schizofrenia respectively, we will refer to them as unaffected and affected distributions or random variables. We will compute estimates of mean, median and variance of those random variables. Median estimation will be chosen as 7th value of sorted sequence of data points also known as midpoint. Estimated mean and variance will be computed with following formulas:

$$\widehat{\mathbb{E}X} = m_1 = \bar{X} = \frac{1}{n}\sum_{k=1}^{n}X_i$$

$$\widehat{\text{var}X} = s^2 = \frac{1}{n-1}\sum_{k=1}^{n}(X_i - \bar{X})^2$$

Additionaly we will compute uncorrected biased variance.

$$S^2 = \frac{1}{n}\sum_{k=1}^{n}(X_i - \bar{X})^2$$

All those computations are done by the following code:

```
print(f'{'Mean:' :<30}{X.mean():.3f}')
print(f'{'Median:' :<30}{X.median():.3f}')
print(f'{'Variance without correction:' :<30}{((X - X.mean()) ** 2).sum() / n :.3f}')
print(f'{'Variance with correction:' :<30}{((X - X.mean()) ** 2).sum() / (n - 1):.3f}')
```

After plugging values in, we obtain such results.

$$\widehat{\mathbb{E}X_{\text{unaff}}} = 1.759 \qquad\qquad \widehat{\mathbb{E}X_{\text{aff}}} = 1.560$$
$$\widehat{F_{X_{\text{unaff}}}^{-1}(0.5)} = 1.770 \qquad\qquad \widehat{F_{X_{\text{aff}}}^{-1}(0.5)} = 1.590$$
$$\widehat{\text{var}X_{\text{unaff}}} = 0.059 \qquad\qquad \widehat{\text{var}X_{\text{aff}}} = 0.091$$
$$S^2_{\text{unaff}} = 0.055 \qquad\qquad S^2_{\text{aff}} = 0.085$$

Let's notice that estimated expected value of unaffected distribution is greater than estimated expected value of affected distribution.

2 Histograms and empirical cumulitive distribution functions

We will plot histograms, width of bins will be 0.2cm^3. By their side we will plot empirical cumulitive distribution functions, they will have a step of $1/n$ at each sample value.

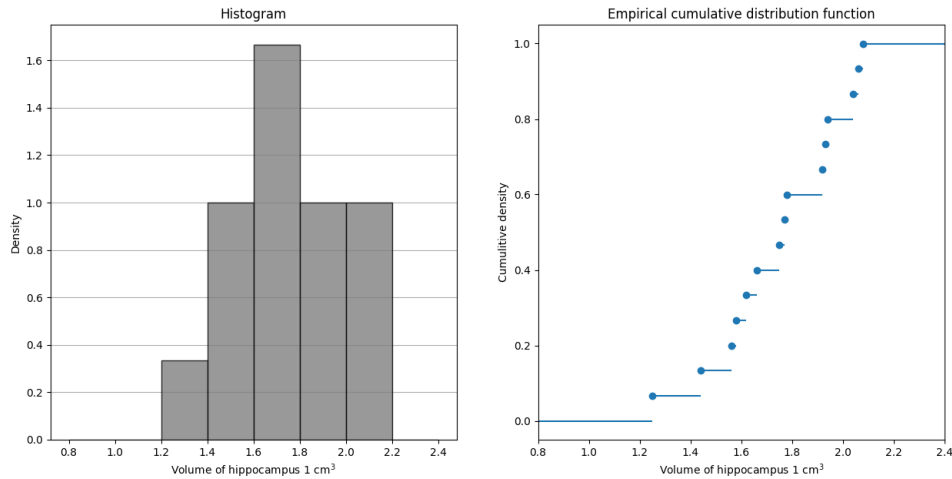Let's plot histogram and ecdf for distribution of unaffected twins.

Figure 1: Unaffected distribution

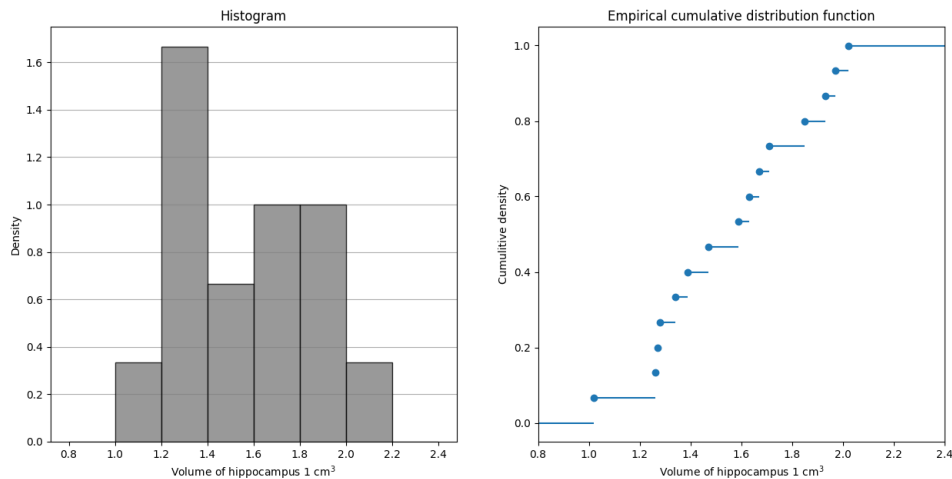Here is the same plot for distibution of schizophrenic twins.

Figure 2: Affected distribution

We can see that later histogram is a little wider, that corresponds with its variance being greater than variance of unaffected distributions. Another observation is that it is altogether slightly shifted towards zero.

3 Choosing between normal, uniform and exponential distributions

We will try to choose distribution for our data that will match it the best. To do that we need to find parameters of distributions, we will calculate them using method of moments.

3.1 Calculating parameters

Normal distribution is parametrized by μ and σ^2, uniform distribution is parametrized by its lower and upper bounds resp. a and b and exponential one is parametrized by the rate λ. All of them can be expressed as functions of at least two moments (exponential can be derived using single moment).

$$\hat{\mu} = m_1$$
$$\hat{\sigma}^2 = m_2 - m_1^2$$
$$\hat{a} = m_1 - \sqrt{3(m_2 - m_1^2)}$$
$$\hat{b} = m_1 + \sqrt{3(m_2 - m_1^2)}$$
$$\hat{\lambda} = \frac{1}{m_1}$$

Those computations are calculated in this code:

```
m1 = X.mean()
m2 = (X ** 2).mean()
mu_hat= m1
sigma2_hat= m2 − m1 ** 2
lambda_hat = 1 / m1
a_hat = m1 − np.sqrt(3 * (m2 − m1**2))
b_hat = m1 + np.sqrt(3 * (m2 − m1**2))
print(f"Normal: {mu_hat=}, {sigma2_hat=}")
print(f"Exp: {lambda_hat=}")
print(f"Uniform: {a_hat=}, {b_hat=}")
```

After we have it run we get following values.

Unaffected	Affected
$\hat{\mu} = 1.758$	$\hat{\mu} = 1.559$
$\hat{\sigma}^2 = 0.054$	$\hat{\sigma}^2 = 0.084$
$\hat{a} = 1.353$	$\hat{a} = 1.055$
$\hat{b} = 2.164$	$\hat{b} = 2.064$
$\hat{\lambda} = 0.568$	$\hat{\lambda} = 0.641$

3.2   Comparing histogram and probability distribution functions
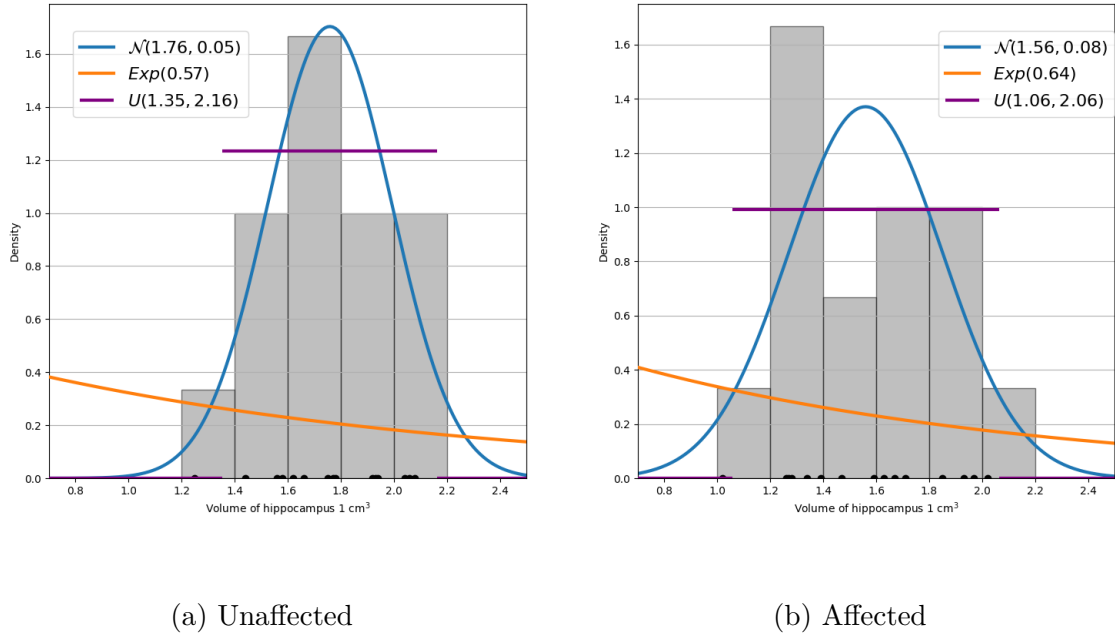

(a) Unaffected   (b) Affected

Figure 3: Histograms and pdfs

In both cases exponential distribution performs horribly, so we will reject it im-mideatly. At both plots there are data points (black dots on x-axis), that lay in zero-probability area for uniform distribution, so it will be rejected too. We are left only with normal distribution, so we naturally choose it, in addition it matches histogram peaks quite nicely.

4   Comparing histograms of 100 generated samples and data

We have computed parameters $\hat{\mu} = 1.76$ and $\hat{\sigma}^2 = 0.05$ for distribution of those, who were not affected, and $\hat{\mu} = 1.56$ and $\hat{\sigma}^2 = 0.08$ for those who were. We will generate a 100 random samples using following code:

```
samples = stats.norm.rvs(
    loc=mu_hat, scale=np.sqrt(sigma2_hat),
    size=100, random_state=42
)
```

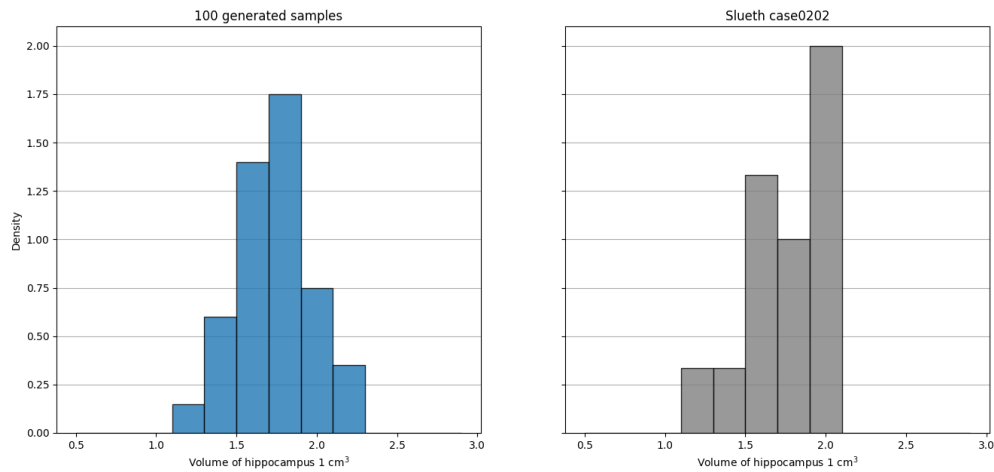Now let's plot histograms and compare them.

Figure 4: Generated (left) and given (right) histograms of unaffected distributions

We can see that artificial one is more gravited towards mean, while given one is skewed right. Besides that they share number of similarities, their ranges match almost perfectly and they both have similar probabilities for peak and tail values.
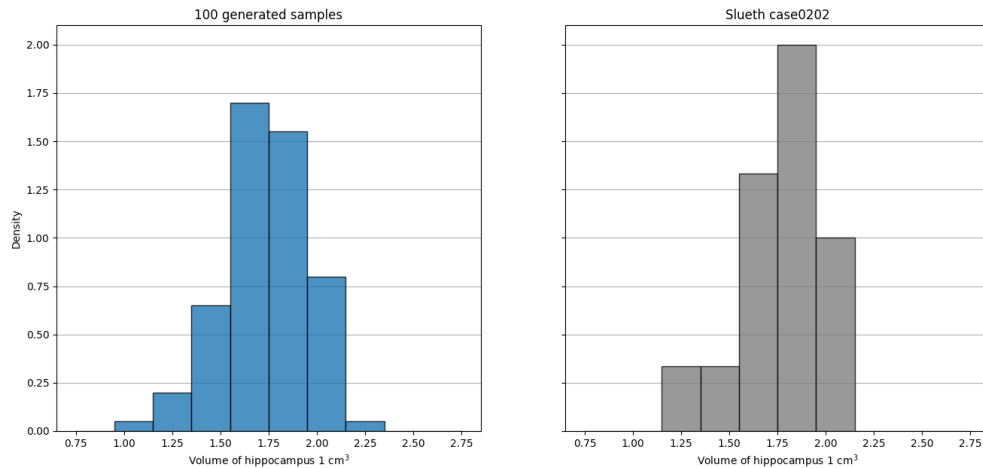
Figure 5: Generated (left) and given (right) histograms of affected distributions

Artificially generated histogram follows shape of data closely, though in this case it tends to be more spread out, with it peak sitting 0.4 points lower and samples reaching further from center.

5 Computing two-sided confidence intervals for expected value at 95% level

We do not have theoretical variance, so we will need to use t distribution to derive our intervals.

From lectures we have formula CI $= \left(\bar{X}_n - \frac{t_{n-1}\left(\frac{\alpha}{2}\right) \cdot s}{\sqrt{n}}, \bar{X}_n + \frac{t_{n-1}\left(\frac{\alpha}{2}\right) \cdot s}{\sqrt{n}} \right)$, we calulate it with next Python code:

```
m = X.mean()
s = np.sqrt(((X - X.mean()) ** 2).sum() / (n - 1))
t_alpha_2 = stats.t.isf(alpha/2, df=n-1)
L = m - t_alpha_2 * s / np.sqrt(n)
R = m + t_alpha_2 * s / np.sqrt(n)
return L, R
```

As a result we obtain these confidence intervals at 95% level for X_{unaff} and X_{aff}:

$$\text{CI}_{\text{unaff}} = (1.6244, 1.8929)$$
$$\text{CI}_{\text{aff}} = (1.3931, 1.7268)$$

6 Testing two-sided hypothesis that expected values equal K at 5% level

Let's formulate hypothesis for unaffected distribution. Null hypothesis will be that expected value of X_{unaff} equals $K = 14$. Hypothesis for affected distribution can be formulated analogically.

$$H_0 : \mathbb{E}X_{\text{unaff}} = K$$
$$H_A : \mathbb{E}X_{\text{unaff}} \neq K$$

To test this hypothesis we would need two-sided confidence intervals for mean. Luckily they were already computed, so we can reject null hypothesis for both affected and unaffected cases, because $K = 14 \notin (1.39, 1.73)$ and $K = 14 \notin (1.62, 1.89)$.

7 Testing if means of affected and unaffected distributionsare equal at 5% level

It is clear that null hypothesis is that expected values are equal. We choose left-sided alternative hypothesis, because estimated mean of affected distribution is lesser than mean of unaffected, so it will be natural to asssume that schizofrenia shrinks one's hippocampus. We have no tests for comparing two different random variables, so we will need to transform our hypothesis in terms of difference of random variables. Let $Y = X_{\text{aff}} - X_{\text{unaff}}$ be such difference, then we can write hypothesis as follows:

$$H_0 : \mathbb{E}X_{\text{aff}} = \mathbb{E}X_{\text{unaff}} \iff \mathbb{E}Y = 0$$
$$H_A : \mathbb{E}X_{\text{unaff}} < \mathbb{E}X_{\text{unaff}} \iff \mathbb{E}Y < 0$$

To test this hypothesis we need to compute left confidence interval for Y. We do not know variance, so we will use formula with t-distribtion $CI_L = (-\infty, \bar{Y} + \frac{t_{n-1}(0.95) \cdot s}{\sqrt{n}})$. We will use almost the same code, that we have used for computing two-sided intervals.

```
Y = X_aff − X_unaff
m = Y.mean()
s = np.sqrt((X ** 2).sum() / (n − 1))
alpha = 0.95

R = m + stats.t.isf(alpha, df=n−1) * s / np.sqrt(n)
print(f"Left CI is: (−inf, {R})")
```

After plugging values in we obtain left confidence interval $(-\infty, -0.34)$. Zero is not covered, so we reject null hypothesis in favor of the alternative.

8 Conclusion

We have conducted analysis of Sleuth case0202 dataset, derived estimates for moments and chose normal distribution for both random variables. Lastly, we computed confidence intervals for means and tested if their expected values are equal by one-sided t-test.

References

[1] BI-PST 24/25, lectures