

this time: exp. design

goal of assignment of experimental subjects to T or C
AMS 7
19 Apr

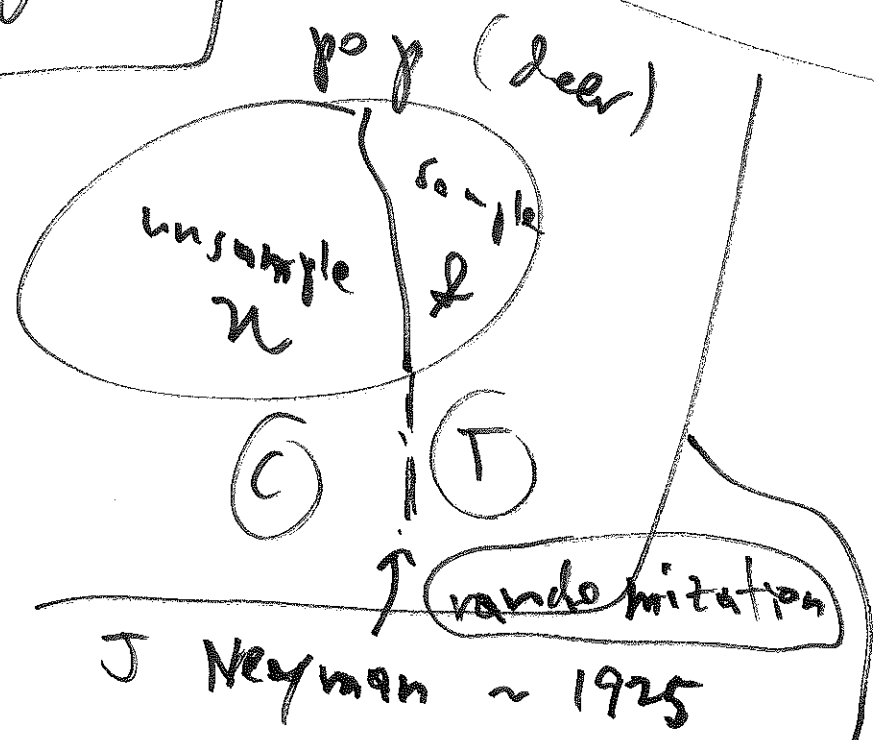
next time: probability

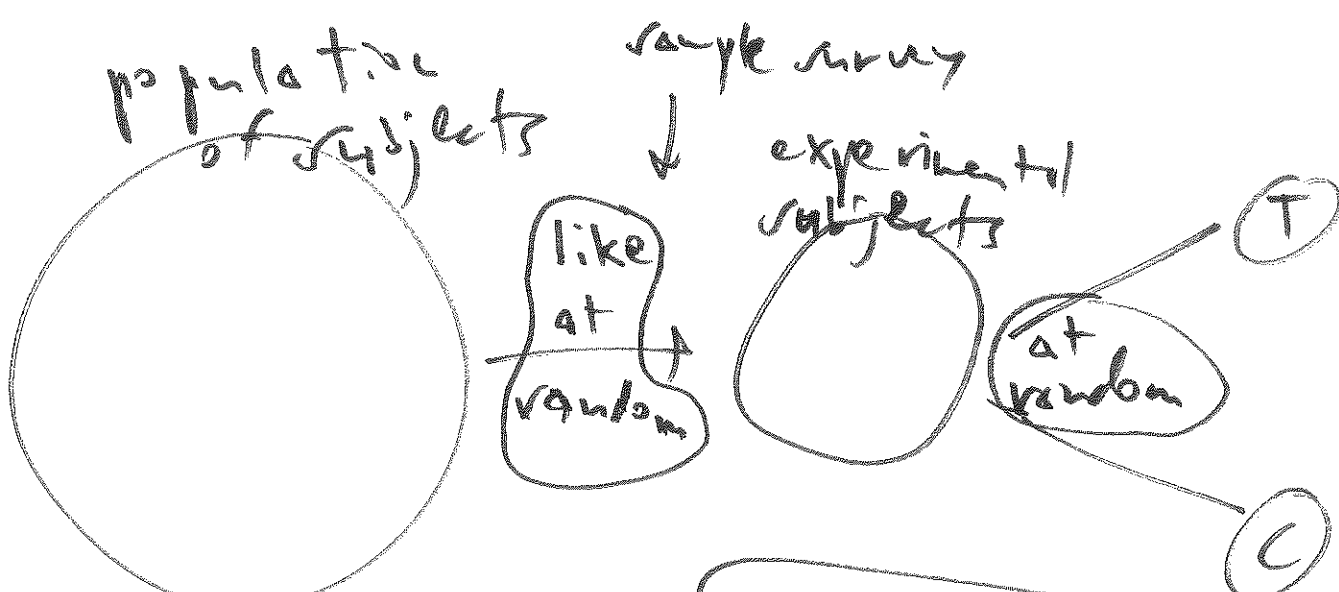
A: T & C groups should be as similar as possible, in all relevant ways, except for T/C distinction

simplest method to achieve this goal

assign to T or C at random

1925
RA Fisher (UK)





to which we wish to generalize

completely randomized design (CRD)

Def: A design is

valid if it's unbiased

Def: A design is unbiased if

- (a) no bias has crept in when (e.g.) assigning subjects to T/C & choosing subjects in first place & (b)

Def: Bias = a systematic tendency to get wrong answer on high or low side

(D) if many people replicated our design, on average across these replications, you would be able to identify truth

Q1: Is the CRD valid?

A1: Yes: no bias anywhere

for drawing causal & effect conclusions

Q2: Might there be other designs that are ~~also~~ also valid but more accurate than CRD?

A2:

Yes

Y (outcome)

X (treatment) (supposedly causal factor (SCF))

Z_1 (potential confounding factor (PCF))

in exp. design, PCFs are the enemy ^④
because without controlling for them
they can bias the causal results

how to test
whether a
 Z_1 is a PCF

① might it be true
that Z_1 and X_1 are
associated (either
positively,
or negatively)

if yes

② might it be
true that Z_1 and X_2 are associated?

if yes to both ① & ②, Z_1 is a
PCF

PLAN AHEAD

two variables U & V are associated

if, when one goes up, the other tends
to go up or down ^{or stay}

psychobiology
care study

I: cortex weight ⑤

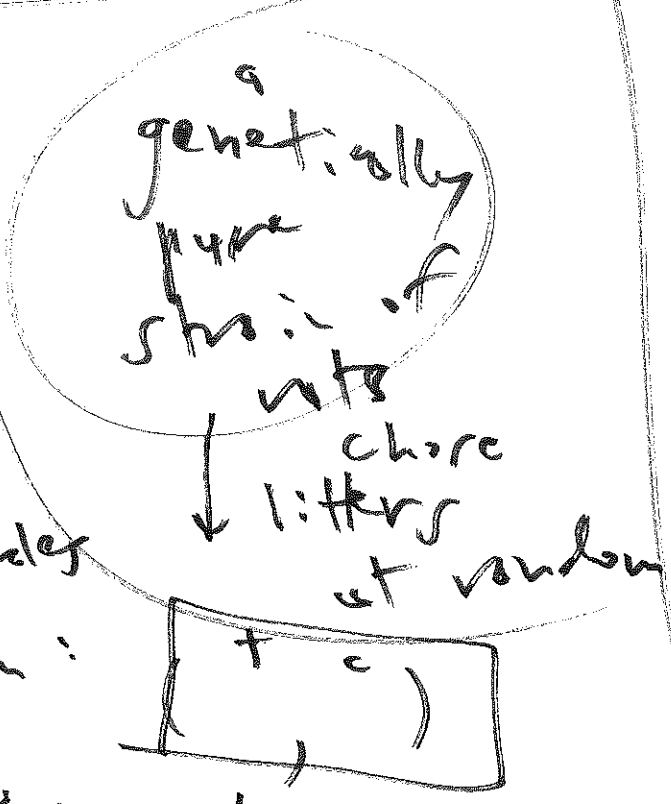
II: psychological environment enriched ① vs. deprived ②

Z: genetics

C/D reflects PCF by breaking link between I & Z

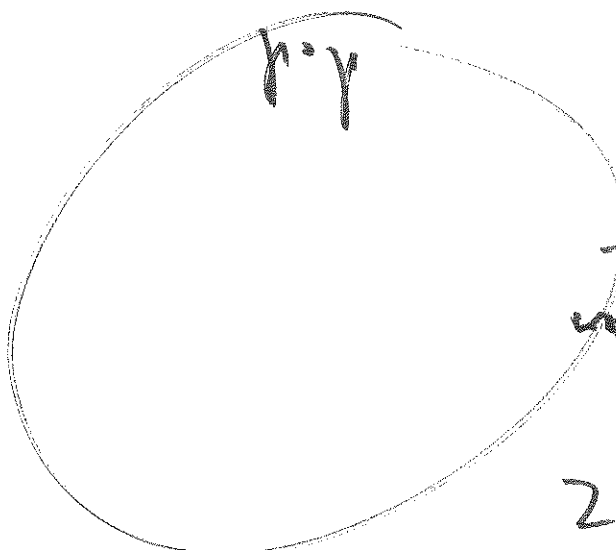
NB all rats were male

A: any given litter with 22 males, chose 2 males at random:



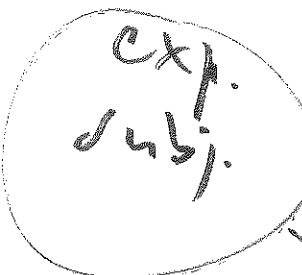
↑ sufficient inside each pair was at random

to defeat PCF we have thoroughly hold them constant in the T/C comparison



CPD

like
at
random

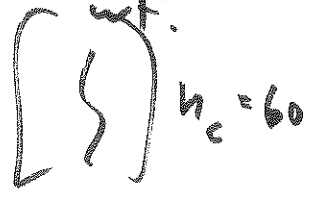


at
random

vertex wt. (6)



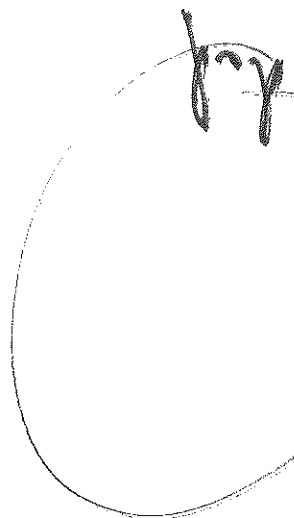
heavy
rd
context
wt.



heavy
sb

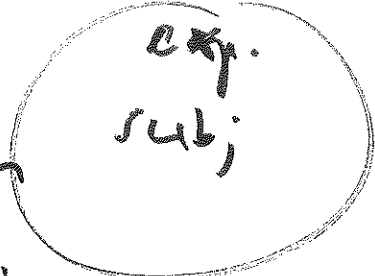
2 - independent
Comparing samples

1 row
for each
at



matched
pairs design

like
at
random



pair #

	T	C
1	708	679
2	695	697

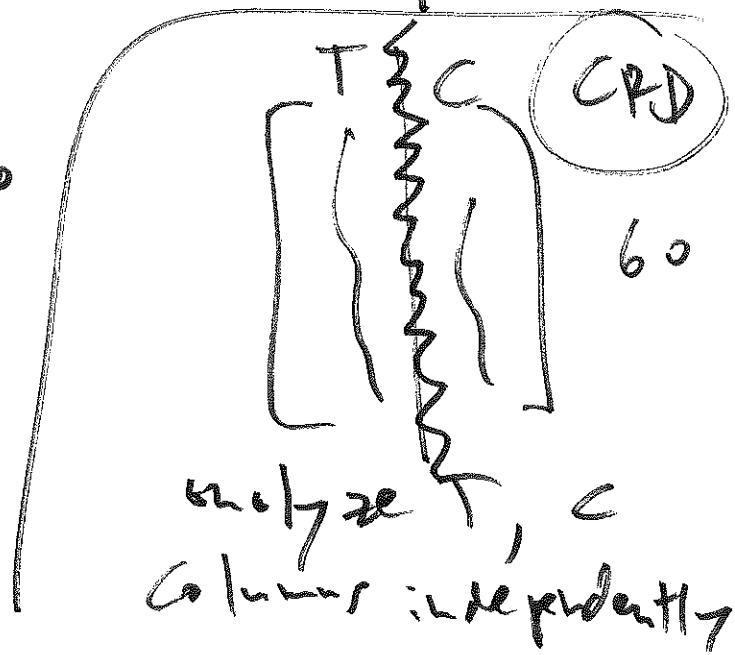
n = 60
1 row
for
each
pair

n = 60

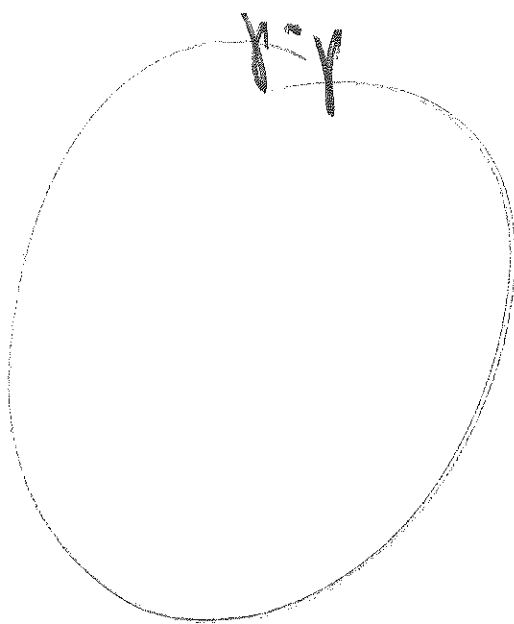
pair	T	C
1	708	679
2	695	697
...		

difference
 $D = T - C$
+ 29
- 2
n = 60

block size = 2
pairs matched on
 $t_2 = \text{genetics}$



analyze T, C
columns independently

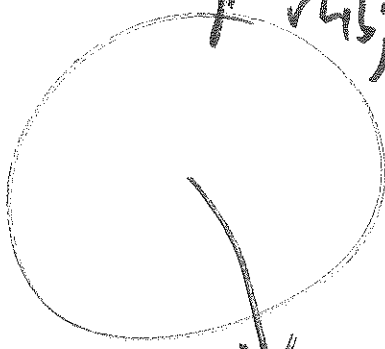


randomized blocks

like at random

exp. rch;

block size = 3



	↑ avoided	↑ normal	↑ deprived
	T	N	C
block 1			

CRD : valid,
not as accurate
as possible

randomized blocks design : valid, likely to be more accurate than CRD

special case:

matched pairs design

(randomized blocks with block size 2)

repeated
measures
design

ex.

T = drug to help with ^{insomnia} (8)

outcome I: ~~less~~ hours of sleep

treatment II: take new drug

make (A) & (B)

pills look identical

or. ~~do it take~~
it

take current
best drug

if (A) pill has no active ingredient,
it's called a ~~place~~ placebo

reason
for why:

placebo effect:

people tend to respond to the
idea of treatment, in addition
to or instead of the (A) itself

Hawthorne
effect:

people change behavior when
they know they're being watched

⊕, ⊙ pill look identical: blinding ⊙
subjects to T/C status

vs. good idea to blind experimenters
to T/C status

both blinds:

double-blind experiment

offer not ethically possible to
randomize people to ⊕^{eg. smoking}
vs. ⊙^{not smoking}

Y: health status (lung cancer? heart disease?)
X: ⊕ smokers
⊙ nonsmokers

Z_1, Z_2, \dots lots of PCFs } observational studies