

Creating decision rules and locating rejection regions

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Evidence?

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Test statistic?

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The sample mean.

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Good? Bad? No idea?

- Arbitrary choice of the rejection region.
- Visualize the regions on a number line.

Example with a continuous random variable: solution

Assume that the standard deviation for the population of weights is $\sigma = 3.6$.

For large samples, we may substitute sample stdev (S) for σ , if not other estimate of σ is available.

- Test statistic?

We will use \bar{X} , since this is a test about μ .

- Sampling distribution?

Sample size is $n = 36$. Central limit theorem \implies distribution of \bar{X} is approximately normal with $\sigma_{\bar{X}} = \frac{3.6}{6} = 0.6$.

- Decision rule / rejection region?

Reject $H_0 : \mu = 68$ if $\bar{X} < 67$ or $\bar{X} > 69$.

Plot: test in action

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Q: What is the probability of rejecting when H_0 is actually true?

$$P(\bar{X} < 67 \text{ or } \bar{X} > 69 \text{ when } H_0 \text{ is true}) = P(Z < a) + P(Z > b),$$

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and we compute a, b as: The **z-scores!**:

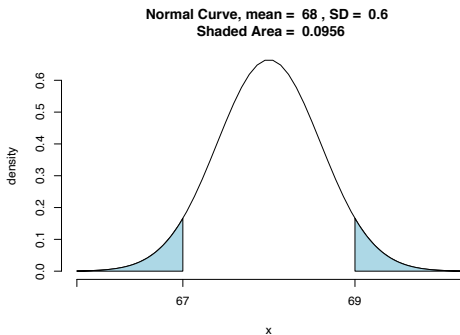
```

true.mu = 68
true.sigma = 3.6
n = 36
a= (67-68)/(true.sigma/sqrt(n)) # a -- the lower cut-off value
b= (69-68)/(true.sigma/sqrt(n)) # b -- the upper cut-off value
c(a,b)

```

```
[1] -1.666667  1.666667
```

```
pnormGC(bound=c(67,69), region="outside",  
        mean=68, sd=true.sigma/sqrt(n),graph=TRUE)
```



```
[1] 0.0955807
```

The level of significance

Uh-oh... 9.5% of all samples of size 36 would lead us to reject $\mu = 68$ kilograms when, in fact, it is true.

Significance level

This error probability is called **level of significance** of the test, and denoted by α .

- Happy? ... seems too high of a chance of error.
- How to fix?
 - Increase the sample size (**try it yourself!**), or
 - Widen the fail-to-reject region.

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But what about H_1 ?

Suppose H_1 is true and $\mu = 70$. What is $P(67 < \bar{X} < 69 \text{ when } \mu = 70)$?

Plot this!

Test significance levels... choices?

Philosophy: Preselection of significance level

Roots of pre-selection of α :

"The maximum risk of making a type I error should be controlled."

- Does not account for values of test statistics that are "close" to the critical region.
- Example: $H_0 : \mu = 10$ vs. $H_1 : \mu \neq 10$. Observed value $z = 1.87$.
 - with $\alpha = 0.05$, value not significant. (no reject)
 - but risk of error:
 $P = 2P(Z > 1.87 \text{ when } \mu = 10) = 2(0.0307) = 0.0614$.
 - 0.0614 is the probability of obtaining a value of z as large as or larger (in magnitude) than 1.87 when in fact $\mu = 10$.
 - \implies Evidence against H_0 is not as strong as that which would result from rejection with $\alpha = 0.05$, but **it is important information to the user**.
 - Indeed, continued use of 'standard' $\alpha = 0.05$ or 0.01 only a result of what standards have been passed down through the generations.

Attained significance level

So how can we tell the user the important information about **strength of evidence**?

The p -value approach, adopted extensively by users of applied statistics, is designed to **give the user an alternative (in terms of a probability) to a mere “reject” or “do not reject” conclusion.**

- The P -value computation also gives the user important information when the z -value falls well into the ordinary critical region.
- For example, if $z = 2.73$, it is informative for the user to observe that $P = 2(0.0032) = 0.0064$, and thus the z -value is significant at a level considerably less than 0.05.
- It is important to know that under the condition of H_0 , a value of $z = 2.73$ is **an extremely rare event.**
 - That is, a value at least that large in magnitude would only occur 64 times in 10,000 experiments!

Graphical representation of p -value

Plot this!

Definition

A p -value is the lowest level (of significance) at which the observed value of the test statistic is significant.