

this time: experimental design

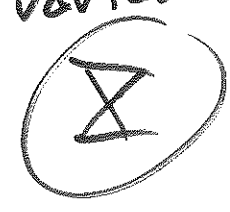
read: DD book (B) ch. 7-8

ANS7 19 Apr 17

next time: probability

today LN pp. L-69 ①

treatment variable (independent variable) (supposedly causal factor)



enriched psychological environment treatment group vs.

control group (deprived environment)

qual. nominal rich

outcome variable (dependent variable) (response variable)



birth neuron (quant. cont. ratio)

cortex weight kg/ after childhood finished

individuals: male rats

how many?

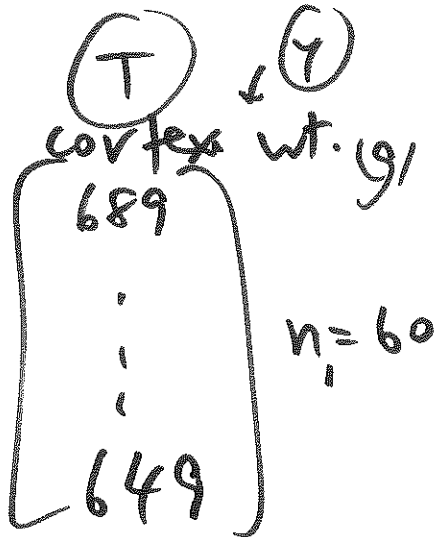
this is a sample size determination problem

A: 120

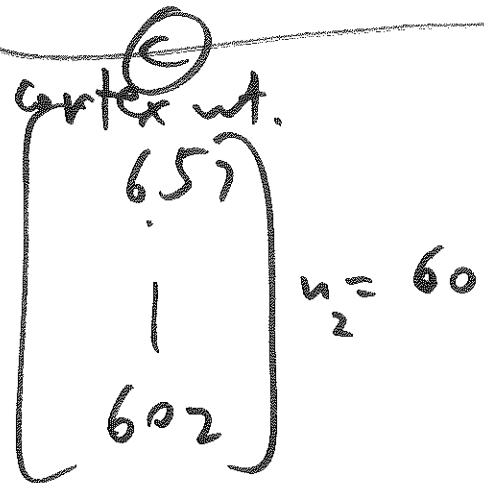
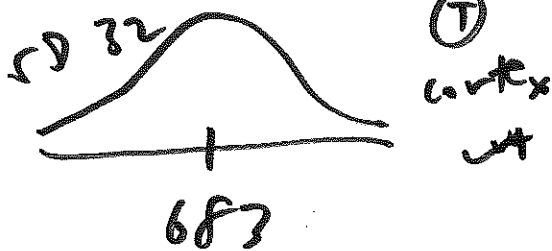
how long? | just wait until adult <sup>(2)</sup>

how many  
in (T), (C)?

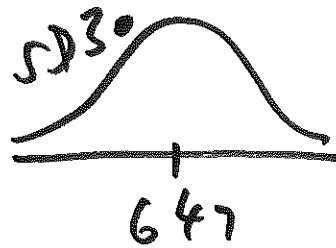
60/60 is best (chain  
is only as strong as  
its weakest link)



mean  $\bar{y}_1 = 683 \text{ mg}$   
SD  $s_1 = 32 \text{ mg}$



mean  $\bar{y}_2 = 647 \text{ mg}$   
SD  $s_2 = 30 \text{ mg}$



Q. Is the difference between  $\bar{y}_1$   
&  $\bar{y}_2$  large in practical terms?  
ie., Is this diff. practically precis significant?

$$\bar{y}_1 - \bar{y}_2 = (683 \text{ mg} - 647 \text{ mg}) = 36 \text{ mg} \quad (3)$$

Q1

is that big enough to matter?

A1

Hard to say with our current knowledge, but:

Q1'

How much bigger is 683 than 647 in relative terms?

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

Ⓣ mean is 5.6% bigger than ⓐ mean

2 heuristics:

① A small relative change (e.g., 1%) can become big if it accumulates over time.

② In many cases a relative change of 5%\* or more is practically sig.  
\* but exceptions exist

5.6% cortex wt  $\rightarrow \approx 5.6^2 \% = 31\%$  ④  
increase in synapses (connections  
between neurons)

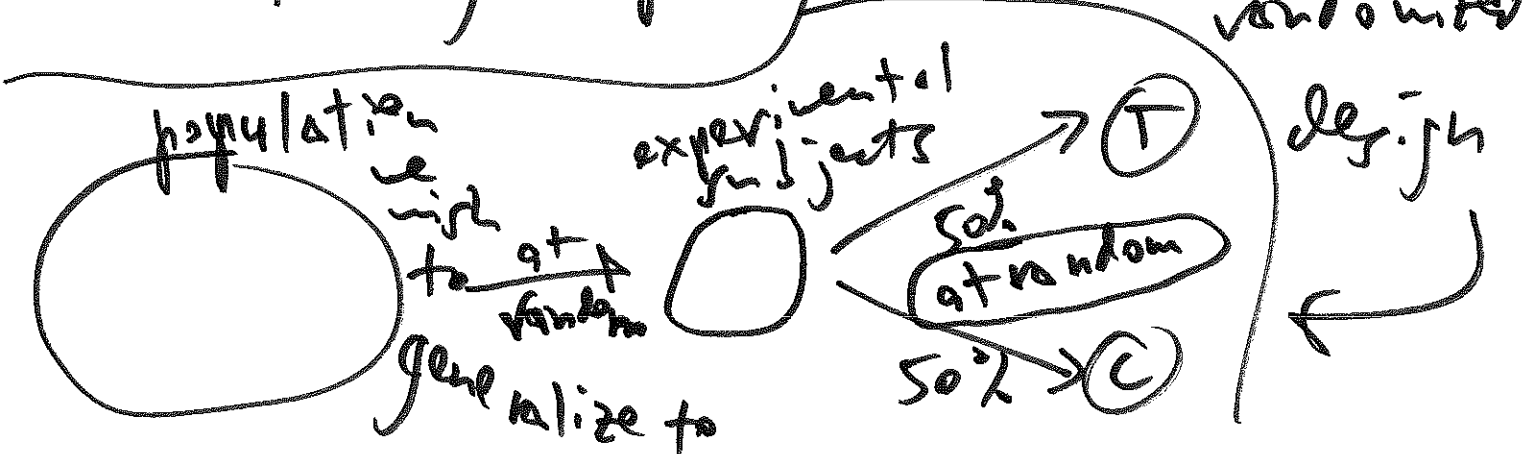
way big

new Q: How assign subjects (rats)  
to T or C?

simplest answer: A at random (just as with  
sample surveys)

The result is called a randomized  
controlled trial (RCT)

T, C groups another name:  
completely randomized



outcome  $Y =$  cortex wt. (5)

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treatment  $X =$  enriched vs. deprived  
(+) (c)

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potential  
confounding  
factors  
(PCFs)

$Z_1 =$  genetics

$Z_2 =$

$\vdots$

$Z_k$

Definition  
of a PCF:

any third variable  $Z_i$

(not  $X$ , no  $Y$ ) that satisfies:

- (1)  $Z_i$  and  $Y$  may be associated
  - (2)  $Z_i$  and  $X$  may be associated
- 

quant.  $Z_i$  & quant  $Y$  associated:

as  $Y \uparrow$ ,  $Z_i$  tends to  $\uparrow$  or  $\downarrow$  on average  
& vice versa

rich  $\mathbb{Z}$  & quant  $\mathbb{Z}$  associated <sup>(6)</sup>:

near  $\frac{1}{2}$  for  $\mathbb{Z} = 0$  (c) &

near  $\frac{1}{4}$  for  $\mathbb{Z} = 1$  (+) are  
different

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can't break link between  $\frac{1}{2}$  &  $\mathbb{Z}$ ,

but can break link between  $\frac{1}{4}$

and  $\mathbb{Z}$ : 2 methods for doing  
so)

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