Chapter 608

Multi-Arm Non-Inferiority Tests for the Ratio of Treatment and Control Proportions in a Cluster-Randomized Design

Introduction

This module computes power and sample size for multi-arm non-inferiority tests of the ratio of treatment and control proportions when binary data are gathered from a cluster-randomized design. The formulas are based on results in Donner and Klar (2000) and Machin, Campbell, Tan, and Tan (2018).

A *cluster (group) randomized design* is one in which whole units, or clusters, of subjects are randomized to the groups rather than the individual subjects in those clusters. The conclusions of the study concern individual subjects rather than the clusters. Examples of clusters are families, school classes, neighborhoods, hospitals, and doctor's practices.

Cluster-randomized designs are often adopted when there is a high risk of contamination if cluster members were randomized individually. For example, it may be difficult for doctors to use two treatment methods in their practice. The price of randomizing by clusters is a loss of efficiency--the number of subjects needed to obtain a certain level of precision in a cluster-randomized trial is usually much larger than the number needed when the subjects are randomized individually. Hence, standard methods of sample size estimation cannot be used.

In this multi-arm design, there are *G* treatment groups and one control group. A proportion is measured in each group. A total of *G* hypothesis tests are anticipated each comparing a treatment group with the common control group using a *z*-test of the ratio between two proportions.

The Bonferroni adjustment of the type I error rate may be optionally made because several comparisons are being tested using the same data. Making a multiplicity adjustment is usually recommended, but not always. In fact, Saville (1990) advocates not applying it and Machin, Campbell, Tan, and Tan (2018) include omitting it as a possibility.

Example

Suppose that the current treatment for a disease works 70% of the time. Unfortunately, this treatment is expensive and occasionally exhibits serious side-effects. Two promising new treatments have been developed and are now ready to be tested. Two non-inferiority hypotheses need to be tested in this study: whether each new treatment is as good as the current treatment. At least three groups are needed to complete this study of the two new treatments.

Because of the many benefits of the new treatment, clinicians are willing to adopt a new treatment even if it is slightly less effective than the current treatment. They must determine, however, how much less effective

the new treatment can be and still be adopted. Should it be adopted if 69% respond? 68%? 65%? 60%? There is a percentage below 70% at which the ratio between the two treatments is no longer considered ignorable. After thoughtful discussion with several clinicians, it was decided that if a response ratio is less than or equal to 0.80, the new treatment will be adopted.

Multiple Treatments Versus a Single Control

Whether you want to test several doses of a single treatment or several types of treatments, good research practice requires that each treatment be compared with a control. For example, a popular three-arm design consists of three groups: control, treatment A, and treatment B. Two tests are run: treatment A versus control and treatment B versus the same control. This design avoids having to obtain a second control group for treatment B. Besides the obvious efficiency in subjects, it may be easier to recruit subjects if their chances of receiving the new treatment are better than 50-50.

Technical Details

Our formulation for cluster-randomized designs comes from Donner and Klar (2000). Suppose you have *G* treatment groups with response probabilities P_i that have samples of size N_i and one control group with response probability P_c that has a sample of size N_c . The total sample size is $N = N_1 + N_2 + ... + N_G + N_c$.

The statistical hypotheses to be tested are

$$H_0: P_i \le P_C(R0)$$
 vs. $H_1: P_i > P_C(R0)$

where *R*0 is the *non-inferiority ratio*.

Cluster-Randomized Designs

Denote a binary (0, 1) observation by Y_{ikj} where *i* is the group, $k = 1, 2, ..., K_i$ is a cluster within group *i*, and $j = 1, 2, ..., M_i$ is an item (often a subject) in cluster *k* of group *i*. The results that follow assume an equal number of items per cluster per group. When the number of items from cluster to cluster are about the same, the power and sample size values should be fairly accurate. In these cases, the average number of items per cluster can be used.

The statistical hypothesis that is tested concerns the ratio between a treatment group proportion and the control group proportion: P_i and P_c . With a simple modification, the large-sample sample size formulas that are listed in the module for testing two proportions can be used here.

When the items are randomly assigned to one of the G + 1 groups, the variance of the sample proportion is

$$\sigma_{S,i}^2 = \frac{P_i(1-P_i)}{N_i}$$

PASS Sample Size Software

When the randomization is by clusters of items, the variance of the sample proportion is

$$\sigma_{C,i}^{2} = \frac{P_{i}(1 - P_{i})(1 + (M_{i} - 1)\rho)}{K_{i}M_{i}}$$
$$= \sigma_{S,i}^{2}[1 + (M_{i} - 1)\rho]$$
$$= F_{i,\rho}\sigma_{S,i}^{2}$$

The factor $[1 + (M_i - 1)\rho]$ is called the *inflation factor*. The Greek letter ρ is used to represent the *intracluster correlation coefficient (ICC)*. This correlation may be thought of as the simple correlation between any two subjects within the same cluster. If we stipulate that ρ is positive, it may also be interpreted as the proportion of total variability that is attributable to differences between clusters. This value is critical to the sample size calculation.

The asymptotic formulas that were used in comparing two proportions (see Chapter 200, "Tests for Two Proportions") may be used with cluster-randomized designs as well, as long as an adjustment is made for the inflation factor. The basic form of the z-test becomes

$$z = \frac{\left|\widehat{D}\right|}{\widehat{\sigma}_{\widehat{D}}(R0)}$$

where

$$\widehat{D} = \widehat{p}_i - \widehat{p}_C R 0$$

$$\hat{\sigma}_{\widehat{D}}(R0) = \sqrt{\frac{\widetilde{p}_i(1-\widetilde{p}_i)F_{i,\rho}}{N_i} + \frac{\widetilde{p}_C(1-\widetilde{p}_C)F_{C,\rho}}{N_C}}$$

The quantities \tilde{p}_i and \tilde{p}_c are the maximum likelihood estimates constrained by $\tilde{p}_i - \tilde{p}_c(R0) = 0$.

Test Statistic

This procedure assumes that Farrington and Manning (1990) likelihood score test statistic is used for the analysis of the data obtained from this study. This statistic tests whether the response ratio is less than or equal to a specified value *R*0. The regular MLE's, \hat{p}_i and \hat{p}_c , are used in the numerator of the score statistic while MLE's \hat{p}_i and \hat{p}_c , constrained so that $\tilde{p}_i/\tilde{p}_c = R0$, are used in the denominator. The significance level of the test statistic is based on the asymptotic normality of the score statistic.

The formula for computing the test statistic is

$$z_{FMD} = \frac{\hat{p}_i - \hat{p}_C(R0)}{\sqrt{\left(\frac{\tilde{p}_i \tilde{q}_i}{N_i / F_{i,\rho}} + \frac{\tilde{p}_C \tilde{q}_C}{N_C / F_{C,\rho}}\right)}}$$

where the estimates \tilde{p}_i and \tilde{p}_c are computed as in the corresponding test of Miettinen and Nurminen (1985) given above.

Multiplicity Adjustment

Because *G* z-tests between treatment groups and the control group are run when analyzing the results of this study, many statisticians recommend that a Bonferroni adjustment be applied. This adjustment is easy to apply: the value of alpha that is used in the test is found by dividing the original alpha by the number of tests. For example, if the original alpha is set at 0.05 and the number of treatment (not including the control) groups is five, the individual tests should be conducted using an alpha of 0.01.

The main criticism of this procedure is that if there are many tests, the value of alpha becomes very small. To mitigate against this complaint, some statisticians recommend separating the treatment groups into those that are of primary interest and those that are of secondary interest. The Bonferroni adjustment is made by the using the number of primary treatments rather than the total number of treatments.

There are some who advocate ignoring the adjustment entirely in the case of randomized clinical trials. See for example Saville (1990) and the discussion in chapter 14 of Machin, Campbell, Tan, and Tan (2018).

Size of the Control Group

Because the control group is used over and over, some advocate increasing the number of clusters in this group. The standard adjustment is to include \sqrt{G} clusters in the control group for each cluster in one of the treatment groups. See Machin, Campbell, Tan, and Tan (2018, pages 231-232). Note that often, the treatment groups all have the same sample size.

Example 1 – Finding the Sample Size

A cluster-randomized, multi-arm trial is being designed to compare two treatments against the standard drug in patients with a specific type of disease. They plan to use the Farrington and Manning likelihood score test based on the response ratio to analyze the data.

Historically, the standard treatment has enjoyed a 70% cure rate. The new treatments both reduce the seriousness of certain side effects of the standard treatment. Thus, the new treatments will be adopted even if they are slightly less effective than the standard treatment. The researchers will recommend adoption of the either of the new treatments that exhibit a cure rate that is no more than 20% less than the cure rate of the standard treatment. The value of R0 is 0.8.

The researchers will recruit patients from various hospitals. All patients at a particular hospital will receive the same treatment. They anticipate an average of 20 patients per hospital. They want to see the impact on cluster count of having cluster sizes ranging for 10 to 30.

The investigators would like a sample size large enough to find statistical significance at the 0.025 level and a power of 0.90 in each test. Based on similar studies, they estimate the intracluster correlation to be 0.002.

Since the control group will be used twice, they set the control group cluster allocation to $\sqrt{G} = \sqrt{2} = 1.414$. The two treatment allocations are set to 1.0.

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
Higher Proportions Are	Better (H1: R > R0)
Power of Each Test	
Overall Alpha	
Bonferroni Adjustment	
Group Allocation	Enter Group Allocation Pattern, solve for group numbers of clusters
M (Items Per Cluster)	
R0 (Non-Inferiority Ratio)	0.8
Control Proportion	
Control Items Per Cluster	M
Control Cluster Allocation	1.414
Set A Number of Groups	2
Set A Proportion	
Set A Items Per Cluster	M
Set A Cluster Allocation	1
Set B Number of Groups	0
Set C Number of Groups	0
Set D Number of Groups	
More	Unchecked
ρ (Intracluster Correlation)	0.002

Output

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

Numeric Reports

Numeric Results	
Solve For:	Sample Size
Group Allocation:	Enter Group Allocation Pattern, solve for group numbers of clusters
Test Type:	Farrington and Manning Likelihood Score Test
Hypotheses:	H0: Ri ≤ R0 vs. H1: Ri > R0
Number of Groups:	3
Bonferroni Adjustment:	Standard Bonferroni (Divisor = 2)

								Rati	io			
	Po	ower	Number of		Items Per			Non-				Alpha
Comparison	Target	Actual	Clusters Ki	Cluster Allocation	Cluster Mi	Size Ni	Proportion Pi		Actual Ri	ICC ρ		Bonferroni- Adjusted
Control			30	1.414	10	300	0.7	0.8		0.002		
vs A1	0.9	0.90773	21	1.000	10	210	0.7	0.8	1	0.002	0.025	0.0125
vs A2	0.9	0.90773	21	1.000	10	210	0.7	0.8	1	0.002	0.025	0.0125
Total			72			720						
Control			16	1.414	20	320	0.7	0.8		0.002		
vs A1	0.9	0.91699	11	1.000	20	220	0.7	0.8	1	0.002		0.0125
vs A2		0.91699	11	1.000	20	220	0.7	0.8		0.002		0.0125
Total			38			760						
Control			11	1.414	30	330	0.7	0.8		0.002		
vs A1	0.9	0.93126	8	1.000	30	240	0.7	0.8	1	0.002	0.025	0.0125
vs A2	0.9	0.93126	8	1.000	30	240	0.7	0.8	1	0.002	0.025	0.0125
Total			27			810						
Comparison]		up that is ir he compari				veen the tre	atment and	d contro	l displ	ayed on	this report
Target Powe	er	The pov		Power is p	orobability	of rejecti	ng a false n	ull hypothe	sis for t	his co	mparisor	n. This power
Actual Powe	er	The pov	ver actually	achieved.								
Ki			mber of clus		ith group.	The total	number of	clusters is I	reported	d in the	e last row	v of the
Allocation			ster allocati		•	p. The v	alue on eac	h row repre	esents f	the rela	ative nun	nber of
Mi		The ave	erage numb	er of items	per cluster	(or aver	age cluster	size) in the	e ith aro	up.		
Ni							imple size i				of the col	umn
Pi							the power					
R0							veen conclu			ont ic r	oon infor	ior or
NU		THE HOI	i-menonity			any Delv		iung that a	ueatin		ion-mer	

The non-inferiority ratio is the ratio boundary between concluding that a treatment is non-inferior or inferior.

RI	I ne ratio of the ith group proportion and the control group proportion at which the power is calculated. The
	formula is Ri = Pi / Pc.
ρ	The intracluster correlation (ICC). The correlation between subjects within a cluster.
Overall Alpha	The probability of rejecting at least one of the comparisons in this experiment when each null hypothesis

Overall Alpha The probability of rejecting at least one of the comparisons in this experiment when each null hypothesis is true.

Bonferroni Alpha The adjusted significance level at which each individual comparison is made.

Summary Statements

A parallel, 3-group cluster-randomized design (with one control group and 2 treatment groups) will be used to test whether the proportion for each treatment group is non-inferior to the control group proportion, with a non-inferiority ratio of 0.8 (H0: R ≤ 0.8 versus H1: R > 0.8, R = Pi / Pc). The hypotheses will be evaluated using 2 one-sided, two-sample, Bonferroni-adjusted (divisor = 2) Farrington and Manning likelihood score tests, with an overall (experiment-wise) Type I error rate (α) of 0.025. The control group proportion is assumed to be 0.7. The intracluster correlation is assumed to be 0.002. The average cluster size (number of subjects or items per cluster) for the control group is assumed to be 10, and the average cluster size for each of the treatment groups is assumed to be 10 and 10. To detect the treatment proportions 0.7 and 0.7 with at least 90% power for each test, the control group cluster count needed will be 30 and the number of needed clusters for the treatment groups will be 21 and 21 (totaling 72 clusters overall).

References

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Gart, John J. and Nam, Jun-mo. 1990. 'Approximate Interval Estimation of the Difference in Binomial Parameters: Correction for Skewness and Extension to Multiple Tables.' Biometrics, Volume 46, Issue 3, 637-643.

Machin, D., Campbell, M.J., Tan, S.B, and Tan, S.H. 2018. Sample Sizes for Clinical, Laboratory, and Epidemiology Studies, 4th Edition. Wiley Blackwell.

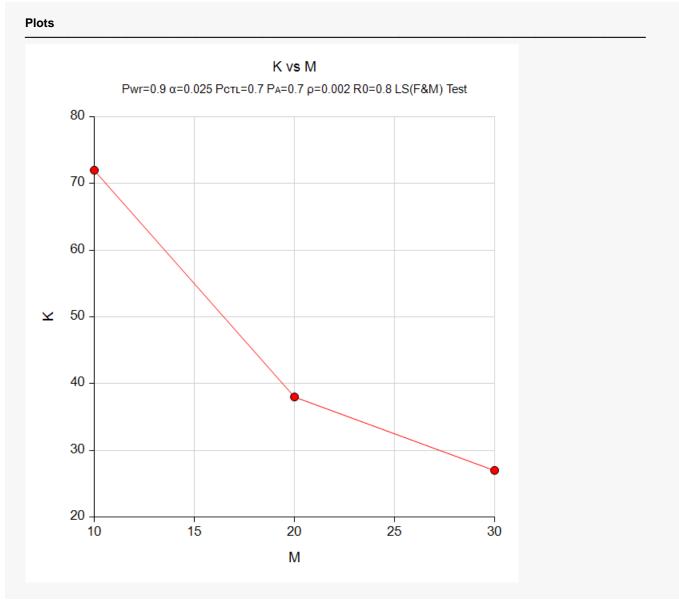
Miettinen, O.S. and Nurminen, M. 1985. 'Comparative analysis of two rates.' Statistics in Medicine 4: 213-226.

This report shows the numeric results of this power study. Notice that the results are shown in blocks of four rows at a time. Each block represents a single design.

PASS Sample Size Software

Multi-Arm Non-Inferiority Tests for the Ratio of Treatment and Control Proportions in a Cluster-Randomized Design

Plots Section



This plot gives a visual presentation to the results in the Numeric Report. We can quickly see the impact on the total cluster count, K, of increasing the cluster size, M.

Example 2 – Validation using a Previously Validated Procedure

We could not find a validation result in the statistical literature, so we will use a previously validated **PASS** procedure (**Non-Inferiority Tests for the Ratio of Two Proportions in a Cluster-Randomized Design**) to produce the results for the following example.

We will use a slightly modified version of the settings from Example 1. Specifically we will set Ki = 12 and M = 20. The Bonferroni adjustment changes the significance level from 0.025 to 0.0125.

The **Non-Inferiority Tests for the Ratio of Two Proportions in a Cluster-Randomized Design** procedure is set up as follows.

Design Tab	
Solve For	Power
Higher Proportions Are	Better (H1: P1/P2 > R0)
Alpha	0.0125
K1 (Clusters in Group 1)	12
M1 (Average Cluster Size)	20
K2 (Clusters in Group 2)	K1
M2 (Average Cluster Size)	M 1
R0 (Non-Inferiority Ratio)	0.8
R1 (Actual Ratio)	1.0
P2 (Group 2 Proportion)	0.7
ICC (Intracluster Correlation)	0.002

This set of options generates the following report.

Solve Fo	r:	Power											
Groups:		1 = 1	Freatr	ment, i	2 = Re	eference							
Test Stat	istic:	Like	lihood	d Scor	e Test	(Farringtor	n & Manning)						
Hypothes	ses:	H0:	P1/F	P2 ≤ R	0 vs.	H1: P1 / I	P2 > R0						
						Total	Pro	portions		Ratio		Intraclustor	
						Total Sample Size	Pro Non-Inferiority	portions Actual	Reference	Ratio	Actual	Intracluster Correlation	
Power						Sample			Reference P2		Actual R1		Alpha

The power is computed to be 0.91116.

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.

Solve For	Power
Higher Proportions Are	Better (H1: R > R0)
Overall Alpha	0.025
Bonferroni Adjustment	
Group Allocation	
Ki (Group Number of Clusters)	12
M (Items Per Cluster)	20
R0 (Non-Inferiority Ratio)	0.8
Control Proportion	0.7
Control Items Per Cluster	M
Set A Number of Groups	2
Set A Proportion	0.7
Set A Items Per Cluster	M
Set B Number of Groups	0
Set C Number of Groups	0
Set D Number of Groups	0
More	Unchecked
ρ (Intracluster Correlation)	0.002

Output

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

Solve For:PowerTest Type:Farrington and Manning Likelihood Score TestHypotheses:H0: Ri \leq R0 vs. H1: Ri > R0Number of Groups:3Bonferroni Adjustment:Standard Bonferroni (Divisor = 2)										
				Sample		Rati	0		Alasha	
		Number of	Items Per			Non-				Alpha
Comparison	Power	Clusters Ki	Cluster Mi	Size Ni	Proportion Pi	Inferiority R0	Actual Ri	ICC ρ	Overall	Bonferroni- Adjusted
		12	20	240	0.7	0.8		0.002		
Control		12	20	240	0.7	0.8	1	0.002	0.025	0.0125
Control vs A1	0.91116									
	0.91116		20	240	0.7	0.8	1	0.002	0.025	0.0125

As you can see, the power is 0.91116 for both treatment groups which match the power found in the validation run above. The procedure is validated.