

Why?

Under (3)

If $\rho_{XY} > 0$ ($\Leftrightarrow \sigma_{XY} > 0 \Leftrightarrow \text{Cov}[\phi^*(Z), \psi^*(Z)] > 0$), then paired sample is more effective than independent samples.

When $\psi^*(Z) = \phi^*(Z) - \Delta$,

$$\begin{aligned}\text{Cov}[\phi^*(Z), \psi^*(Z)] &= \text{Cov}[\phi^*(Z), \phi^*(Z) - \Delta] \\ &= \text{Cov}[\phi^*(Z), \phi^*(Z)] \\ &= \text{Var}[\phi^*(Z)] > 0.\end{aligned}$$

1/13

Q: Why are independent samples more effective than paired samples when $\sigma_{XY} < 0$?

If $\sigma_X^2 = \sigma_Y^2 = \sigma^2$, then in the paired case

$$\sigma_D^2 = \text{Var}(\bar{D}) = [2\sigma^2(1 - \rho_{XY})]/n \quad \text{and} \quad \sigma_{\bar{X}-\bar{Y}}^2 = \text{Var}(\bar{X}-\bar{Y}) = 2\sigma^2/n$$

in the unpaired case. The relative efficiency is $\sigma_D^2 / \sigma_{\bar{X}-\bar{Y}}^2 = 1 - \rho_{XY}$.

If $\rho_{XY} = 0.5$, a paired design with n pairs of subjects yields the same precision as an unpaired design with $2n$ subjects per treatment.

From now on, the analyses of paired data are based on

$$D_i = X_i - Y_i, \quad i = 1, \dots, n.$$

Statistical modeling for D_i 's: $D_1, \dots, D_n \sim \text{i.i.d. } F \Leftarrow \text{one-sample model}$

• paired
n X_i 's
n Y_i 's
• 2-indep.
2n X_i 's
2n Y_j 's

σ_D^2
 $\sigma_{\bar{X}-\bar{Y}}^2$
under (3)

Example 7 (Effect of cigarette smoking on platelet aggregation, Levine, 1973)

- Blood samples were drawn from 11 individuals before and after they smoked a cigarette to measure the extent to which the blood platelets aggregated.
- data: maximum percentage of all platelets that aggregated after being exposed to a stimulus.

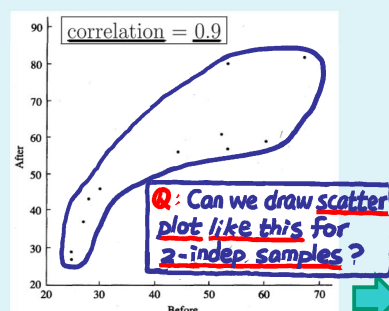
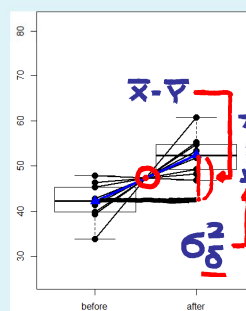
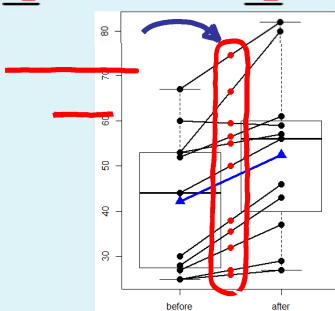
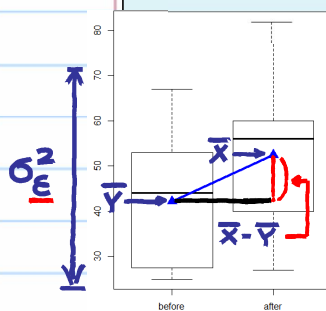
Q: Can we take this difference on the X 's, Y 's data from 2-indep. samples?

block (LNp.46) →

paired

	1	2	3	4	5	6	7	8	9	10	11
before (Y)	25	25	27	44	30	67	53	53	52	60	28
after (X)	27	29	37	56	46	82	57	80	61	59	43
difference (D)	2	4	10	12	16	15	4	27	9	-1	15

- Q: Do the differences D_i 's indicate a clear pattern of $\Delta = \mu_X - \mu_Y \neq 0$?
- The two-sample (unpaired) t -test for the before and after data gives a p -value = 0.1721 \Rightarrow the null $H_0: \mu_X = \mu_Y$ is not rejected under $\alpha = 0.1$ ($s_p^2 = 289.34$).
 - Q: Why did the 2-sample t -test not reject H_0 when the differences showed such a clear pattern of $\mu_X > \mu_Y$?
 - Note that in 2-sample t -test, the test statistic is $|T| = \frac{|\bar{X} - \bar{Y}|}{s_p \sqrt{\frac{1}{n} + \frac{1}{m}}}$, where s_p^2 estimates σ_ϵ^2 , rather than σ_D^2 .



Q: Can we draw scatter plot like this for 2-indep. samples?

- Figure 11.7 (textbook, p.447) plots after-values vs. before-values. They are positively correlated with a sample correlation coefficient 0.9. Pairing was a natural and effective experimental design in this case: relative efficiency = 0.1.

Methods based on normality assumptions

- Recall.** $D_i = X_i - Y_i$, $i = 1, \dots, n$, and $D_1, \dots, D_n \sim \text{i.i.d. } F$. **1-sample model**
- Assume that F is Normal.
- Thus, the statistical model for D_i 's is

where $\mu_D = \mu_X - \mu_Y$. $D_1, \dots, D_n \sim \text{i.i.d. } N(\mu_D, \sigma_D^2)$, **1-sample normal model** (Δ)

– This model contains two parameters: $\mu_D (\in \mathbb{R})$ and $\sigma_D^2 (> 0)$.

Under this model, we can only examine whether there exists “difference” between the means of the two paired samples, i.e.,

$$\mu_D = \mu_X - \mu_Y = 0 \Rightarrow \text{no difference or no effect}$$

cannot examine $\sigma_x^2 \neq \sigma_y^2$

Theorem 18 (test and confidence interval for μ_D , 1-sample normal model, paired data)

Consider the model (Δ).

- Recall** (Review 1 in LNp.6-7).
 - $\bar{D} = \frac{1}{n} \sum_{i=1}^n D_i \xrightarrow{e} \mu_D$, and $\bar{D} \sim N(\mu_D, \sigma_D^2/n) \Rightarrow \sqrt{n}(\bar{D} - \mu_D)/\sigma_D \sim N(0, 1)$

standardization \rightarrow *statistic* \rightarrow *pivotal quantity for μ_D if σ_D known*
 - $s_D^2 = \frac{\sum_{i=1}^n (D_i - \bar{D})^2}{n-1} \xrightarrow{e} \sigma_D^2$, and $(n-1)s_D^2 \sim \sigma_D^2 \chi_{n-1}^2 \Rightarrow (n-1)s_D^2/\sigma_D^2 \sim \chi_{n-1}^2$

statistic \rightarrow *pivotal quantity for σ_D^2*
 - \bar{D} and s_D^2 are independent

- Pivotal quantity for μ_D

– σ_D known: $Q_{Z, \mu_D} \equiv \frac{\bar{D} - \mu_D}{\sigma_{\bar{D}}} = \frac{\bar{D} - \mu_D}{\sigma_D/\sqrt{n}} = \frac{\sqrt{n}(\bar{D} - \mu_D)}{\sigma_D} \sim N(0, 1)$

– σ_D unknown: $Q_{T, \mu_D} \equiv \frac{\bar{D} - \mu_D}{s_{\bar{D}}} = \frac{\bar{D} - \mu_D}{s_D/\sqrt{n}} = \frac{\sqrt{n}(\bar{D} - \mu_D)/\sigma_D}{\sqrt{\frac{(n-1)s_D^2/\sigma_D^2}{n-1}}} \sim \frac{t_{n-1}}{\sim \chi_{n-1}^2}$

μ_D : fixed, data: changed

- Test the null and alternative hypotheses at significance level α :

$$H_0 : \mu_D = \mu_{D,0} \quad \text{vs.} \quad H_A : \mu_D \neq \mu_{D,0}$$

where $\mu_{D,0}$ is a known constant.

(Note. if $\mu_{D,0} = 0$, this is equivalent to $H_0 : \mu_X = \mu_Y$ vs. $H_A : \mu_X \neq \mu_Y$.)

- σ_D known: reject H_0 if

The two tests are likelihood ratio tests (exercise)

$$|Z| \equiv \frac{|\bar{D} - \mu_{D,0}|}{\sigma_{\bar{D}}} > z(\alpha/2) \Leftrightarrow |\bar{D} - \mu_{D,0}| > z(\alpha/2) \sigma_{\bar{D}} = z(\alpha/2) \frac{\sigma_D}{\sqrt{n}}$$

- σ_D unknown: reject H_0 if

$$|T| \equiv \frac{|\bar{D} - \mu_{D,0}|}{s_{\bar{D}}} > t_{n-1}(\alpha/2) \Leftrightarrow |\bar{D} - \mu_{D,0}| > t_{n-1}(\alpha/2) s_{\bar{D}} = t_{n-1}(\alpha/2) \frac{s_D}{\sqrt{n}}$$

- A $100(1 - \alpha)\%$ confidence interval for μ_D is **(data fixed, μ_D : changed)**

– σ_D known: $\bar{D} \pm z(\alpha/2) \times \sigma_{\bar{D}} \Leftrightarrow \bar{D} \pm z(\alpha/2) \times (\sigma_D/\sqrt{n})$

– σ_D unknown: $\bar{D} \pm t_{n-1}(\alpha/2) \times s_{\bar{D}} \Leftrightarrow \bar{D} \pm t_{n-1}(\alpha/2) \times (s_D/\sqrt{n})$

estimate difference between μ_D & $\mu_{D,0}$

scale

2-sided tests

paired samples

one-sample Z-test
one-sample T-test

Example 8 (Effect of smoking, t -test for paired data, cont. Ex.7 In LNp.52)

- $n = 11$, $D_i = \text{after}_i - \text{before}_i$ estimate of population variance σ_D^2
- $\bar{D} = 10.27$, $s_{\bar{D}} = 2.405$ cf. $\Rightarrow s_D^2 = 11 \times 2.405^2 = 63.62$
 $\Rightarrow 63.62/2 = 31.81 \xrightarrow{\text{cf.}} \sigma_D^2 \xrightarrow{\text{cf.}} s_p^2 = 289.34 \xrightarrow{\text{cf.}} \sigma_\epsilon^2$ in Ex.7) include variation of Z_i 's

If $\sigma_{\delta_1}^2 = \sigma_{\delta_2}^2 = \sigma_\delta^2$,
 then (LNp.49)
 $\text{Var}(D_i) = 2\sigma_\delta^2$

Same as $\bar{x} - \bar{y}$
 in 2-sample
 T-test (Ex.7)

$\alpha = 0.1$

- A 90% confidence interval for μ_D is cf.
 $\bar{D} \pm t_{10}(0.05) s_{\bar{D}} = 10.27 \pm 1.812 \times 2.40 = (5.9, 14.6)$, → reject $\Delta = \mu_x - \mu_y = 0$
 which does not contain zero (cf. H_0 not rejected in Ex.7 using 2-sample t -test)
- The (one-sample) t -statistic is $T = (10.27 - 0)/2.40 = 4.28 > t_{10}(0.005) = 3.169$.
 The p -value of a two-sided test is less than 0.01. There is little doubt that smoking increases platelet aggregation. ↑ α

Note 9 (Some notes about one-sample t -test when normality assumption does not hold)

- Consider the model: $D_1, \dots, D_n \sim \text{i.i.d. } F$,
 where F can be any continuous distributions with finite variance. ← can use 1-sample T-test
- By CLT and LLN, when $n \rightarrow \infty$ (sample size is large), → can use 1-sample T-test
 $\bar{D} \xrightarrow{D} N(\mu_D, \sigma_D^2/n)$ and $s_D^2 \xrightarrow{P} \sigma_D^2$. ← $\bar{D} \xrightarrow{D} N(0.1)$
- Thus, by Slutsky's Thm,
 $Q_{T, \mu_D} = \frac{(\bar{D} - \mu_D)/(\cancel{\sigma_D}/\sqrt{n})}{\sqrt{s_D^2/\cancel{\sigma_D^2}}} \xrightarrow{D} \frac{N(0,1)}{1} \approx N(0,1)$ ← not a function of σ_D^2
 and t_{n-1} tends to $N(0,1)$ as $n \rightarrow \infty$. ← cf.
- Q: What if the sample size n is small or population variance $= \infty$? ← cf.

Check
 Thm 12
 in LN,
 CH7, p.29