

Correlated Data

4/12/22

Paired Data - Binary response

- SPAT 216 → Paired t-test:
- Quantitative (normal) response
 - Measured in pairs
- Pre / Post
- Husband / Wife

2x2 Table:

Paired

		2 nd member of pair (Post)		
		Yes	No	
1 st member of pair (Pre)	Yes	n_{11}	n_{12}	n_{1+}
	No	n_{21}	n_{22}	n_{2+}
		n_{+1}	n_{+2}	n

own "Σ" per

Independent Samples

Y = go for walks more than 2x/week

		Y		
		0	1	
X	0			
	1			
				n

↳ "Y" on both rows & cols

(See 8.1)

McNemar's Test of Independence for Correlated Proportions:

$$H_0: P(Y_1 = \text{Yes}) = P(Y_2 = \text{Yes})$$

$$H_a: P(Y_1 = \text{Yes}) \neq P(Y_2 = \text{Yes})$$

one of $<$, $>$, \neq depending on research quest.

$$\pi_{ij} = P(\text{being in the } i^{\text{th}} \text{ row} \cdot j^{\text{th}} \text{ col})$$

$$\Rightarrow P(Y_1 = \text{Yes}) = \pi_{11} + \pi_{12} = \pi_{1+}$$

$$P(Y_2 = \text{Yes}) = \pi_{11} + \pi_{21} = \pi_{+1}$$

- Only interested in "discordant" pairs -
individuals in $(1,2)^{\text{th}}$ cell or $(2,1)^{\text{th}}$ cell

We say there is marginal homogeneity if

$$\pi_{12} = \pi_{21}$$

Why? $\pi_{1+} = \pi_{+1} \rightarrow$

$$\pi_{1+} - \pi_{+1} = (\cancel{\pi_{11}} + \pi_{12}) - (\cancel{\pi_{11}} + \pi_{21}) = \pi_{12} - \pi_{21}$$

Let $n^* = n_{12} + n_{21} \rightarrow$ total # of discordant pairs

Then under H_0 , $n_{12} \sim \text{Bin}(n^*, 0.5)$.
 \rightarrow Use to get p-value

or $n^* > 10$ (arbitrary rule of thumb),

$$Z = \frac{n_{12} - n^*(0.5)}{\sqrt{n^*(0.5)(1-0.5)}} = \frac{n_{12} - n_{21}}{\sqrt{n_{12} + n_{21}}} \sim N(0,1)$$

or $Z^2 \sim \chi^2(1) \leftarrow R$

Example: (Pediatrics, 2006) - Fire Safety
Researchers compared children's reactions to two kinds of alarms:

① Conventional smoke alarm

② Recording of mother's voice

Says the child's name & urges them to wake up.

\rightarrow each child tested by each alarm \rightarrow 2 measurements per child

		Mother's Voice		
		Awoke	Did Not Awoke	
Conventional Alarm	Awoke	14	0	14
	Did not awake	9	1	10
		23	1	24

(Note: In the original image, the cell '0' is circled in pink and the cell '9' is circled in red. A red arrow points to the '0' with the text 'observed test stat').

$$H_0: \pi_{12} = \pi_{21} \quad \left(P(\text{Awoke} | \text{Alarm 1}) = P(\text{Awoke} | \text{Alarm 2}) \right)$$

$$H_a: \pi_{12} \neq \pi_{21}$$

$$p\text{-value} = P(\bar{X} = 0 \text{ or } \bar{X} = 9 \mid \bar{X} \sim \text{Bin}(9, 0.5))$$

Let $\bar{X} = n_{12}$
(prior to data collection)

$$P: \text{sum}(d\text{binom}(c(0,9), 9, 0.5))$$

$$= 0.0039063$$

\Rightarrow Strong evidence that the probability of waking differs between the two alarm types.

Approximate p-value:

$$Z = \frac{n_{12} - n_{21}}{\sqrt{n_{12} + n_{21}}} = \frac{0 - 9}{\sqrt{9}} = -3$$

$$P: 2 \times \text{pnorm}(-3) = 0.0026998$$

$$\text{or } \text{pchisq}((-3)^2, 1, \text{lower.tail} = \text{FALSE})$$

		Mother's Voice		
		Awoke	Did Not Awake	
Conventional Alarm	Awoke	14	0	14
	Did not awake	9	1	10
		23	1	24

observed test stat

How would the data be organized in a 2x2 table if they were incorrectly treated as independent measurements.

ID	Alarm Type	Awoke or Not
1	Conv.	Yes
1	Mother	Yes
2	.	.
2	.	.
:	.	.
:	.	.

		Awoke or Not		
		Yes	No	
Alarm Type	Conv			
	Mother			48

(Note: A large red 'X' is drawn over the 2x2 table above.)

Asymptotic CI for $\pi_{1+} - \pi_{+1}$: (dependent data)

$$SE(\hat{\pi}_{1+} - \hat{\pi}_{+1}) = \frac{1}{n} \sqrt{(n_{12} + n_{21}) - \frac{(n_{12} - n_{21})^2}{n}}$$

$$CI: \hat{\pi}_{1+} - \hat{\pi}_{+1} \pm z^* \cdot SE(\hat{\pi}_{1+} - \hat{\pi}_{+1})$$

↑
from $N(0,1)$

Asthma Example \rightarrow 95% CI (0.01891, 0.03709)

We are 95% confident that the true difference in proportions of 13-year-olds having asthma \rightarrow 20-year-olds having asthma (13-20) is between (0.019, 0.037), among children similar to those in the sample.

We are 95% confident that the risk of asthma at age 13 is between 0.019 \times 0.037 higher than the risk of asthma at age 20

Notes on "correct = FALSE" argument in R.

- When using a normal approximation (continuous) on discrete data, we often employ a "continuity correction" -

$$\text{McNemar: } z_c^2 = \frac{(n_{12} - n_{21} - 0.5)^2}{n_{12} + n_{21}}$$

Tails

$$\sum_{H_0}^0 \chi_1^2$$

$$\text{Chi-squared: } \chi_c^2 = \sum \frac{(\text{obs} - \text{exp} - 0.5)^2}{\text{exp}}$$

More Generally - Modeling Framework

Option 1: Population-averaged models - Marginal models

$$P(Y_1=1) = \alpha + \delta$$

$$P(Y_2=1) = \alpha$$

$$\delta = P(Y_1=1) - P(Y_2=1)$$

$$P(Y_t=1) = \alpha + \delta X_t$$

$$X_t = \begin{cases} 1 & t=1 \\ 0 & t=2 \end{cases}$$

or another common model:

$$\text{logit}(P(Y_t=1)) = \log\left(\frac{P(Y_t=1)}{1-P(Y_t=1)}\right) = \alpha + \delta X_t$$

→ But data are not independent. "Clustered" data.

- Need different method for fitting these models → generalized estimating equations (GEE)

→ "Naive SEs" → treating data as independent
* "Robust SEs" → account for correlation

