Counter matching

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COUNTER-MATCHING IN Cincinnati

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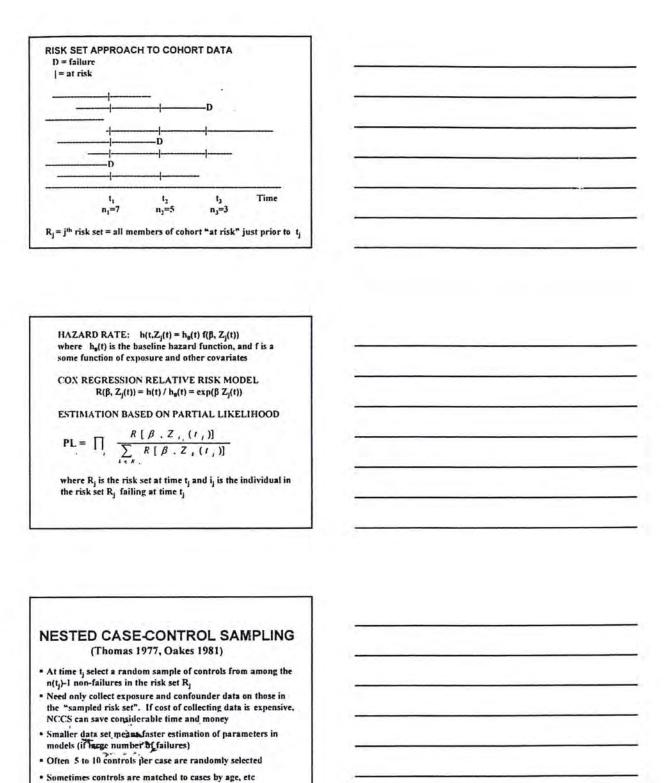
INCREASED PRECISION USING COUNTER-MATCHING IN NESTED CASECONTROL STUDIES, EPIDEMIOLOGY, 8(1997):238-242, COMMENTARY PP-227-229



MORTALITY/MORBIDITY STUDIES INVOLVING TIME UNTIL AN EVENT

- OUTCOME OF INTEREST (eg, DEATH FROM LUNG CANCER)
- EXPOSURE OF INTEREST (eg, CUMULATIVE EXPOSURE TO DUST)
- OFTEN THE EXPOSURE OF INTEREST IS EXPENSIVE TO MEASURE ON ALL SUBJECTS

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NESTED CASE-CONTROL SAMPLING

- . Estimation is done using the usual Partial Likelihood
- Usually one forms the risk sets, selects the controls, and uses PROC PHREG with STRATA=risk set, and defining the case's failure time to occur before the time assigned to the controls
- * Efficiency is known to be (m-1)/m when the exposure response coefficient β is 0, where efficiency is defined as the ratio of the variance of the estimator of β using the full cohort to the variance of the estimator of β using Nested Case-Control Sampling

$$EFF = \sigma_{FULL}^{2} / \sigma_{NCCS}^{2}$$

COUNTER-MATCHING IN NESTED CASE-CONTROL(CM) (Langholz & Clayton 1994, Langholz & Borgan, etc.)

GOAL: FURTHER IMPROVE EFFICIENCY OF NCCS

SUPPOSE for all individuals in the cohort one knows a covariate X(t) (surrogate) that is correlated with the true exposure Z(t)

- Suppose each risk set R_j is divided into k strata (based on X(t)) containing n₁, n₂, n₃,..., n_k subjects. One then randomly selects m₁, m₂, m₃,..., m_k subjects from each strata in such a way that if the case is in strata i one selects only m_i-1 controls from strata i.
- 1-1 counter-matching requires dividing each risk set into 2 strata and then picking 1 control from the strata not containing the case.
- "Maximize the heterogeneity of exposure in the sampled risk sets
- Since counter-matching does not involve random sampling, one must compensate by introducing weights into the partial likelihood equation: For each risk set R₁ let:

weight =
$$W_i(t_i) = n_i / m_i$$

where \mathbf{n}_i is the number of subjects in strata \mathbf{i}_i and \mathbf{m}_l is the number of controls selected from strata \mathbf{i}_i .

Estimates are then based on the modified partial likelihood.

PL
$$\prod_{i=1}^{k} \frac{R[\beta, Z_i(t_i)] \cdot W_i(t_i)}{\sum_{i=k}^{k} R[\beta, Z_i(t_i)] \cdot W_i(t_i)}$$

 Weights can be incorporated into the estimation process using the offset function in PROC PHREG

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. If the surrogate X(t) is a 0/1 variable, then 1-1 countermatching requires picking a control with X(t)=0 if the case has X(t)=1, and visa versa. . This maximizes the number of "discordant" pairs with respect to the surrogate X(t) and possibly with respect to Z(t). . If the surrogate is a 0/1 variable, then 3-1 counter-matching might consist picking 2 controls with X(t)=0 and 1 control with X(t)=1 if the case has X(t)=1, etc ($m_1=2$, $m_2=2$), etc . If the surrogate X(t) is a continuous variable, then one might create 4 strata for each risk set R1, based on cutpoints of X(t) For 3-1 counter-matching one would pick one control from each strata except the strata of the case. For 7-1 counter-matching one might pick 1 control from the case's strata, and 2 from each of the other three strata (m1= m2= m3= m4=2). Of course, there are many other possible choices of m1, m2, m3, m4 summing to 8 . The question is then "How to form the strata in each risk set?" . The natural method would be to use "quartiles" (or percentiles) There are several ways to pick the percentiles. For illustration purposes suppose one does a 3-1 counter-matching with 4 strata. METHOD 1: Pick quartiles of the covariate X(t) within each risk set. If equal numbers are sampled from each quartile, then all weights are equal, and hence cancel out of the partial likelihood. METHOD 2: Pick quartiles of the covariate X(t) for the cases. Then weights will vary, since the size of the strata will vary within each risk set and between risk sets. In our study METHOD 2 proved to be more efficient.

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ppose X(t) and 2 (X=1 Z=1) spec gree of associationsure	cificity = $P(X=0)$	Z=0), 1.e.,		_		
fine the asympt	totic relative ef	ficiency (A	RE):	-		
ARE	$E = \sigma_{NCCS}^2 / \sigma_{CN}$	2		0.2		
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nder Ha: B=0	0, Langholz	and Clay	ton showed:	1.2		
				100		
ARE		CIFICITY	0.0			
ENSITIVITY 0.4	0.80	0.9	0.8 0.88			
0.5	1.00(1)	1.00	1.00			
0.7	1.40(2)	1.32	1.24	-		
0.8	1.60(4)	1.48	1.36			
0.9	1.80(19)	1.64	1.48	1 1		
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Langholz & Cl and P(Z=1)=.1		wed that if	exp(p)=4			
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ARE	SPEC	IFICITY		100		
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ARE	SPECIFICITY		
SENSITIVITY	1.0	0.9	0.8
0.4	1.79	1.38	1.14
0.5	2.17	1.65	1.33
0.7	2.89	2.17	1.69
0.8	3.23	2,41	1.86
0.9	3.54	2.64	2.02

 Langholz & Clayton also showed that relative efficiency increased with rarity of exposure.

In general the efficiency of CM versus NCCS depends on:	
(Borgan & Olsen 1999)	
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• sensitivity and specificity	
• fraction exposed	
baseline hazard	
* relative risk coefficient β	
• censoring distribution	
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Most general results have been developed for dichotomous (0/1) exposures Z(t) and their dichotomous (0/1) surrogates X(t).	
exposures 2(1) and men denotomous (0/1) surrogates refs.	
We were interested in continuous exposures and surrogates.	-
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CRYSTALLINE SILICA EXPOSURE AND SILICOSIS	
IN GOLD MINERS	
 cohort of 3,300 gold miners, 1940-1965 	
170 cases of silicosis	
 cumulative silica exposure estimated via repeated dust 	,
sampling	
a case control study would avoid much cost of dust	
collection for whole cohort	
a surrogate for cumulative exposure, duration employed	
is known AND highly correlated with cumulative silica exposure	
Calmanic	
Steenland & Brown, Silicosis among gold miners, Amer.	,
J. Public Health, 85(1995)1372-1377	
GOLD MINERS COHORT	-
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We wanted to see the effect of various sampling strategies: full cohort, nested case-control sampling (m=3,10,20,100),	-
and counter-matching (m=3).	
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We performed 50 sample selections for each method	
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We performed the analysis on the full cohort, which showed a very strong dose-response, and also on a	
reconstructed data set with a reduced dose response.	
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RESULTS : Average estimates of the dose response coefficient β and its standard error:

DESIGN	ORIGINAL	REDUCED
FULL	1.56(.110)	0.844(.078)
NCCS(100)	1.56(.112)	0.843(.078)
NCCS(20)	1.54(.119)	0.845(.081)
NCCS(10)	1.54(.127)	0.840(.084)
NCCS(3)	1.46(.155)	0.876(.103)
CCM(3) M1	1.53(.157)	0.842(.093)
CCM(3) M2	1.61(.125)	0.865(.085)

^{*} averages based on 50 sample selections

RESULTS : Estimated relative efficiency = $100^{+}\sigma_{\text{FULL}}^{2}/\sigma_{\text{CM}}^{2}$

DESIGN	ORIGINAL	REDUCED
NCCS(100)	96.4%	98.8%
NCCS(20)	85.4%	91.9%
NCCS(10)	73.9%	85.7%
NCCS(3)	50.0%	59.1%
CCM(3) M1	49.0%	71.6%
CCM(3) M2	77.7%	83.1%

- averages based on 50 sample selections
- counter-matching m=3 equivalent to matching m=10 to 15
- NCCS and CM are more efficient for reduced dose-response effect

CONCLUSIONS:

- Method 2 (strata formed based on cases' exposure) is superior to Method 1 (strata formed based on all members of risk set)
- * Counter-matching is superior to NCCS
- Counter-matching with 3 controls is about the same as NCCS with 10-15 controls, i.e., countermatching achieves same efficiency as NCCS using fewer selected controls (LESS EXPENSIVE)
- Counter-matching is even more efficient for reduced dose response situation

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GENERAL CONSIDERATIONS

- Counter-matching requires knowing some information on all subjects in the cohort, before randomly sampling
- if the information (eg, duration) is correlated with exposure then one should use it to COUNTER-MATCH
- on the other hand, if the information is a confounder, then one should use it to MATCH
- counter-matching on a confounder will result in a loss of efficiency for estimating exposure
- matching on a variable related to exposure will result in being unable to estimate the exposure effect

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