Chapter 153

Tests for Two Proportions in a Cluster-Randomized Design with Clustering in Only One Arm

Introduction

Cluster-randomized designs are those in which whole clusters of subjects (classes, hospitals, communities, etc.) are put into a treatment group or a control group. In this case, the proportions of the two arms (groups), where the first arm is made up of K_i clusters of M_{ij} individuals each and the second arm is made up of individuals, are to be tested using a modified *z* test.

In the trials analyzed by this procedure, subjects are treated together (nested) within clusters in one arm, but not in the other. In the other arm, subjects are treated individually. For example, one arm might receive individual intervention (such as medicine) while the other receives this medicine plus some type of group therapy session. The fact that they attend this therapy session implies that they are "clustered". Thus the clustering occurs in only one arm.

Technical Details

Our formulation comes from Moerbeek and Wong (2008). They combine two mixed models: one for the arm made up of individuals (group 2) and a second, more complicated model for the arm (group 1) which accounts for the clustering. Let y_i be a binary variable that is one of a certain outcome is observed and zero otherwise. The treatment effect is measured by $D = P_1 - P_2$. This can be estimated by

$$\widehat{D} = \overline{y}_1 - \overline{y}_2$$

where \bar{y}_1 and \bar{y}_2 are the estimators of the probabilities of the outcome $y_i = 1$, P_1 and P_2 .

The treatment effect may be tested for statistical significance using

$$\widehat{D}/\sqrt{\operatorname{var}(\widehat{D})}$$

The variance of this estimator is

$$\operatorname{var}(\widehat{D}) = P_2(1 - P_2) \left(\tau \frac{(\overline{m} - 1)\rho + 1}{\overline{m}k} + \frac{1}{N_2} \right)$$

where \overline{m} is the average cluster size, N_2 is the number of subjects in the non-clustered arm, ρ is intracluster correlation coefficient, and τ is the ratio of the variances of the outcome when $\rho = 0$,

$$\tau = \frac{P_1(1-P_1)}{P_2(1-P_2)}.$$

Assume that $D = P_1 - P_2$ is to be tested using a z-test. The statistical hypotheses are $H_0: D = 0$ vs. $H_a: D \neq 0$. The test statistic

$$t = \frac{\bar{Y}_1 - \bar{Y}_2}{\sqrt{\operatorname{var}(\widehat{D})}}$$

has an approximate normal distribution for a *subject-level* analysis.

Using the above, a large sample formula for computing the power of a two-sided test of the significance of the treatment effect at significance level α can be derived from

$$\operatorname{var}(\widehat{D}) = \left(\frac{D}{z_{1-\frac{\alpha}{2}} + z_{Power}}\right)^2$$

The power of a one-sided test can be calculated similarly.

Example 1 – Calculating Sample Size

Suppose that a cluster randomized study is to be conducted in which one arm (group 2) will receive an individual medical intervention while the other arm (group 1) receives this medicine plus a special group therapy session conducted by a trained therapist. These therapy sessions will be treated as clusters. Here, group 2 (the non-clustered subjects) is assigned to the 'control group' and group 1 (the clustered subjects) is assigned to the treatment group. The researchers want to explore what happens as *R* is varied from 1 to 2.

The parameter values are set as follows: *alpha* = 0.05, *power* = 0.9, R = 1.0 1.5 2.0, *P1* = 0.25, *P2* = 0.4, ρ = 0.01, and *M1* = 10. Sample size is to be calculated for a two-sided test.

Setup

If the procedure window is not already open, use the PASS Home window to open it. The parameters for this example are listed below and are stored in the **Example 1** settings file. To load these settings to the procedure window, click **Open Example Settings File** in the Help Center or File menu.

Design Tab	
Solve For	Sample Size
Alternative Hypothesis	Two-Sided (H1: P1 - P2 ≠ 0)
Power	0.90
Alpha	0.05
Group Allocation	Enter R = (K1 × M1) / N2, solve for K1 and N2
R (Allocation Ratio)	1 1.5 2
M1 (Average Cluster Size)	10
Input Type	Proportions
P1 (Group 1 Proportion H1)	0.25
P2 (Group 2 Proportion)	0.4
ρ (Intracluster Correlation, ICC)	0.01

Output

Click the Calculate button to perform the calculations and generate the following output.

Numeric Reports

Numeric	Results												
Solve For Clustering Group Allo Hypothese	: g: ocation: es:	Samp Group Enter H0: D	le Size 1 has Clustering R = (K1 × M1) / N = 0 vs. H1: D	l, Group 2 ha V2, solve for I ≠ 0	s No Clusteri <1 and N2	ng							_
	G	roup 1	Clusters	Comm	e Cine (Cubi								
	Numbe	er of	Average	Sampi			Allocation	Propo	ortions	-			
Power	Clus	K1	Cluster Size M1	Group 1 N1	Group 2 N2	l otal N	Ratio	P1	P2	Difference D1	ρ	Alpha	
0.90326 0.90665 0.90027		21 27 32	10 10 10	210 270 320	210 180 160	420 450 480	1.0 1.5 2.0	0.25 0.25 0.25	0.4 0.4 0.4	-0.15 -0.15 -0.15	0.01 0.01 0.01	0.05 0.05 0.05	
Power K1 M1 N1, N2, a R P1 P2 D1 ρ Alpha	and N	The p The p The p The p The p The p The p The p	probability of re- number of clust average cluster number of subj target value of R = (K1 × M1) / value of the res value of the res difference in the intracluster com- probability of re-	ejecting a fa ters in group size (numb ects in grou the allocatic 2 N2. The ac sponse prop e response relation (ICC ejecting a tru	lse null hyp o 1. Der of subje- ps 1 and 2, n ratio of th ctual ratio va ortion of gro ortion of gro proportions C). The corr ue null hypo	othesis v cts) of th and the ne numb alue may oup 1 as oup 2 as at which relation b thesis.	when the alter e clusters in <u>c</u> ir total. Note the er of subjects or not be exact sumed by the sumed by bot in the power is between a pair	native h group 1. hat N1 = in group ly R bec alterna h H0 ar calcula r of subj	ypothes = K1 × N > 1 and ause N tive hyp Id H1. ted. D1 ects wit	sis is true. M1. number of su 1 and N2 are bothesis, H1. = P1 - P2. hin a cluster.	bjects in integer	n group s.	

Summary Statements

A parallel, two-group design with cluster-randomized subjects in Group 1 only (and no clustering in Group 2) will be used to test whether the Group 1 proportion (P1) is different from the Group 2 proportion (P2) (H0: D = 0 versus H1: $D \neq 0$, D = P1 - P2). The comparison will be made using a two-sided mixed model Z-test with a Type I error rate (α) of 0.05. The intracluster correlation coefficient for Group 1 is assumed to be 0.01. To detect a difference (P1 - P2) of -0.15 (P1 = 0.25 and P2 = 0.4), with 10 subjects per cluster in Group 1 and 210 subjects in Group 2, with 90% power, the number of needed clusters in Group 1 is 21 (totaling 210 subjects in Group 1).

References

Moerbeek, M. and Wong, W.K. 2008. 'Sample size formulae for trials comparing group and individual treatments in a multilevel model.' Statistics in Medicine, Vol. 27, pages 2850-2864.

Donner, A. and Klar, N. 2000. Design and Analysis of Cluster Randomization Trials in Health Research. Arnold. London.

This report shows the results for each of the scenarios.

Plots Section



This plot shows the number of clusters versus the sample size allocation ratios.

Example 2 – Validation using Moerbeek and Wong (2008)

Moerbeek and Wong (2008) pages 2858 and 2859 provide an example which we will used to validate this procedure. When alpha is 0.05, *P2* is 0.243, P1 is 0.397, ρ is 0.05, N2 is 146, M1 is 8, and K1 is 23, they calculate a power of approximately 0.80.

Setup

If the procedure window is not already open, use the PASS Home window to open it. The parameters for this example are listed below and are stored in the **Example 2** settings file. To load these settings to the procedure window, click **Open Example Settings File** in the Help Center or File menu.

Design Tab	
Solve For	Power
Alternative Hypothesis	Two-Sided (H1: P1 - P2 ≠ 0)
Alpha	0.05
Group Allocation	Enter K1 and N2 individually
K1 (Number of Clusters)	23
M1 (Average Cluster Size)	8
N2 (Number of Subjects)	146
Input Type	Proportions
P1 (Group 1 Proportion H1)	0.397
P2 (Group 2 Proportion)	0.243
ρ (Intracluster Correlation, ICC)	0.05

Output

Click the Calculate button to perform the calculations and generate the following output.

Number of Average ————————————————————————————————————	Solve For: Clustering Group Allo Hypothese	: Powe p: Grou pcation: Ente es: H0: [r o 1 has Clustering, 0 K1 and N2 individu 0 = 0 vs. H1: D ≠ 0	Group 2 has N ally)	o Clustering						
		Grou	1 Clusters	Samp	le Size (Subje	ects)					
	Power	Grou Number of Clusters K1	0 1 Clusters Average Cluster Size M1	Samp Group 1 N1	le Size (Subje Group 2 N2	ects) Total N	Propo P1	P2	Difference D1	ICC P	Alpha

PASS calculates a power of approximately 0.80. Thus the procedure is validated.