

Testing for Differences: One sample, Independent Samples and Paired Samples t-tests

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Objectives:

- Recognise when to use a one sample t-test, independent samples t-test, paired samples t-test;
- Explain and check the assumptions and conditions for each test;
- Run and interpret a 'One sample t-test' to show a sample mean is different from some hypothesised value;
- Run and interpret an 'independent samples t-test' to show the difference between two groups on one attribute;
- Run and interpret a 'paired samples t-test' to show the difference between two attributes as assessed by one sample

One Sample t-test

Hypothesis Testing for a Single Population Mean;

- The One-Sample T Test compares the mean score of a sample to a known value. Usually, the known value is a population mean.

One-sample t-test for the mean

We test the hypothesis $H_0: \mu = \mu_0$ using the statistic

$$t_{n-1} = \frac{\bar{y} - \mu_0}{SE(\bar{y})},$$

where the standard error of \bar{y} is: $SE(\bar{y}) = \frac{s}{\sqrt{n}}$.

When the conditions are met and the null hypothesis is true, this statistic follows a Student's t -model with $n - 1$ degrees of freedom. We use that model to obtain a P-value.

Assumptions and Conditions

- Independence Assumption The data values should be independent.
- Randomisation Condition The data arise from a random sample.
- 10% Condition The sample size should be no more than 10% of the population.
- Normal Population Assumption The data should be from a population that follows a normal model.

Why do we need to ensure the data is not extremely skewed?
How can we check these assumptions?

Example: One Sample t-test

The mean yield for a certain crop is 8 kg/m². This year, a new pesticide was applied and the yield from nine randomly selected plots are:

9.3 8.2 7.9 8.8 9.4 8.6 8.9 9.5 9.0

Any evidence that yield has increased? - Test the $H_0: \mu = 8.0$ against $H_a: \mu > 8.0$

We create a simple vector of yield using the function `c()`

```
yield<-c(9.3,8.2,7.9,8.8,9.4,8.6,8.9,9.5,9.0) # create an object  
yield      # to show contents of object 'yield'
```

```
## [1] 9.3 8.2 7.9 8.8 9.4 8.6 8.9 9.5 9.0
```

Exploring the data

Before undertaking any statistical analysis it is good to explore the data. We start with numerical summaries.

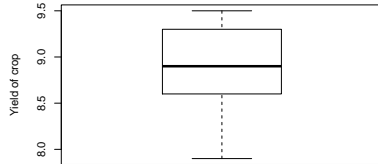
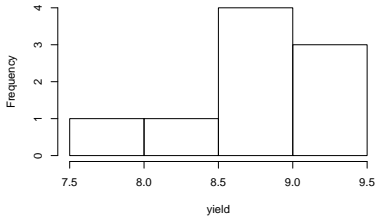
```
summary(yield)  # to summarise the variable 'yield'
```

##	Min.	1st Qu.	Median	Mean	3rd Qu.	Max.
##	7.900	8.600	8.900	8.844	9.300	9.500

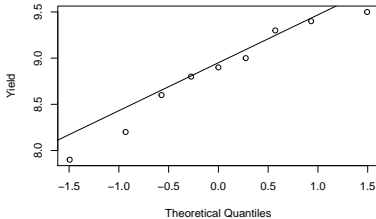
What is the estimated mean yield from the sample?

Before we do the t-test we check the distribution of the data using histogram, boxplot, normal qqplot or stem and leaf.

Yield of crop



Normal Q-Q plot for Yield



What is your comment regarding the normality assumption on the data?

Cont

Finally, we might want a more formal test of agreement with normality (or not). R provides the Shapiro-Wilk test:

```
shapiro.test(yield)           # do a Shapiro-Wilk test on 'yield'
```

Alternatively, we use the Kolmogorov-Smirnov test using `ks.test()` function

```
ks.test(yield, "pnorm", mean = mean(yield),  
        sd = sqrt(var(yield)))
```

Here we are testing the null hypothesis: the data are sampled from the normal distribution. What is your comment on the test results?

Note: Use normality tests with caution:

Let us perform the one sample t-test on the data 'yield' specifying the alternative hypothesis is "greater" as follows:

Cont

```
t.test(yield,mu=8,alt="greater")
```

```
##  
## One Sample t-test  
##  
## data: yield  
## t = 4.6819, df = 8, p-value = 0.000789  
## alternative hypothesis: true mean is greater than 8  
## 95 percent confidence interval:  
## 8.509051 Inf  
## sample estimates:  
## mean of x  
## 8.844444
```

```
# specify hypothesized mean mu=8  
#by default alternative is 2-sided
```