

this time: 2-sample paired comparisons
next time: 2 independent samples

read:
LN pp. L-180+
L-213
today:
LN pp. L-188 →

2-Sample Problem
① paired comparisons

CORTEX STUDY	pair #	(difference)		
		T	C	D = T - C
	1	689	657	+32
	2	656	623	+33
	59	649	602	+47
	mean	$\bar{y}_T = 683$	$\bar{y}_C = 647$	$\bar{d} = +36.2$
	SD	$s_T = 32$	$s_C = 30$	$s_d = 31.5$

* here, there are $2(59) = 118$ individuals grouped into $n = 59$ pairs by holding PCF (genetics) constant

* 2-sample problem converts into a 1-sample problem when attention in inference is focused on the difference column

← blood pressure →

SYSTOLIC BP STUDY	person	before	after	diff. (A - B)
		1	}	}
	2			
	⋮			
	n			
	mean	$\bar{y}_B = 174$	$\bar{y}_A = 160$	$\bar{d} = -14$ ← statsig
	SD	$s_B = 20$	$s_A = 21$	$s_d = 11$ ←

* here, there are only n individuals, not $2n$ as in matched pairs

* likely to be accurate because 2 measurements per person, which holds entire person constant

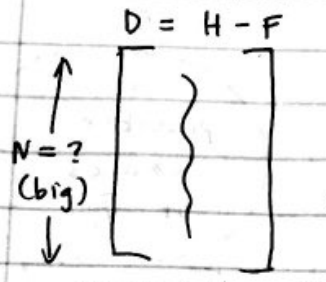
$$\frac{\bar{y}_A - \bar{y}_B}{\bar{y}_R} = \frac{160 - 174}{174} = -0.08 = -8\% \text{ (practsig)}$$

DEER STUDY

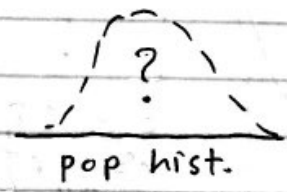
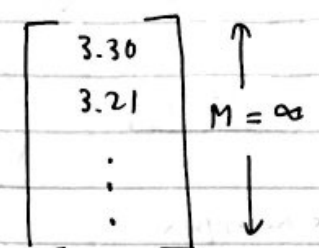
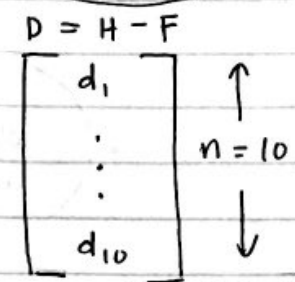
pop all relevant deer for generalization

sample the observed deer

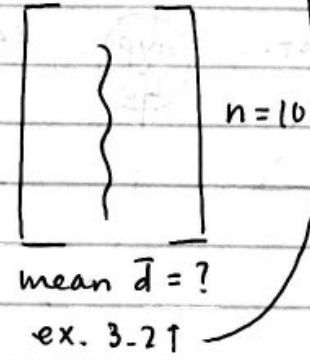
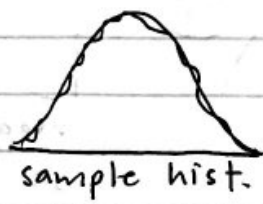
I.D. all possible values of \bar{d}



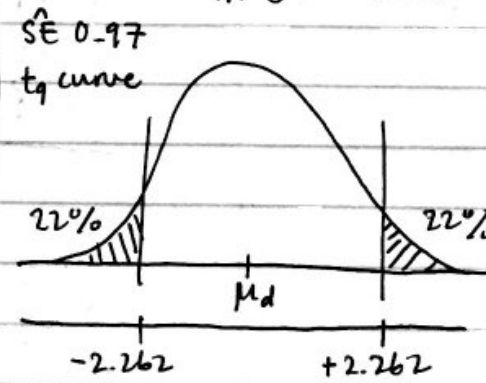
actual like SRS = IID



hyp IID



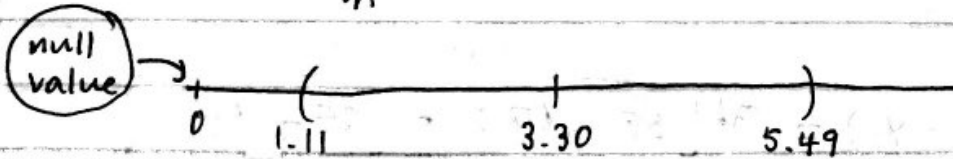
long run mean EV of $\bar{d} = \mu_d$
 est. long run SD \hat{SE} of $\bar{d} = 0.97$
 long run hist. of \bar{d} , accounting uncertainty in σ



Inferential Summary

Ⓟ	unknown pop quantity of main interest	$\mu_d = \text{pop. mean diff. in } \textcircled{H} \text{ vs. } \textcircled{F} \text{ length}$
Ⓢ	estimate	$\bar{d} = 3.30 \text{ cm}$
Ⓜ	give or take for \bar{d} as est. of μ_d	$\hat{SE}(\bar{d}) = 0.97 \text{ cm}$
Ⓜ	95% CI for μ_d	$\bar{d} \pm 2.262 \hat{SE}(\bar{d}) = (1.11, 5.49)$

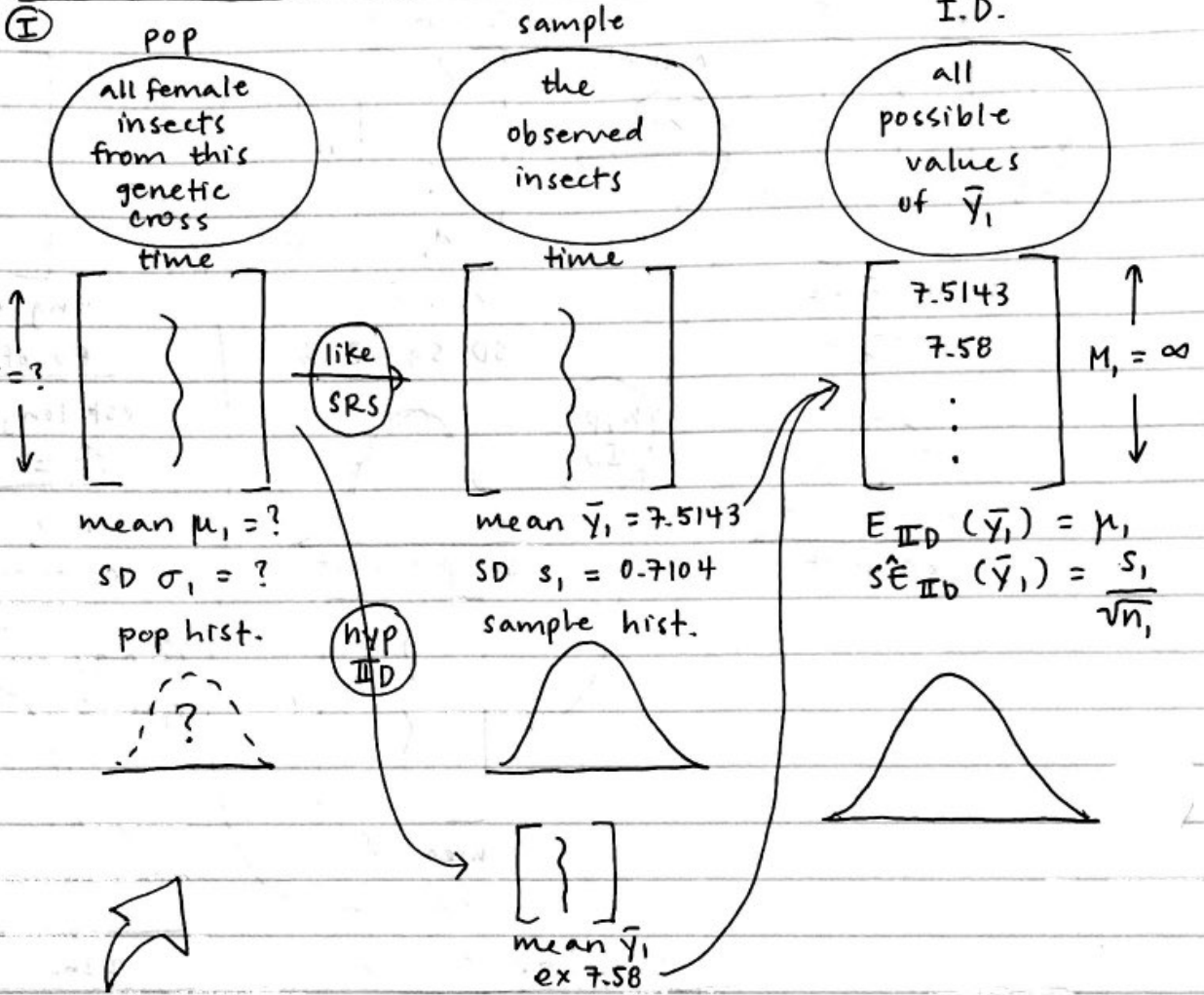
$$\bar{d} \pm (t_{n-1}^{0.95}) \frac{s}{\sqrt{n}} = 3.30 \pm (2.262)(0.97) = 3.30 \pm 2.19$$



* 0 is not in 95% CI, so diff is statsig.

2 independent samples

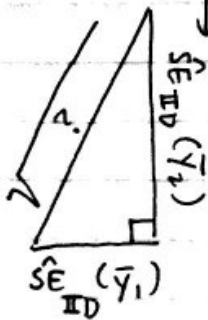
Sokal & Rohlf (1995)



II repeat (ditto for genetic cross II)

Inferential Summary

(P) unknown pop. quantity of interest	$(\mu_2 - \mu_1) =$ pop mean diff in time to reproduce b/t gen cross I & II
(S) estimate	$(\bar{y}_2 - \bar{y}_1) = 7.5571 - 7.5143 = +0.0428$ days
give or take	$\hat{SE}_{ID}(\bar{y}_2 - \bar{y}_1) = 0.3614$ days
$(\bar{y}_2 - \bar{y}_1)$ as est of $(\mu_2 - \mu_1)$	
95% CI $(\mu_2 - \mu_1)$	$(\bar{y}_2 - \bar{y}_1) \pm (2.179) \hat{SE}_{ID}(\bar{y}_2 - \bar{y}_1)$



$$\hat{SE}_{ID}(\bar{y}_2 - \bar{y}_1) = \sqrt{[\hat{SE}_{ID}(\bar{y}_1)]^2 + [\hat{SE}_{ID}(\bar{y}_2)]^2}$$

$$= \sqrt{\left(\frac{s_1}{\sqrt{n_1}}\right)^2 + \left(\frac{s_2}{\sqrt{n_2}}\right)^2} = \sqrt{\frac{s_1^2}{n_1} + \frac{s_2^2}{n_2}}$$

$$\hat{SE}_{\text{ID}}(\bar{y}_1) = \frac{S_1}{\sqrt{n_1}} = \frac{0.7104 \text{ days}}{\sqrt{7}} = 0.2685 \text{ days}$$

$$\hat{SE}_{\text{ID}}(\bar{y}_2) = \frac{S_2}{\sqrt{n_2}} = \frac{0.6399 \text{ days}}{\sqrt{7}} = 0.2419 \text{ days}$$

* uncertainty (SE) in $(\bar{y}_2 - \bar{y}_1)$ is related to uncertainty in \bar{y}_1 (SE) & \bar{y}_2 (SE)

$$\hat{SE}_{\text{ID}}(\bar{y}_2 - \bar{y}_1) = \sqrt{(0.2685)^2 + (0.2419)^2} = \boxed{0.3614 \text{ days}}$$

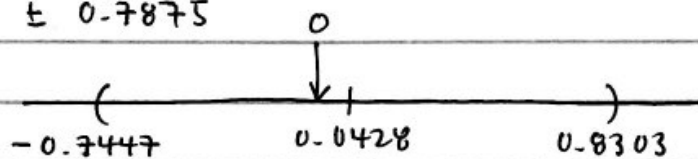
OR

$$\hat{SE}_{\text{ID}}(\bar{y}_2 - \bar{y}_1) = \sqrt{\frac{0.7104^2}{7} + \frac{0.6399^2}{7}} = \boxed{0.3614 \text{ days}}$$

$$(\bar{y}_2 - \bar{y}_1) \pm 2.179 \hat{SE}(\bar{y}_2 - \bar{y}_1)$$

$$= 0.0428 \pm 2.179 (0.3614)$$

$$= 0.0428 \pm 0.7875$$



* not statsig *