What to do today (Feb 16, 2023)?

Part I. Introduction Part II. Epidemiologic Concepts and Designs Part III. Clinical Trials

Part IV. Modern Biostatistical Approaches

Part IV.1 Incomplete Data Analysis

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Part IV.3 Selected Widely-Used Algorithms

Mismeasurement of exposure status or level is "present to at least some degree in nearly every epidemiologic study, since nearly every means of ascertaining the presence or level of exposure is imperfect" – Koepsell and Weiss (2003)

- Measure refers broadly to any way of capturing data on a certain characteristic of study subjects.
- Measurement error is the discrepancy between the true value and the measured value.
- The scale of measurement is usually categorized into
 - continuous: e.g. body weight; any positive real number
 - categorical: ordinal vs nominal; e.g. disease serverity mild, moderate, severe vs gender – male, female

Misclassification; Fine to Coarse Measurement Scales

Assessing Measurement Error

- Reliability. A good measurement should yield the same value if applied repeatdly under circumstances in which the underlying characteristic is believed to remain the same.
 - e.g. for binary measures and 2 × 2 table of outcomes, concordance [percent agreement]: p_O = n₁₁/n₊₊ + n₂₂/n₊₊
 - e.g. for binary measures, Kappa: κ = p₀-p_e/1-p_e with p_e = (n₁₊/n₊₊)(n₊₊/n₊₊) + (n₂₊/n₊₊)(n₊₊/n₊₊), expected overlap by chance
 e.g. for continuous measures, intraclass correlation coefficient (reliability ratio): λ = σ²/σ²_x+σ²_y ≤ 1

 Validity A good measurement method should yield the correct value. [Being consistent is not good enough if the results are consistently wrong.]

A gold standard (a criterion measure) is required to evaluate the validity of a measure.

- Sensitivity and specificity: 2 × 2 table of outcomes with a diagnosis test and the condition presence Sensitivity=P(T₊|C₊), estimated by n₁₁/n₊₁ Specificity=P(T₋|C₋), estimated by n₂₂/n₊₂
- when a test yields an ordinal or continuous scale, often is to select a cutoff value => receiver operating characteristic (ROC) curve: (1-specificity, sensitivity) at different cutoff values

uninformative test; good test; perfect test

Consequences of Measurement Error

- with Continuous Variables
 - when the variable is the response: if the errors sum up to zero? if the errors don't sum up to zero?
 - when the variable is explanatory: if the errors sum up to zero? [Part IV.1.4] if the errors don't sum up to zero?
- with Categorical Variables (misclassification)
 - non-differential (non-selective) a form of random measurement errors?
 - differential bias to a particular direction?

"Nondifferential misclassification of exposure is ubiquitous in epidemiology, and usually leads to an attenuation of the estimated size of a true association between exposure and disease." (Thomas, 1995)

Example. In a case-control study

A. When t	пе ехро	sule was	measured perfectly
Exposure	case	control	Odds Ratio
yes	150	75	$\frac{150}{150} \div \frac{75}{225}$
no	150	225	= 3.0
Total	300	300	-

A. When the exposure was measured perfectly

Example. (cont'd)

	1 /0	c .					· C' I
B. When	1/3	ot ex	nosured	subjects	were	miscla	ssitied
D. Winch	1/5	OI CA	posuicu	Jubjects	were	mocia	JJIIICu

D. When I		Surea Subjet	
Exposure	case	control	Odds Ratio
yes	150-50	75-25	$\frac{100}{200} \cdot \frac{50}{250}$
no	150 + 50	225 + 25	= 2.5
Total	300	300	

C. In addition to B., 20% of non-exposured subjects were misclassified

Exposure	case	control	Odds Ratio
yes	150-50+30	75-25+45	$\frac{130}{170} \div \frac{95}{205}$
no	150 + 50 - 30	225+25-45	= 1.65
Total	300	300	-

What is confounding?

"Confounding occurs in epidemiologic research when the measured association between an exposure and disease occurrence is distorted by an imbalance between exposed and non-exposed persons with regard to one or more other risk factors for the disease."

- Koepsell and Weiss (2003)

Example. Crude Death Rate (per 100,000 person-years):

 $\frac{total \ deaths \ in \ a \ year}{average \ popluation \ in \ the \ year} \times 10^5$

 U.S. Global Health Policy: (http://www.globalhealthfacts.org/data/topic/map.aspx?ind=90) Crude Death Rate (per 100,000 people) in 2012: Canada 8.09; Mexico 4.90

Mexican age specific moratlity rates are greater: The World Bank (http://data.worldbank.org/indicator/SH.DYN.MORT) age 5 or under group: Canada 6; Mexico 16 Why?

-						
	(Community A		Community B		
	No. of	Mid-Year		No. of	Mid-Year	
Age	Deaths	Population	$Rate^a$	Deaths	Population	$Rate^a$
young	1	1000	1	10	5000	2
middle	15	3000	5	40	4000	10
old	50	5000	10	20	1000	20
Total	66	9000	7.3	70	10,000	7.0

Example. Mortality Rates in Two Hypothetical Communities

^aDeaths per 1000 person-year

- Crude Death Rates (1000 per-year): A, 7.3; B, 7.0
- Mortality Rates in A and B both sharply increase with increasing age
- Difference in the age distributions on average: people in A older

A has higher proportion of older people and is "penalized" in comparison to B: the Simpson's Paradox

Methods of Accounting for Confounding Variables:

in the Study Design:

- random assignment
- matching select matched pairs (sets) from each age group in Mexico and Canada

restriction - compare death rate within a specific age group

as Part of Data analysis:

 stratification - obtain separate comparisons of death in each selected age groups using age-specific mortality rates

covariate adjustment

Advantage vs disadvantage for each?

Standardization: to calculate what would have been the overall mortality rates in A and B if they had the same age composition (i.e. by using a common set of weights).

- Step 1. Pick a reference population to construct weights Choice of a Standard Population:
 - regional comparisons may use the combined population of a specified date as the standard
 - the non-exposed group
- Step 2. Calculate weighted average using age-specific rates in each population and the selected weights.

The common confounding factor distn is taken from the standard population; hence, the term of "standardization".

Example. Mortality Rates in Two Hypothetical Communities (cont'd)

Step 1. Select the combined mid-year population of Community A and B to construct the reference population:

Age	Standard Weig	Standard Weights				
young		= 0.316				
middle	(3000+4000)/19,000	=0.368				
old	(5000+1000)/19,000	=0.316				
Total		1.000				

Step 2. Calculate weighted average

	Community A			C	ommunity	иB
Age	rate	weight		rate	weight	
young	$1 \times$.316	=.316	2 ×	.316	=.632
middle	$5 \times$.368	=1.84	10 $ imes$.368	=3.68
old	10 \times	.316	=3.16	$20 \times$.316	=6.32
		5.3 ^a			10.6 ^a	
				-		

^aAge standardized mortality rates in Community A and B

Direct vs Indirect Standardization

- Direct Standardization: all disease rates from strata are (weighted) averaged using the distribution of the standard population for the weights
 - It gives the crude rate would have been if the study population(s) had the same distribution as the standard population.

Other adjusted measures e.g. $\hat{\theta}_{XY,MH} = \frac{\sum_k N_{11k} N_{22k} / N_{++k}}{\sum_k N_{12k} N_{21k} / N_{++k}}$ [adjusted OR]

It may be inefficient when there are few events per stratum

- Indirect Standardization:
 - "Multivariate regression analysis" (Multiple regression?)
 e.g. multiple logistic regression analysis [adjusted log-OR]: an additional covariate to adjust for the effect of a confounder
 - Propensity scores to control multiple potential confounders simultaneously by using a propensity score:
- First modeling the exposure variable as a function of the potential confounders by logistic regression or a related method:

to calculate an expected probability ("propensity") of exposure for each study subject

Then examining the exposure-outcome association while controlling for the propensity score by stratification, matching, or covariate adjustment

Stratification:

to separate data into several subgroups (e.g. by age and sex)

1st step in standardization

stratified analysis: rationale for reporting it vs a combined result?

"conditioning"

Conditional vs Marginal Associations

X-Y conditional odds ratios: [describe conditional X-Y association] For Z = k, k = 1, ..., K,

$$\theta_{XY(k)} = \frac{\pi_{11k}\pi_{22k}}{\pi_{12k}\pi_{21k}} = \frac{\mu_{11k}\mu_{22k}}{\mu_{12k}\mu_{21k}}$$

If $\theta_{XY(k)} \equiv \text{constant}$, \implies "homogeneous" conditional X-Y association

X-Y marginal odds ratios: [describe marginal X-Y association]

$$\theta_{XY} = \frac{\pi_{11+}\pi_{22+}}{\pi_{12+}\pi_{21+}} = \frac{\mu_{11+}\mu_{22+}}{\mu_{12+}\mu_{21+}}$$

- Homogeneous conditional association
 If θ_{XY(k)} = c for all k, not necessarily θ_{XY} = c
 e.g. the Simpson's Paradox
- Marginal vs conditional independence

•
$$X \perp Y | Z \leftrightarrow (iff) \ \theta_{XY(k)} = 1$$
 for all k

$$\blacktriangleright X \bot Y \leftrightarrow \text{(iff)} \ \theta_{XY} = 1$$

$$\triangleright X \bot Y | Z \not\leftrightarrow X \bot Y$$

Cohran-Mantel-Haenszel Test. with a 2 × 2 × K table, to test X⊥Y|Z − H₀ : "θ_{XY(k)} = 1 for all k = 1,..., K" vs H₁ : otherwise

CMH-test works well if conditional X-Y associations are similar

- ► Mantel-Haenszel Estimator. with a 2 × 2 × K table, when $\theta_{XY(1)} = \ldots = \theta_{XY(K)}$, to estimate the common conditional odds ratio: $\hat{\theta}_{XY,MH} = \frac{\sum_k N_{11k} N_{22k} / N_{1+k}}{\sum_k N_{12k} N_{21k} / N_{1+k}} \neq \frac{N_{11+} N_{22+}}{N_{12+} N_{21+}}$
- Breslow-Day Test. with a 2 × 2 × K table, to test for homogeneity of conditional odds ratios H₀: θ_{XY(1)} = ... = θ_{XY(K)} vs H₁: otherwise

Confounding vs Mediating Variables

- Mediators are also known as intervening or intermediate variables.
- Confounders are associated with but not caused by exposure; adjusting for variables on the causal pathway biases estimated odds ratios towards one (Leon, 1993).
- e.g. Birth weight is on the causal pathway between maternal smoking and infant mortality:

Maternal		Birth		Infant
Smoking	\Rightarrow	Weight	\Rightarrow	Mortality
(Exposure)		(Mediator)		(Outcome)

The odds ratio for infant mortality comparing smokers to non-smokers was:

- 1.3 (95% CI (1.2,1.4)), after adjusting for marital status, education, maternal age and parity;
- ▶ 1.0 (95% CI (0.9,1.1)), after further adjustment for infant birth weight!

Residual Confounding

Our ability to obtain unconfounded estimates for the effect of exposure in observational studies is limited by residual confounding due to:

- unknown confounding variables,
- known confounders are not measured,
- random measurement error (non-differential misclassification) of confounders biasing adjusted estimates of the exposure-disease association towards estimates of the unadjusted association.

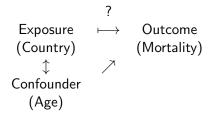
For example,

- Mothers who smoke while pregnant tend to have smaller babies.
- Male babies tend to be bigger than female babies.
- To what extent could the observed association between maternal smoking and infant birth weight be confounded by infant gender?

When is confounding present?

classical criteria

A variable is a confounder if it is associated with exposure and causally related to the outcome:



the question mark ? about the association of Country and Mortality

collapsibility criterion

Confounding is present when there is a substantive difference between the crude and adjusted odds ratios.

A common application of the collapsibility criterion concern for the effects of confounding occur when the crude and adjusted estimates of excess risk differ by at least 10%.

How to Use the Criteria for Confounding?

The classical criteria may be used when designing a study to:
 (i) develop a conceptual framework and (ii) identify potential confounding variables.

The classical criteria may also prove useful in identifying the source of confounding.

The collapsibility criteria is most useful when deciding how best to describe study results.

What to study next?

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Part IV. Analytic Epidemiology

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IV.2.3 Confunding and Its Control

IV.2.4 Causation vs Association

Part IV.3 Selected Widely-Used Algorithms

IV.3.1 Bootstrap and Related

IV.3.2 EM Algorithm and Related