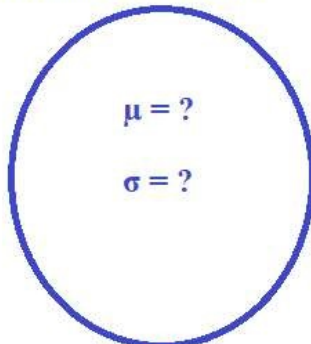


Topic 16 example 2

Hypothesis test for Population Mean, based on the sample mean, σ unknown

Start with a population with unknown mean and standard deviation



We have a **null hypothesis** that the mean is **135.3** and an **alternative hypothesis** that the mean is **not equal to 135.3**.

We want to test that null hypothesis at the **0.03 level of significance**.

H₀: $\mu = 135.3$
H₁: $\mu \neq 135.3$
 $\alpha = 0.03$

We take a random sample of size **32** and we compute the sample mean \bar{x} and the sample standard deviation s_x .

$n = 32$ $\bar{x} = 138.8$ $s_x = 9.2$

Critical value approach: Because this is a two-tailed test we need to find the **t** value such that, for 31 degrees of freedom, $P(X < -t \text{ or } X > t) = 0.03$. We can do this via `high_t <- qt(0.03/2, 31, lower.tail=FALSE)` and, by symmetry of the Student's-t distribution, `low_t` is just the opposite value.

Now we need to translate those values back to the case where the mean is **135.3** and the standard deviation of the sample mean is $9.2/\sqrt{32}$. We do this via `135.3 + low_t*9.2/sqrt(32)` and `135.3 + high_t*9.2/sqrt(32)` giving a critical low value of **131.6007** and a critical high value of **138.9993**.

Then, because the sample mean, **138.8**, is **neither less than the critical low nor greater than the critical high values** we say that we do not have enough evidence to reject **H₀** in favor of **H₁**.

Attained significance approach: We just ask how strange would it be, if **H₀** is true, to get a sample mean this **extreme or more extreme**? To find this we first note that the sample mean is higher than the null hypothesis. Thus, we can find the **probability of getting that sample mean or higher**. First we "standardize" our sample mean, \bar{x} , via `standard_t <- (138.8-135.3)/(9.2/sqrt(32))` to get the test statistic **2.152064**. Then we can use the `pt()` function, with `lower.tail=FALSE`, to find the probability of getting that value or higher.

`p_val <- pt(standard_t, 31, lower.tail=FALSE)`

But that value needs to be **doubled** to account for low values that extreme or more extreme. This gives the value **0.3929236**.

That value is not less than the level of significance, $\alpha=0.03$, so we say that we do not have enough evidence to reject **H₀** in favor of **H₁**.

Of course, we could just use our function, `hypoth_test_unknown()` to do both approaches at once.

```
74 # Or we could have just used our
75 #     hypoth_test_unknown function to compute both
76 #     approaches
77 source("../hypo_unknown.R")
78 #     note the 0 to indicate that H1: mean != 135.3
79 hypoth_test_unknown( 135.3, 0, 0.03, 32, 138.8, 9.2)
```



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```

HO_mu	H1:	std. error
"135.3"	"mu != 135.3"	"1.62634559672906"
n	sig level	t
"32"	"0.03"	"2.27461386471811"
samp mean	samp stdev	test stat
"138.8"	"9.2"	"2.15206411665471"
how far	critical low	critical high
"3.69930824314317"	"131.600691756857"	"138.999308243143"
attained	decision	
"0.0392923649996298"	"do not reject"	