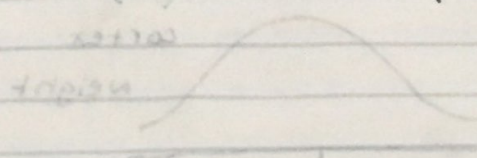


April 19, 2017

This time: experimental designs
Next time: probability
read: OD book



Treatment Variable

(independent variable)

\bar{X}

(supposed causal factor)?
(supposed causal factor)

phy

quant, continuous, ratio
cortex weight (g) after

histogram ✓
mean ✓

Outcome variable

(dependent variable)

\bar{Y}

childhood finished

(response variable)

*enriched ~~psy~~ psychological environment → treatment group (T)

deprived environment → controlled group (C)

Individuals: lab male rats / how many

This is a sample size determination problem

A: 120

How long? → just wait until adulthood

How many in (T), (C)? → 60/60 is the best choice (chain is only as strong as its weakest link)

(T) → cortex width (mg)

689
⋮
649

$n_1 = 60$

(C) → cortex width (mg)

657
⋮
602

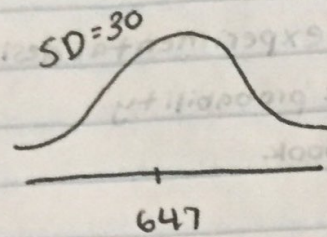
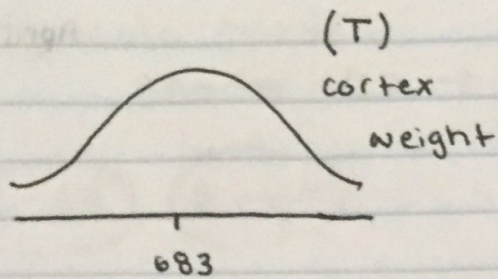
$n_2 = 60$

mean $\bar{y}_1 = 683 \text{ mg}$

SD $s_1 = 32 \text{ mg}$

mean $\bar{y}_2 = 647 \text{ mg}$

$s_2 = 30 \text{ mg}$



Q1: Is the difference between \bar{y}_1 + \bar{y}_2 large in practical terms?

i.e. Is this diff. practically significant? (pratsig)

$$\bar{y}_1 - \bar{y}_2 = (683\text{mg} - 647\text{mg}) = 36\text{mg}$$

Q1:

Is that big enough to matter?

A1: Hard to say w/ our current knowledge

but: Q1 How much bigger is 683 than 647 in relative terms?

$$\frac{683\text{mg} - 647\text{mg}}{647\text{mg}} = \frac{36}{647} = 5.6\%$$

T mean is 5.6% bigger than C means

2 heuristics: ① a small relative change (ex. 1%) can become big if it accumulates over time.

② In many cases, a relative change of 5% diff. is practg., but exceptions exist

$$5.6\% \text{ cortex weight} \rightarrow \approx 5.6^2\% = 31\%$$

increase in synapses (connections between neurons)

new Q

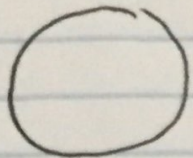
How assign subjects (rats) to T or C?

A] at random (just as w/ sample survey)

the result is called a randomized controlled trial (RCT)

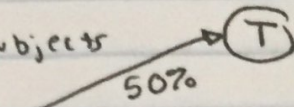
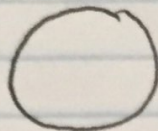
T, C groups

pop. we wish to generalize to

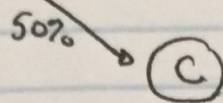


at random

experimental subjects



at random



completely randomized design

outcome \bar{Y} = cortex weight

treatment \bar{X} = enriched vs. deprived

(T)

(C)

Z = genetics

Potential

$Z_1, Z_2 =$

confounding factors

:

(PCFs)

Z_{1k}

↓

def: any third variable Z_i , not \bar{X} or \bar{Y} that satisfies:

1.) $Z_i + \bar{Y}$ may be associated

2.) $Z_i + \bar{X}$ may be associated

quant. $Z_i +$ quant. \bar{Y} associated:

as $\bar{Y} \uparrow$, Z_i tends to \uparrow or \downarrow on average + vice versa

dich $\bar{X} +$ quant. Z_i associated: mean Z_i for $X = 0$ (C) Z_i

mean Z_i for $\bar{X} = 1$ (T) are different

Can't break link between $Z_i + \bar{Y}$ but can break

link between $Z_i + \bar{X}$