Topic 2: Sampling distribution of the mean Background for inference of location parameter in location/scale families

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Context

Statistics:

- Functions of random variables
- Therefore, are random variables themselves.
 - In particular, they have their own distributions, called sampling distributions.
 - Meaning of sampling distribution (Review)
- How does inference relate to analytics? (Review)

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 - In particular, they have their own distributions, called sampling distributions.
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Do you know population variance?

Recall the random variable which has a normal distribution by the CLT:

$$Z=\frac{\overline{X}-\mu}{\sigma/\sqrt{n}}.$$

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What happens when you do not know σ ?

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Recall the random variable which has a normal distribution by the CLT:

$$Z = \frac{\overline{X} - \mu}{\sigma / \sqrt{n}}.$$

What happens when you do not know σ ?

Question

What do we know about

$$\frac{\overline{X}-\mu}{S/\sqrt{n}}?$$

Suppose we just take one sample, X, of size 100 from a normal population with mean $\mu = 25$ and standard deviation $\sigma = 10$. Here's our friend, the random variable Z:

```
get.Z.value <- function(sample.size,mu,sigma){
    x <- rnorm(n=sample.size,mean=mu,sd=sigma)
    (mean(x)-mu)/(sigma/sqrt(sample.size))
}</pre>
```

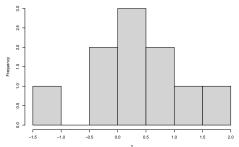
Notice a **function** up there, in the code?

```
my.mu = 25
my.sigma = 10
my.sample.size = 100
get.Z.value(sample.size=my.sample.size,mu=my.mu,sigma = my.s
```

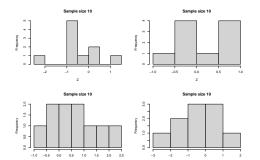
[1] -0.4039858

Of course this is just *one* value of the staistic Z. Now, repeat!



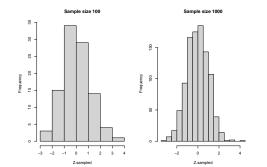


sigma =my.sigma)),main="Sample size 10",xlab="Z")



... What about more reps?

```
par(mfrow = c(1, 2))
Z.sampled <- replicate(100,get.Z.value(
    sample.size=my.sample.size,mu=my.mu,sigma = my.sigma))
hist(Z.sampled, main="Sample size 100")
Z.sampled <- replicate(1000, get.Z.value(
    sample.size=my.sample.size,mu=my.mu,sigma = my.sigma))
hist(Z.sampled, main="Sample size 1000")</pre>
```

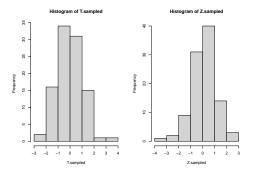


```
Now let's replace \sigma by S.
```

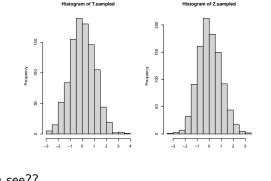
```
get.T.value <- function(sample.size, mu, sigma){
    x <- rnorm(n=sample.size,mean=mu,sd=sigma)
    (mean(x)-mu)/(sd(x)/sqrt(sample.size))
}
my.mu = 25
my.sigma = 10
my.sample.size = 100
get.T.value(sample.size=my.sample.size,mu=my.mu,sigma=my.sigma=my.sigma=my.sigma=my.sigma=my.sigma=my.sigma=my.sigma=my.sigma=my.sigma=my.sigma=my.sigma=my.sigma=my.sigma=my.sigma=my.sigma=my.sigma=my.sigma=my.sigma=my.sigma=my.sigma=my.sigma=my.sigma=my.sigma=my.sigma=my.sigma=my.sigma=my.sigma=my.sigma=my.sigma=my.sigma=my.sigma=my.sigma=my.sigma=my.sigma=my.sigma=my.sigma=my.sigma=my.sigma=my.sigma=my.sigma=my.sigma=my.sigma=my.sigma=my.sigma=my.sigma=my.sigma=my.sigma=my.sigma=my.sigma=my.sigma=my.sigma=my.sigma=my.sigma=my.sigma=my.sigma=my.sigma=my.sigma=my.sigma=my.sigma=my.sigma=my.sigma=my.sigma=my.sigma=my.sigma=my.sigma=my.sigma=my.sigma=my.sigma=my.sigma=my.sigma=my.sigma=my.sigma=my.sigma=my.sigma=my.sigma=my.sigma=my.sigma=my.sigma=my.sigma=my.sigma=my.sigma=my.sigma=my.sigma=my.sigma=my.sigma=my.sigma=my.sigma=my.sigma=my.sigma=my.sigma=my.sigma=my.sigma=my.sigma=my.sigma=my.sigma=my.sigma=my.sigma=my.sigma=my.sigma=my.sigma=my.sigma=my.sigma=my.sigma=my.sigma=my.sigma=my.sigma=my.sigma=my.sigma=my.sigma=my.sigma=my.sigma=my.sigma=my.sigma=my.sigma=my.sigma=my.sigma=my.sigma=my.sigma=my.sigma=my.sigma=my.sigma=my.sigma=my.sigma=my.sigma=my.sigma=my.sigma=my.sigma=my.sigma=my.sigma=my.sigma=my.sigma=my.sigma=my.sigma=my.sigma=my.sigma=my.sigma=my.sigma=my.sigma=my.sigma=my.sigma=my.sigma=my.sigma=my.sigma=my.sigma=my.sigma=my.sigma=my.sigma=my.sigma=my.sigma=my.sigma=my.sigma=my.sigma=my.sigma=my.sigma=my.sigma=my.sigma=my.sigma=my.sigma=my.sigma=my.sigma=my.sigma=my.sigma=my.sigma=my.sigma=my.sigma=my.sigma=my.sigma=my.sigma=my.sigma=my.sigma=my.sigma=my.sigma=my.sigma=my.sigma=my.sigma=my.sigma=my.sigma=my.sigma=my.sigma=my.sigma=my.sigma=my.sigma=my.sigma=my.sigma=my.sigma=my.sigma=my.sigma=my.sigma=my.sigma=my.sigma=my.sigma=my.sigma=my.sigma=my.sigma=my.sigma=my.sigma=my.sigma=my.sigma=my.sigma=my.sigma=
```

[1] -0.7673518

```
Again, need to repeat:
par(mfrow = c(1, 2))
T.sampled <- replicate(100, get.T.value(sample.size=my.sample.size,
hist(T.sampled)
Z.sampled <- replicate(100, get.Z.value(sample.size=my.sample.size,
hist(Z.sampled)
```



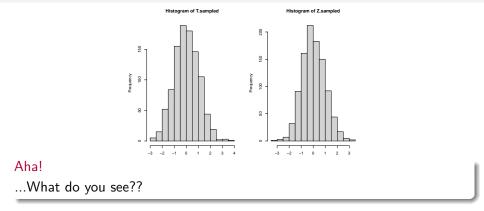
A comparative simulation: the Z vs. T random variable



Aha!

...What do you see??

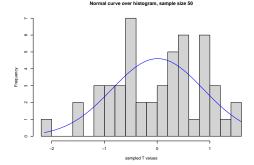
A comparative simulation: the Z vs. T random variable



- The T distribution looks just like Z. Let's use the CLT! ←??
- I don't get it. We just replaced σ by S so... big deal?? \leftarrow ??
- T is a different random variable than Z. The histograms look similar... but can we prove it's the same shape?? ←??

How do we compare the two sampling distributions?

One way: let us plot the T histogram and overlay the the Z density curve on top.



*For HW, you will explore different ways to plot a normal curve over a histogram.

Why?

Because we know the theoretical distribution of Z!

The *t* distribution

Theorem

Let X_1, \ldots, X_n be independent random variables that are all normal with mean μ and standard deviation σ . Let \overline{X} and \S^2 be sample mean and sample standard deviation, respectively. Then Then the random variable

$$T = \frac{\overline{X} - \mu}{\frac{S}{\sqrt{n}}}$$

has a *t*-distribution with $\nu = n - 1$ degrees of freedom.

- What are 'degrees of freedom' and how does n-1 change the shape of $\mathcal{T} \sim t_{n-1}$?
- How is this distribution defined?
 {(By the way, have you ever wondered how is the normal N(μ, σ) defined?
 It's not "just a curve", there's a prob. formula, right?)}

We need some standard notation.

- Discuss: the notation t_{α} .
- Discuss aslo: the notation z_{α} .

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Aha!

Do we get the meaning of these?

An example that needs the *t*-distribution computation

Problem.

A chemical engineer claims that the population mean yield of a certain batch process is 500 grams per milliliter of raw material. To check this claim he samples 25 batches each month. If the computed t-value falls between $-t_{0.05}$ and $t_{0.05}$, he is satisfied with this claim. What conclusion should he draw from a sample that has a mean $\overline{X} = 518$ grams per milliliter and a sample standard deviation s = 40 grams? Assume the distribution of yields to be approximately normal.

An example with t in Python

```
from scipy.stats import t
mean=518
n=25
s=40
u=500
#computing the t-value from the sample
tvalue=(mean-u)/(s/5)
#computing the t0.5,... remember the degree of freedom is 24
interval=t.cdf(0.5,24)
print(-interval)
```

-0.6891856388430067

print(interval)

0.6891856388430067

print(tvalue)

What is next?

- The F-distribution (... almost the last one!)
- The sampling distribution of S^2 .
- Putting it all together, and doing formal statistical tests:
 - Data scenarios
 - Flowchart: which distribution to use when and why?
 - Play with examples and test things out in lab work.

Aha!

Let's get started on some basic questions for "what to use when", shall we? :)

This document is created for ITMD/ITMS/STAT 514, Spring 2021, at Illinois Tech.

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