1 GENSTAT and S-PLUS Codes For Recovering Interblock and Intergradient Information

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Abstract

In order to facilitate the computational needs of researchers, computer programs using the GENSAT and S-PLUS software packages are presented. In particular, codes are presented for both fixed and mixed model analyses for an incomplete block design, for a lattice rectangle design, and for differential gradients within incomplete blocks or within the rows (columns) of a lattice rectangle design. The block or row and column effects are treated as random effects in order obtain a more efficient analysis of experimental data and to make use of the information contained in inter-effect mean squares. Also, differential gradients may occur within incomplete blocks or rows (columns) and need to be treated as random effects. The GENSTAT package is widely used by experimenters, especially outside the United States. The S-PLUS package is gaining in popularity among researchers, especially statisticians.

## 1. Introduction

 When analyzing data from experiments designed as incomplete block or row-column designs, an efficient analysis recovers the treatment information contained in the incomplete blocks or in the rows and columns of the design and uses a mixed model procedure. The blocks and the rows and columns are considered to be random effects., resulting in the so-called mixed model situation. The resulting adjusted means containing this additional information have a smaller mean square error than means ignoring this information.

Textbook analyses may be appropriate for certain types of experimental variation but quite inappropriate for other types. Patterns of spatial variation not anticipated may arise during the course of conducting the experiment. For example, invasion of insects may not occur in the manner used for blocking. Water or freezing damage to experimental units may occur in patterns not anticipated by the experimenter. Differential gradients may occur within the complete blocks or within the incomplete blocks. In such instances, textbook analyses will be inappropriate and need to be replaced by an appropriate analysis. The differential gradients should be considered as random effects.

This paper demonstrates how to program analyses on GENSTAT and S-PLUS for a triple lattice (incomplete block) designed experiment, for a lattice square (lattice rectangle) designed experiment, and for differential linear gradients within the rows (or incomplete blocks) of a lattice square designed experiment. GENSTAT codes are presented for the ANOVA, regression (fixed effects model), and variance component using REML (restricted maximum likelihood) analyses. Comparable S-PLUS codes are given for the variance component analysis using REML solutions for the variance components. It should be noted that different solutions for the variance components are obtained from the two programs and hence, the adjusted means will differ to some extent. S-PLUS results are the same as obtained by SAS PROC MIXED. It appears that the developers of the packages used different boundary conditions for their REML solutions of variance components.

2. Recovery of Interblock Information in Incomplete Block Designs

 The GENSTAT regression program is a fixed effects or intrablock modeling of the responses. The information in the blocks is ignored and intrablock treatment means are obtained. The code for the incomplete block data set named fed933.dat, Example XI-3 in Federer (1955), is:

```
55
56
       "gn-1-1 Triple lattice"
57
       units [27]
58
       factor r, b, t
59
       open 'fed933.dat'; channel = 2
60
       read [channel = 2] y, r, b, t;
61
       model v
62
       terms r/b + t
63
       fit [print=accumulated; fprobability=yes] r/b + t
64
       predict t
65
       stop
```

The first line names the analysis, the second gives the number of responses to be analyzed, and the third line lists the factors (class variables in SAS notation). The fourth line opens the data set named fed933.dat. The fifth line denotes the input from fed933.dat and the entries must be in the correct order with Y = YIELD or RESPONSE, R = REPLICATE or COMPLETE BLOCK, B = INCOMPLETE BLOCK, and T = TREATMENT. The sixth line states that response Y is being modeled and the seventh line gives the terms of the linear model used *sans* the error term. The eighth line requests a nested or accumulated (Type I) analysis of variance table with probability statements for F-values. The ninth line obtains intrablock (fixed effects) means with their standard errors.

The GENSTAT analysis of variance (ANOVA) program recovers interblock information using ANOVA solutions for the variance components which are the ones usually used in textbooks but are being replaced by some form of maximum likelihood solutions. The code is:

```
80
       "gn-1-0 Triple lattice"
81
      unit [27]
82
      factor r, b, t
83
      open 'fed933.dat'; channel = 2
84
      read [channel=2] y, r, b, t;
85
      factor [levels=3] pa, pb, pc
86
      calc pa = newlevels(t; ! ((1...3)3)
87
      & pb = newlevels(t; ! (3(1...3))
88
      & pc = newlevels(t; ! (1,2,3,3,1,2,2,3,1))
89
      blocks r/b
90
      treatments t//(pa+pb+pc)
91
      anova [print=aovt, info, cbmeans; pse=means; fprobability=yes] y
92
```

An analysis of variance table with F statistics and probabilities are the same as for the regression analysis. The adjusted means recovering interblock information along with standard errors are obtained, using ANOVA solutions for the variance components.

The GENSTAT variance component procedure uses REML solutions for the variance components to obtain the adjusted treatment means. The variance component solutions are constrained to be positive unless specified otherwise. The program for this analysis is:

```
"gn-1-2.n Triple lattice"

103 units [27]

104 factor r, b, t

105 open 'fed933.dat'; channel=2

106 read [channel=2] y, r, b, t;

107 vcomponents [fixed=t] r/b

108 reml [print=means, comp,stra; pse=estimates] y
```

```
109
       stop
110
       In the vcomponents statement the fixed effect is treatment (T) and the random effects are indicated by
111
112
       R/B. i.e., R and B nested within R are random effects. The output from this program contains the REML
       solutions for the variance components and their approximate standard errors, a matrix of coefficients of
113
114
        variance components, the adjusted means and their standard error.
115
       The corresponding S-PLUS code for the above analysis is:
116
117
       # SP-1 Triple lattice
118
119
       #Read data
                dat <- read.table('fed933.dat', col.names=c('y', 'r', 'b', 't'))
120
121
                attach (dat)
122
       # Convert r,b,t to factors
123
                r <- as.factor(r)
                b <- as.factor(b)
124
125
                t <- as.factor(t)
        # Fit model & print results
126
127
                e < -rinfo(y \sim r/b + t, t)
128
                print ('Variance components & standard errors")
                print (cbind (e$vc,e$se.vc,deparse.level=2))
129
                print ("REML means & standard errors')
130
131
                print (cbind (e$mn,e$se.mn,deparse.level=2))
132
        #Clean up
133
                detach ()
134
                rm(dat,r,b,t,e)
135
136
        This program prints out the variance components and their standard errors (questionable) and the REML
        means and their standard errors. For this particular example, the adjusted means are equal to those
137
        obtained from the GENSTAT variance component analysis because the ANOVA and REML solutions for
138
139
        the variance components are equal. In general, they will not be the same but the S-PLUS and SAS PROC
        MIXED results will be the same.
140
141
142
        3. Recovery of Interrow and Intercolumn Information in Lattice Rectangle Designs
143
144
        The data set used to illustrate the program for recovering interrow and intercolumn information for a
145
        lattice rectangle designed experiment is the lattice square design given in Table 12.5 of Cochran and Cox
146
        (1957). This data set is given the name lsgr1645.dat in the program. The GENSTAT regression code for
147
        a fixed effects analysis is:
148
149
        "gn-2.1 gn Lattice square"
150
        units [80]
151
        open 'lsgr1645.dat'; channel = 2
        read [channel=2] yield, rep, row, col, grad, treat
152
153
        model yield
154
        terms rep/(row + col) + treat
155
        fit [print=accumulated; probability=yes] rep/(row + col) + treat
156
        predict treat
157
158
        The GRAD (gradient) column was included in the data file and is used for the analysis in the next section.
159
```

The GENSTAT ANOVA program which recovers interrow and intercolumn information is

160

161 162

"gn-2-0 gn Lattice square"

```
163
       units [80]
        factor yield, rep, row, col, grad, treat
164
165
       open 'lsgr1645.dat'; channel = 2
       read [channel=2] yield, rep, row, col, grad, treat
166
        blocks rep/(row + col)
167
        treatments treat
168
        anova [print=aovt, info, cbmeans; pse=means; fprobability=yes] yield
169
170
171
        The GENSTAT variance component program for this data set is
172
        "gn-2-2 Lattice square"
173
174
        units [80]
        factor rep, row, col, treat
175
        open 'lsgr1645.dat'; channel = 2
176
        read [channel=2] yield, rep, row, col, grad, treat
177
        vcomponents [fixed=treat] rep/(row+col); constraints = positive
178
        reml [print=means,comp,stra; pse=estimates] yield
179
180
        stop
181
        To obtain unconstrained solutions for variance components, simply omit the statement "constraints =
182
183
        positive" in the above.
184
         The corresponding S-PLUS code for recovering interrow and intercolumn information for the above
185
186
        example is
187
188
        #sp-2 Lattice square
189
        #Read data
190
          dat <- read.table('lsgr1645.dat', col.names=c('yield', 'rep', 'row', 'col', 'gard', 'treat'))
191
          attach(dat)
192
        #Convert rep,row,col,treat to factors
193
          rep <- as.factor(rep)
194
          row <- as.factor(row)
195
          col <- as.factor(col)
196
        treat <- as.factor(treat)
197
        #Fit model & print results
198
          e <- rinfo(yield ~ rep/(row+col) + treat, treat)
199
200
        The adjusted means and variance components obtained here agree with SAS PROC MIXED but not with
201
        the preceding GENSTAT results.
202
203
204
        4. Recovery of Interblock, or Interrow, and Intergradient Information in Incomplete Block or Lattice
205
        Rectangle Designs
206
207
        The data used to illustrate the recovery of interrow and intergradient information are those of the previous
208
        section where differential linear gradients within each row are considered rather than considering column
209
        effect within each complete block. The values included in the grad data column are the centered linear
        regression coefficients of position. The gradients and rows are considered to be random effects. Note that
210
        this analysis is equally applicable to an incomplete block design with differential gradients within each
211
212
        incomplete block. This form of spatial analysis may be a more appropriate analysis than the standard
213
        textbook one. The form of the gradient is linear but additional polynomial or other terms may be added if
214
        deemed appropriate.
215
```

A GENSTAT regression program for fixed effects is

216

```
217
218
       units [80]
219
       factor rep, row, treat
220
       open 'lsgr1645.dat'; channel = 2
       read [channel=2] yield, rep, row, col, grad, treat
221
222
       model yield
223
       terms rep/row/grad + treat
224
       fit [print=accumulated; fprobability=yes] rep/row/grad + treat
225
       predict treat
226
       stop
227
228
       A GENSTAT ANOVA program for this data set is
229
230
       units [80]
231
       factor rep, row, col, treat
232
       open 'lsgr1645.dat'; channel = 2
233
       read [channel=2] yield. rep, row, col, grad, treat
234
       blocks rep/(row+col)
235
       treatments treat
236
       anova [print=aovt,info,cbmeans; pse=means; fprobability=yes] yield
237
       stop
238
        A GENSTAT variance component code for REML solutions with positive restraints on the solutions is
239
240
241
       units [80]
242
       factor rep, row, treat
243
       open 'lsgr1645.dat'; channel = 2
       rear [channel = 2] yield, rep, row, col, grad, treat
244
245
        vcomponents [fixed=treat] rep/row/grad; constraints = positive
246
        read [print=means, comp, stra; pterms=treat; pse=estimates] yield
247
       stop
248
249
        If no restraints are to be placed on variance component solutions, omit the statement "constraints =
250
        positive".
251
252
        The S-PLUS code for recovering interrow and intergradient information for the above example is
253
254
       #sp-3 Lattice square with gradients
255
        #Read data
256
          dat <- read.table('lsgr1645.dat', col.names=c('yield', 'rep', 'row', 'col', 'grad', 'treat'))
257
          attach (dat)
258
        #Convert rep,row,treat to factors
259
          rep <- as.factor(rep)
260
          row <- as.factor(row)
261
         treat <- as.factor(treat)
262
        #Fit model & print results
263
          e <- rinfo(yield ~ rep/row/grad+treat, treat)
         print ('Variance components')
264
265
         print (e$vc)
266
         print ("REML means & standard errors')
267
         print (cbind(e$mn,e$se.mn,desparse.level=2))
268
        #Clean up
269
         rm(dat,rep,row,treat,e)
270
```

Function RINFO is called the model formula YIELD~REP/ROW/GRAD+TREAT where YIELD is the response variate, REP is replicate, ROW is row, GRAD is gradient, TREAT is treatment, and REP/ROW/GRAD indicates that rows are nested within replicates, and gradients are nested within rows.

5. Some Comments

GENSTAT regression and SAS PROC GLM programs produce the intra-effect or fixed-effect analyses and give the same results. GENSTAT ANOVA and VCOMPONENTS produce the same adjusted means and variance component solutions but different standard errors. SAS PROC Mixed differs from the preceding two programs in adjusted means, variance component solutions, and standard errors. If no constraints are put on the variance component solutions, then the three programs produce the same results. The adjusted means for S-PLUS agree with SAS but not with GENSTAT output when positive restraints are used. The results from the S-PLUS programs are the same as for SAS. The question of which REML solution for variance components to use is unresolved. Technical Reports (Barnard and Federer, 1996; Federer, 1995) illustrating the programs with examples and annotated computer outputs are available for individuals requiring more than the above programs.

6. Literature Cited

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