

## A Macro for Computing the Mantel-Fleiss Criterion

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### ABSTRACT

When using chi-square statistics to evaluate the relationship between two dichotomous variables, the test statistics approach the chi-square distribution as the sample size increases. In practice, the validity of the chi-square approximation depends on the expected cell counts exceeding five. For examining three-way tables using the Cochran-Mantel-Haenszel (CMH) statistic, the same rule does not apply. Mantel and Fleiss (1980) proposed an extension to the 'minimum of five' rule for  $2 \times 2 \times H$  tables. We propose a user-friendly macro to supplement an analysis that uses the CMH one-degree-of-freedom test. The macro outputs all calculated components of the criterion to a SAS<sup>®</sup> data set and accompanying log.

### INTRODUCTION

It is common in many areas of statistical practice to examine the relationship between a binary response variable and a binary predictor variable in the presence of a potentially confounding categorical factor. Data are typically presented in a series of partial  $2 \times 2$  tables segmented by a stratification factor. Cochran (1954) first proposed a test of conditional independence in  $2 \times 2 \times H$  tables, treating the rows in each  $2 \times 2$  table as independent binomials. Mantel and Haenszel (1959) proposed a similar, but more generalized test based on the hypergeometric distribution (Agresti, 1996).

The CMH method tests whether the conditional odds ratio between the response and predictor variables equals one in each partial  $2 \times 2$  table. The approach conditions on the row and column totals in each partial  $2 \times 2$  table and, without loss of generality, utilizes the cell in the first row and column ( $n_{h11}$ ) (Agresti, 1996).

The test statistic

$$\frac{\{\sum_{h=1}^q n_{h11} - \sum_{h=1}^q m_{h11}\}^2}{\sum_{h=1}^q v_{h11}}$$

(where  $m_{h11} = \frac{n_{h1+}n_{h+1}}{n_h}$  is the expected value of  $n_{h11}$  and  $v_{h11} = \frac{n_{h1+} n_{h2+} n_{h+1} n_{h+2}}{n_h^2(n_h-1)}$  is its variance under the null hypothesis) has a large-sample chi-square distribution with one degree of freedom (Stokes, Davis and Koch, 2000).

In general, the CMH statistic is valid with sparse data but requires a large sample size. Mantel and Fleiss showed that it is appropriate to compute the CMH p-value from a chi-square distribution with one degree of freedom if

$$\min \left\{ \left[ \sum_{h=1}^q m_{h11} - \sum_{h=1}^q (n_{h11})L \right], \left[ \sum_{h=1}^q (n_{h11})U - \sum_{h=1}^q m_{h11} \right] \right\}$$

exceeds 5, where  $(n_{h11})L = \max(0, n_{h1+} - n_{h+2})$  and  $(n_{h11})U = \min(n_{h+1}, n_{h1+})$  (Stokes, Davis and Koch, 2000).

When performing an analysis using the CMH test, PROC FREQ produces asymptotic p-values for the general association (CMH) test statistic. We propose supplementing this analysis with the %MantelFleiss macro presented in this article.

### TABLE STRUCTURE

Visually, the table structure follows where  $X=2$ ,  $Y=2$  and  $h$  runs from 1 to  $q$ .

For  $h=1$

X	Y		
	$n_{111}$	$n_{112}$	$n_{11+}$
	$n_{121}$	$n_{122}$	$n_{12+}$
	$n_{1+1}$	$n_{1+2}$	$n_1$

For  $h=2$

X	Y		
	$n_{211}$	$n_{212}$	$n_{21+}$
	$n_{221}$	$n_{222}$	$n_{22+}$
	$n_{2+1}$	$n_{2+2}$	$n_2$

⋮  
⋮  
⋮

For  $q^{\text{th}}$  table

X	Y		
	$n_{q11}$	$n_{q12}$	$n_{q1+}$
	$n_{q21}$	$n_{q22}$	$n_{q2+}$
	$n_{q+1}$	$n_{q+2}$	$n_q$

## MACRO

**MACRO CALL:** %MantelFleiss(In,Strat,Var1,Var2,Count)

The macro consists of five parameters. Parameters &IN, &STRAT, &VAR1 and &VAR2 are required, whereas &COUNT is optional. The &IN parameter denotes the input data set and requires unique observations. &STRAT, &VAR1 and &VAR2 represent the stratification, row, and column variables respectively. &COUNT issues the WEIGHT statement in PROC FREQ. If the &COUNT parameter is omitted, all observations in the input data set are assumed to have a weight of one.

For illustration, we divide the criterion into four components. We compute each component using the following syntax:

- 1) Summing the expected values in each  $n_{h11}$  cell for  $h=1,2,\dots,q$ .

$$\sum_{h=1}^q m_{h11}$$

## SYNTAX:

```

%*get expected counts and totals in each partial table;
PROC FREQ data = &In;
  tables &Strat*&Var1*&Var2 / expected outexpect out = _expect;
  %if %nrbquote(&Count.) ne %then weight &Count.;;
  ODS output CrossTabFreqs = _counts;
RUN;

%*get sum of expected cell counts in cell n11 in each partial table
(assign to macro var SUMEXP);
DATA _null_;
  set _expect end = eof;
  by &Strat;
  retain sumexp 0;
  if first.&Strat then do;
    sumexp = sumexp + expected;
  end;
  if eof then call symput('sumexp',put(sumexp,8.4));
RUN;

```

- 2) Summing the maximum difference between the row one total and the column two total for each of the  $h$  strata. The calculations span two data steps, but we combine the syntax here for illustration. See APPENDIX for unified method.

$$\sum_{h=1}^q (n_{h11}) L = \max(0, n_{11+} - n_{1+2}) + \max(0, n_{21+} - n_{2+2}) + \dots + \max(0, n_{q1+} - n_{q+2})$$

**SYNTAX:**

```
retain n11_ n1_1 n1_2;

if first.&Strat._ then do;
  n11_ = 0;
  n1_1 = 0;
  n1_2 = 0;
end;

%*get each total;
if &Var1._ = 1 then n11_ = frequency; %*for row one in the hth strata;
if &Var2._ = 1 then n1_1 = frequency; %*for column one in the hth strata;
if &Var2._ = 2 then n1_2 = frequency; %*for column two in the hth strata;

%*define (nh11)L for final computation;
max_ = max(0,n11_-n1_2);

%*****;
%*get sum of maximum for each of the hth strata*;
%*****;

retain mf_sum1;

if _n_ = 1 then do;
  mf_sum1 = 0;
end;

mf_sum1 = mf_sum1 + max_;
```

- 3) Summing the minimum of the column one total and the row one total across all strata. Except for using a different formula (for maximum), the abbreviated code below is nearly identical to step two. See APPENDIX for unified method.

$$\sum_{h=1}^q (n_{h11}) U = \min(n_{1+1}, n_{11+}) + \min(n_{2+1}, n_{21+}) + \dots + \min(n_{q+1}, n_{q1+})$$

**SYNTAX:**

```
%*define (nh11)U for final computation;
min_ = min(n1_1,n11_);

retain mf_sumu;
if _n_ = 1 then do;
  mf_sumu = 0;
end;

mf_sumu = mf_sumu + min_;
```

- 4) Computing the criterion by combining all steps. Compare summed expected values (step one) to maximum (step two) and minimum (step three), then find the minimum. If minimum is greater than five, then the criterion is satisfied.

$$R = \min \left\{ \left[ \sum_{h=1}^q m_{h11} - \sum_{h=1}^q (n_{h11})L \right], \left[ \sum_{h=1}^q (n_{h11})U - \sum_{h=1}^q m_{h11} \right] \right\} > 5$$

**SYNTAX:**

```

if eof then do;
  mflendpt = &sumexp - mf_suml;
  mfrendpt = mf_sumu - &sumexp;
  mf_r      = min(mflendpt,mfrendpt);

  put "Sum of lower bound (sum[(nh11)L]) = " mf_suml;
  put "Sum of upper bound (sum[(nh11)U]) = " mf_sumu;
  put "Left end-point: sum[mh11] - sum[(nh11)L] = " mflendpt;
  put "Right endpoint: sum[(nh11)U] - sum[mh11] = " mfrendpt;
  put "Mantel-Fleiss R = min[" mflendpt "," mfrendpt "] = " mf_r;

  if mf_r>5 then put "Mantel-Fleiss criterion satisfied (since R > 5)";
  else          put "Mantel-Fleiss criterion NOT satisfied (since R <= 5)";
  output;
end;

```

**EXAMPLE1**

The following is a hypothetical scenario comparing treatments A and B within three sites (Arkansas, Indiana and Illinois). In this example, the criterion is not satisfied. Therefore the chi-square approximation is not valid:

**DATA SETUP:**

```

PROC FORMAT;
  value trtf
    1 = 'TreatmentA'
    2 = 'TreatmentB'
  ;

  value sitef
    1 = 'Arkansas'
    2 = 'Indiana'
    3 = 'Illinois'
  ;

  value respf
    1 = 'Present'
    2 = 'Absent'
  ;
RUN;

DATA test;
  do site = 1 to 3;
    do treat = 1 to 2;
      do resp = 1 to 2;
        input counts @@;
        output;
      end;
    end;
  end;
  label site = "Site" treat = "Treatment" resp = "Response";
  format treat trtf. site sitef. resp respf.;
cards;
12 5 7 3
1 2 5 2
0 9 5 6
;
RUN;

```

**TABLE STRUCTURE**

*site=Arkansas (h=1)*

Treatment	Response		Total
	Present	Absent	
A	12	5	17
B	7	3	10
Total	19	8	27

*site=Indiana (h=2)*

Treatment	Response		Total
	Present	Absent	
A	1	2	3
B	5	2	7
Total	6	4	10

*site=Illinois (h=3)*

Treatment	Response		Total
	Present	Absent	
A	0	9	9
B	5	6	11
Total	5	15	20

**MACRO CALL:** %MantelFleiss(test,site,treat,resp,counts)

- 1) Summing the expected values in each  $n_{h11}$  cell for  $h=1,2,\dots,q$ .

**OUTPUT TO LOG:**

Sum of expected values in cells nh11: sum(mh11) = 20.5130

- 2) Summing the maximum difference between the row one total and the column two total.

**OUTPUT TO LOG:**

Sum of lower bound (sum[(nh11)L]) = 9

- 3) Summing the minimum of the column one total and the row one total.

**OUTPUT TO LOG:**

Sum of upper bound (sum[(nh11)U]) = 25

- 4) Computing the criterion by combining steps two and three.

**OUTPUT TO LOG:**

Left end-point: sum[mh11] - sum[(nh11)L] = 11.513  
 Right endpoint: sum[(nh11)U] - sum[mh11] = 4.487  
 Mantel-Fleiss R = min[11.513 , 4.487 ] = 4.487  
 Mantel-Fleiss criterion NOT satisfied (since R <= 5)

**EXAMPLE2**

The following example illustrates a case in which the criterion is satisfied:

**TABLE STRUCTURE**

*site=Arkansas*

Treatment	Response		Total
	Present	Absent	
A	12	5	17
B	7	3	10
Total	19	8	27

site=Indiana			
Treatment	Response		Total
	Present	Absent	
A	1	2	3
B	5	2	7
Total	6	4	10

site=Illinois			
Treatment	Response		Total
	Present	Absent	
A	5	9	14
B	5	6	11
Total	10	15	25

- 1) Summing the expected values in each  $n_{h1}$  cell for  $h=1,2,\dots,q$ .

**OUTPUT TO LOG:**

Sum of expected values in cells nh11: `sum(mh11) = 19.3630`

- 2) Summing the maximum difference between the row one total and the column two total.

**OUTPUT TO LOG:**

Sum of lower bound (`sum[(nh11)L]`) = 9

- 3) Summing the minimum of the column one total and the row one total.

**OUTPUT TO LOG:**

Sum of upper bound (`sum[(nh11)U]`) = 30

- 4) Computing the criterion by combining steps two and three.

**OUTPUT TO LOG:**

Left end-point: `sum[mh11] - sum[(nh11)L] = 10.363`  
 Right endpoint: `sum[(nh11)U] - sum[mh11] = 10.637`  
 Mantel-Fleiss R = `min[10.363 ,10.637 ] = 10.363`  
 Mantel-Fleiss criterion satisfied (since  $R > 5$ )

**CONCLUSION**

The macro presented in this paper determines the validity of the chi-square approximation for the CMH one-degree-of-freedom test. If the Mantel-Fleiss criterion is satisfied, the chi-square approximation is valid. However, if the criterion is not satisfied, other techniques are necessary.

Techniques for producing exact p-values are illustrated by Agresti (1996). Both PROC LOGISTIC and PROC MULTTEST provide tools for obtaining exact CMH p-values. The LOGISTIC approach uses the STRATA and EXACT statements. For MULTTEST, the STRATA statement and PERM option will produce similar results. See <http://support.sas.com/kb/32/711.html> for detailed examples.

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## APPENDIX

```
%macro MantelFleiss(In,Strat,Var1,Var2,Count);

%*-----;
%*  &In      - input data set                      ;
%*  &Strat   - stratification variable (e.g., site, race, etc.);
%*  &Var1    - row variable                        ;
%*  &Var2    - column variable                    ;
%*  &Count   - optional weight variable           ;
%*-----;

%*reset output data sets;
PROC DATASETS nodetails nolist;
  delete _counts _expect _mantel;
RUN;
QUIT;

%*determine number of strata levels;
PROC SQL noprint;
  select count(distinct &strat) into: stratflag from &In;
QUIT;
%put **Number of strata levels &stratflag;

%*get expected counts and totals in each partial table;
PROC FREQ data = &In;
  tables &Strat*&Var1*&Var2 / expected outexpect out = _expect;
  %if %nrbquote(&Count.) ne %then weight &Count.;;
  ODS output CrossTabFreqs = _counts;
RUN;

%*get sum of expected cell counts in cell n11 in each partial table
  (assign to macro var SUMEXP);
DATA _null_;
  set _expect end = eof;
  by &Strat;
  retain sumexp 0;
  if first.&Strat then do;
    sumexp = sumexp + expected;
  end;
  if eof then call symput('sumexp',put(sumexp,8.4));
RUN;
%put Sum of expected values in cells nh11: sum(mh11) = %cmpres(&sumexp);

PROC SORT data = _counts;
  by &Strat &Var1 &Var2;
RUN;
```

```

%*Subset to the needed totals output by PROC FREQ
number the &Strat, &Var1, and &Var2 for use below
0 - totals
1 - first &Strat/&Var1/&Var2
2 - second &Strat/&Var1/&Var2
...;

DATA _mantelf1;
%*only include marginal totals from ODS;
set _counts(where = (_type_ not in ('100' '111')));
by &Strat &Var1 &Var2;
retain &Strat._ 0 &Var1._ &Var2._;

%*reset for change in strata;
if first.&Strat then do;
    &Var1._ = 0; &Var2._ = 0;
end;

%*get numeric version of strata variable;
if first.&Strat then &Strat._ = &Strat._ + 1;

%*define zero as total for any &Var1/&Var2;
if missing(&Var1) then &Var1._ = 0;
    else &Var1._ + 1;

if missing(&Var2) then &Var2._ = 0;
    else &Var2._ + 1;

keep &Strat._ &Var1._ &Var2._ &Strat. &Var1. &Var2. frequency;
RUN;

DATA _mantelf2;
set _mantelf1;
by &Strat._;

retain n11_ n1_1 n1_2;

if first.&Strat._ then do;
    n11_ = 0;
    n1_1 = 0;
    n1_2 = 0;
end;

%*get each total;
if &Var1._ = 1 then n11_ = frequency; %*for row one in the hth strata;
if &Var2._ = 1 then n1_1 = frequency; %*for column one in the hth strata;
if &Var2._ = 2 then n1_2 = frequency; %*for column two in the hth strata;

%*define (nh11)L and (nh11)U for final computation;
max_ = max(0,n11_-n1_2);
min_ = min(n1_1,n11_);

if last.&Strat._;

RUN;

%*sum across min and max to get lower/upper bounds of criterion;
DATA _mantelf3(keep = mf_suml mf_sumu mflendpt mfrendpt mf_r);
set _mantelf2 end = eof;

retain mf_suml mf_sumu;

if _n_ = 1 then do;
    mf_suml = 0; mf_sumu = 0;

```

```

end;

mf_suml = mf_suml + max_;
mf_sumu = mf_sumu + min_;

if eof then do;
    mflendpt = &sumexp-mf_suml;
    mfrendpt = mf_sumu-&sumexp;
    mf_r      = min(mflendpt,mfrendpt);

    put "Sum of lower bound (sum[(nh11)L]) = " mf_suml;
    put "Sum of upper bound (sum[(nh11)U]) = " mf_sumu;
    put "Left end-point: sum[mh11] - sum[(nh11)L] = " mflendpt;
    put "Right endpoint: sum[(nh11)U] - sum[mh11] = " mfrendpt;
    put "Mantel-Fleiss R = min[" mflendpt "," mfrendpt "] = " mf_r;

    if mf_r>5 then put "Mantel-Fleiss criterion satisfied (since R > 5)";
    else          put "Mantel-Fleiss criterion NOT satisfied (since R <= 5)";
    output;
end;

label
mf_suml = "Summation of Maximium (sum[(nh11)L])"
mf_sumu = "Summation of Minumum (sum[(nh11)U])"
mflendpt = "Left End-Point: sum[mh11] - sum[(nh11)L]"
mfrendpt = "Right End-Point: sum[(nh11)U] - sum[mh11]"
mf_r      = "R Value: min[sum[mh11] - sum[(nh11)L],sum[(nh11)U] - sum[mh11]]";
RUN;
%mend;

```