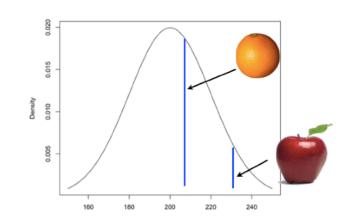
z-scores allow you to compare different variables. reflect distance from the mean in terms of # of standard deviations.



$$z = \frac{X - \mu}{\sigma}$$

z-score of +2 means it is two standard deviations above the mean

Two-sided one-sample t-test in R

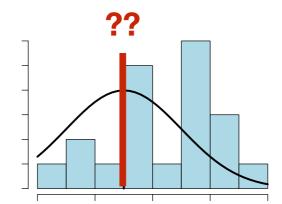
t.test (x=rawdata, mu=M)

rawdata is the rawdata N is the mean you're comparing it against

Cohen's d: measure of effect size

one-sample t-test:

compares numerical data against a theoretically predicted mean



test statistic reflects sampling distribution the distance of your

N-1 degrees of freedom

is a t-distribution with

cohensD(x=rawdata,mu=M)

$$d = \frac{\text{"mean 1"} - \text{"mean 2"}}{\text{"std dev"}}$$

writing up: stats block

0.2

0.5

0.8

interp

small

medium

large

$$t(19) = 3.42, p = .003$$
df t-statistic p-value

$$t = \frac{X - \mu}{\hat{\sigma}/\sqrt{N}}$$

data from the mean



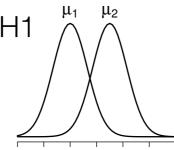
independent samples t-test:

compares two groups (two means)

Assumptions:

- normality
- independent sampling
- Student: groups have same standard deviation
- Welch: groups have different s.d.s

H0 $\mu_1 \quad \mu_2$ $\mu_1 \quad \mu_2$



test statistic: t sampling distribution: t, with N1+N2-2 degrees of freedom

Welch's t-test in R

long form: t.test(outcome~predictor,dataset)

wide form: t.test(x=var1,y=var2)

Student: add var.equal=TRUE



paired samples t-test: just like a onesample t-test on the difference in means

wide form: t.test(x=var1,y=var2,paired=TRUE)

long form: t.test(outcome~predictor,dataset,paired=TRUE)

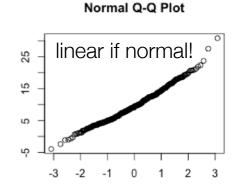


normality: Shapiro-Wilk test. if p<0.05, not normal. but if sample size > 50, may be significant even if normal. look at QQplot / histogram

shapiro.test(var)

<u>Independent-samples</u>: test each variable separately

<u>paired-samples</u>: test the difference variable



Wilcoxon: use if data is not normal. Compares data by "ranks" rather than actual values

wilcox.test(outcome~predictor,dataset)

Add paired=TRUE for paired-sample

Wilcoxon effect size: same interp as Cohens D

library(rstatix)
wilcox.effsize(outcome~predictor,dataset)

Add paired=TRUE for paired-sample