

An APL program for calculating pairwise and nonpairwise a posteriori tests on means

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When using experimental designs with more than three treatments, an investigator is often confronted with the problem of locating mean differences following rejection of the overall null hypothesis. That is, if no specific hypotheses have been proposed in advance that are amenable to analysis by a priori procedures (e.g., orthogonal comparisons using either the *t* or *F* ratio), it becomes necessary to use "data snooping" procedures a posteriori (Hays, 1973; Kirk, 1968; Lindquist, 1956; Winer, 1971).

Conceptual and procedural differences between a priori and a posteriori multiple comparisons arise, in part, out of concern with error rate. That is, "Should the probability of committing a Type I error be set at α for each individual comparison or should the probability of an error equal α or less for some larger conceptual unit such as the collection of comparisons?" (Kirk, 1968, p. 78). Conceptually, the problem is one of decision—a decision on the "conceptual unit for error rate" (e.g., individual comparison, hypothesis, family of comparisons, or the experiment). Confusion arises because, as the number of comparisons increases, so does the probability of making a false positive (Type I error) (Kirk, 1968; Ryan, 1962). Procedurally, the problem is one "of regulating and apportioning the Type I error rate" (Winer, 1971, p. 199). "For planned orthogonal comparisons, contemporary practice in the behavioral sciences favors setting the Type I error probability at α for each comparison. For planned and unplanned nonorthogonal comparisons it is suggested

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that the Type I error probability should be set at α for the collection of comparisons" (Kirk, 1968, p. 78). Several a posteriori multiple-comparison procedures exist which enable an investigator to set the Type I error probability equal to or less than α for the collection of comparisons. Familiarity with these a posteriori procedures is assumed, and the present discussion will be limited to a method by which these tests may be easily calculated. Kirk (1968) offers an excellent review of a posteriori procedures; the work of Hays (1973) and Winer (1971) complement that of Kirk. Use of the method (computer program) discussed below should result in substantial time savings, the alleviation of computational difficulties, and greater flexibility in choice of a posteriori tests.

The present paper describes a computer program (APL) that can be used to compute several commonly used a posteriori multiple-comparison tests: (1) Newman-Keuls, (2) Duncan's new multiple-range (NMR) test, (3) Tukey's honestly significant difference, Type a [h.s.d.(a)], (4) Tukey's honestly significant difference, Type b [h.s.d.(b)], (5) Fisher's least significant difference (l.s.d.) test, (6) Dunn's procedure (also known as the Bonferroni *t* statistic, and (7) Scheffé's method. Each of these tests is described in detail by Kirk (1968) with the exception of Tukey's Type b test, which is described by Winer (1971).

The a posteriori multiple-comparison APL program is written "interactively" (conversationally) and is intended for users who are only vaguely familiar with APL. It is presented in Table 1 and explained below.

Function 1: RBD. RBD is a function (fns) used in the analysis of a randomized block design, that is, a mixed model in which subjects (blocks) represent a random effect and treatments represent a fixed effect. Fns RBD, the output of which is a summary ANOVA table, is used to illustrate the a posteriori multiple-comparison APL program described herein. All that is necessary to use the fns is to type the name of the fns (RBD), followed by a space, and then the name of the data set. The data set may be built easily through the use of the general-purpose data building fns BUILDMTX, described by McGowan and Appel (1977). For illustrations of the use of fns RBD, see Table 2.

Function 2: POSTHOC. The purpose of fns POSTHOC is to build left and right arguments for fns APPC and ANPC (below) and to organize the necessary fns for the a posteriori analysis. To use the fns, one must type the name POSTHOC and respond appropriately to the queries (Qs) generated by the fns. The first query (Q-1) requests the type of a posteriori test desired (a posteriori select value). A response of any real number between one and nine will "select" the respective a priori test; "zero" causes a listing of all the available a posteriori tests to be printed with their respective

TABLE 1

A Posteriori Functions

Fns I RBD

```

▽ RBD XV
[1] RANDOMIZED BLOCK DESIGN
[2] MIXED MODEL [SUBJECTS=RANDOM; TREATMENTS=FIXED]
[3] V+ '
[4] SST+(BS+/, (XV*2))-(X+((+/, XV)*2)+((Q+1+ρXV)×(N+1+ρXV)))
[5] MSB+(SSB+(SB+(+/(+([1] XV)*2)+N))) -X)+(DFB+Q-1)
[6] MSS+(SSS+(SS+(+/(+/, XV)*2)+Q))) -X)+(DFS+N-1)
[7] MSR+(SSR+(BS-SB)-SS)+X)+(DFR+(N-1)×(Q-1))
[8] ZBAR+(+/[1] XV)+N
[9] (25ρV), 'ANALYSIS OF VARIANCE TABLE'
[10] (19ρV), 'RANDOMIZED BLOCK DESIGN: MIXED MODEL'
[11] 1 75 ρ'-
[12] (4ρV), 'SOURCE', (19ρV), 'SS', (10ρV), 'DF', (12ρV), 'MS', (11ρV), 'F'
[13] 1 75 ρ'-
[14] 'BETWEEN TREATMENTS', (3ρV), (14 5 ▼SSB), (2ρV), (6 0 ▼DFB), (4ρV), (14 5 ▼MSR), V, (10 3 ▼(MSB+MSR))
[15] V
[16] 'BETWEEN BLOCKS', (7ρV), (14 5 ▼SSS), (2ρV), (6 0 ▼DFS), (4ρV), (14 5 ▼MSS), V, (10 3 ▼(MSS+MSR))
[17] V
[18] (4ρV), 'RESIDUAL'
[19] '[BLOCK×TREATMENT]', (4ρV), (14 5 ▼SSR), (2ρV), (6 0 ▼DFR), (4ρV), (14 5 ▼MSR)
[20] 1 75 ρ'-
[21] 'TOTAL', (16ρV), (14 5 ▼SST), (2ρV), (6 0 ▼(Q×N)-1)
[22] 1 75 ρ'-

```

Fns II POSTHOC

```

▽ POSTHOC;AS;C;Q;V
[1] C+''''
[2] Q+10ρ(V+ ' )
[3] UP1:'TYPE A POSTERIORI SELECT VALUE [0=LISTING]'
[4] →DN1×1(AS+□)=0
[5] 'TYPE: 1) MEAN SQUARE ERROR, 2) COMPARISON SAMPLE SIZE'
[6] NP+□,AS
[7] 'TYPE MEAN VALUES OR VARIABLE MEAN NAME'
[8] ZBAR+|
[9] NP APPC ZBAR
[10] +0
[11] DN1:(11ρV), 'A POSTERIORI TEST', (24ρV), 'SELECT VALUE', (5ρV), 'SAMPLING DISTRIBUTION'
[12] 90ρ'-
[13] 'NEWMAN-KEULS', (46ρV), '1', Q, 'STUDENTIZED RANGE'
[14] 'DUNCAN', C, 'S NEW MULTIPLE RANGE [NMR]', (25ρV), '2', Q, 'DUNCAN', C, 'S QR'
[15] 'TUKEY', C, 'S HONESTLY SIGNIFICANT DIFFERENCE(A) [HSD(A)]', (7ρV), '3', Q, 'STUDENTIZED RANGE'
[16] 'TUKEY', C, 'S HONESTLY SIGNIFICANT DIFFERENCE(B) [HSD(B)]', (7ρV), '4', Q, 'STUDENTIZED RANGE'
[17] 'FISHER', C, 'S LEAST SIGNIFICANT DIFFERENCE [LSD]', (15ρV), '5', Q, 'STUDENT', C, 'S T'
[18] 'DUNN', C, 'S PROCEDURE [BONFERRONI T STATISTIC]', (17ρV), '6', Q, 'DUNN', C, 'S T', C, 'D'
[19] 'SCHEFFE', C, 'S S METHOD', (40ρV), '7', Q, 'F'
[20] 'SCHEFFE', C, 'S PROCEDURE [PAIRWISE/NONPAIRWISE]', (16ρV), '8', Q, 'F'
[21] 'DUNN', C, 'S PROCEDURE [PAIRWISE/NONPAIRWISE]', (19ρV), '9', Q, 'DUNN', C, 'S T', C, 'D'
[22] 90ρ'-
[23] 2 1 ρV
[24] →UP1

```

Fns III APCC

```

▽ NP APPC ZBAR;C;D;G;I;IA;IB;MS;N;NH;Q;T;Z;V;Y;W;Z
[1] MS+NP[IA+I+1]
[2] IB+φ(i(N+ρZBAR))
[3] Y+▲ZBAR
[4] C+''''
[5] V+ '
[6] →(DN10, DN10, DN6, DN5, DN4, DN3, DN3, DN2, DN1)[(9, 8, 7, 6, 5, 4, 3, 2) \ NP[3]]
[7] DN1: NH+ 'NEWMAN-KEULS'
[8] 'TYPE Q VALUES FOR NEWMAN-KEULS TEST [R=2-' ; N; ]'
[9] →DN7
[10] DN2: NH+ 'DUNCAN', C, 'S NMR TEST'
[11] 'TYPE Q VALUES FOR DUNCAN', C, 'S TEST [R=2-' ; N; ]'
[12] →DN7
[13] DN3: NH+ 'TUKEY', C, 'S HSD(A) TEST'
[14] 'TYPE Q/K VALUE FOR TUKEY', C, 'S HSD TEST [K=' ; N; ]'
[15] →DN7×1 NP[3]=3

```

TABLE 1 (continued)

```
[16] Q+[]
[17] NH+'TUKEY',C,'S HSD(B) TEST'
[18] 'TYPE Q/R VALUES FOR TUKEY',C,'S HSD TEST [R=2-' ;N;']
[19] Q+(Q+[])+2
[20] +DN8
[21] DN4:NH+'FISHER',C,'S LSD TEST'
[22] 'TYPE T VALUE FOR LSD TEST [α/2]'
[23] +DN7
[24] DN5:NH+'DUNN',C,'S PROCEDURE'
[25] 'TYPE T VALUE FOR DUNN',C,'S TEST [C=';(0.5X(NX(N-1))):]'
[26] +DN7
[27] DN6:NH+'SCHEFFE',C,'S S METHOD'
[28] 'TYPE F VALUE FOR SCHEFFE',C,'S TEST [K=' ;N;']
[29] DN7:Q+[]
[30] α(NP[3]ε5,6,7) / 'MS+2×NP[1]'
[31] α(NP[3]=7) / 'Q+(((N-1)×Q)*.5)'
[32] DN8:W+(N,N)ρ(((MS+NP[2])*0.5)×Q)
[33] α(NP[3]ε12) / 'W+(N,N)ρ(0,(((MS+NP[2])*0.5)×Q))'
[34] D+40ρ'-1
[35] T+(T-((QZ)-(Z+(N,N)ρ(ZBAR[4]ZBAR))))>W
[36] 'MULTIPLE COMPARISON: ',NH
[37] D
[38] 'SIGNIFICANT CONTRAST',(10ρV),'DIFFERENCE'
[39] D
[40] UP1:+DN9×1(+/T[I;])=0
[41] G+φ(T[I;]/Y)
[42] UP2:(7ρV),'M',(2 0 √K[I]),'M',(2 0 √G[IA]),(16ρV),(10 3 √(T[I;](IB[IA'])))
[43] +UP2×1(IA+IA+1)≤ρG
[44] +UP1×1(I+I+(IA+1))<N
[45] DN9:D
[46] +0
[47] DN10:NP ANPC ZBAR
```

Fns IV ANPC

```
∇ NP ANPC ZBAR;A;A1;A2;A3;A4;C;Q;CC;QQ;CV;I;IM;Q;M;MTX;MX;N;NC;NH;NP;S;SV;TC;V
[1] NE+(N+ρZBAR)ρNP[2]
[2] S+(I+1)ρ'+
[3] C+'''
[4] V+' '
[5] 'HOW MANY CONTRAST? [TYPE LONGEST CONTRAST FIRST]'
[6] M+((NC+[]),100)ρ(V+SV+' ')
[7] +DN1×1NP[3]=8
[8] NH+'DUNN',C,'S PROCEDURE'
[9] 'TYPE T VALUE FOR DUNN',C,'S PROCEDURE [C=' ;NC;']
[10] +DN2×1(Q+[])>0
[11] DN1:NH+'SCHEFFE',C,'S PROCEDURE'
[12] 'TYPE F VALUE FOR SCHEFFE',C,'S PROCEDURE [K=' ;N;']
[13] Q+(((N-1)×[])*0.5)
[14] DN2:+DN3×1NP[2]≠0
[15] 'TYPE SAMPLE SIZE FOR EACH TREATMENT'
[16] NE+[]
[17] DN3:'TYPE MEAN INDEXES [C-' ;I;']
[18] IM+[]
[19] 'TYPE CONTRAST COEFFICIENTS [C-' ;I;']
[20] CV+Q×((NP[1]×(+/(((CC+αQQ+[])*2)+NP[IM])))*0.5)
[21] α(|(TC+/(CC×(ZBAR[IM])))|>CV) / 'SV+Q'
[22] MX+(((Q,1)ρ'M'),(2 0 √(Q,1)ρIM),((Q,1)ρ(((Q+ρIM)-1)ρ' |'),V))
[23] α(I=1) / 'A3+(A1+ρMX)+(A2+ρQQ)+47'
[24] MTX+(((A1-(ρMX))+5)ρV),MX,(((A2-(ρQQ))+10)ρV),QQ,(7ρV),(10 3 √TC),(13ρV),SV
[25] +DN5×1(ρMTX)>100
[26] M[I;]+(MTX,(100-(ρMTX))ρ(SV+V))
[27] +DN3×1(I+I+1)≤NC
[28] (5ρV),'MULTIPLE COMPARISON: ',NH
[29] ;A-(1,100)ρ((5ρV),((A3)ρ'-'),((100-(A3+5))ρV))
[30] ((A1-6)ρV),'CONTRASTED',((A4+(A2-8)+9)ρV),'CONTRAST',(11ρV),'DIFFERENCE',(6ρV),'SIGNIFICANT'
[31] ((A1-4)ρV),'MEANS',((A4+1)ρV),'COEFFICIENTS',(12ρV),'MEANS',(8ρV),'CONTRASTS[*]'
[32] A,[1](@M),[1] A
[33] +0
[34] DN5:'CHARACTER LINE > 100 CHARACTERS[' ;ρM;']: ADJUST FORMAT'
```

TABLE 2
RBD

XVAR				
67.24	93.3	85.71	94.69	75.36
65.36	93.75	83.93	93.31	77.84
69.84	96.41	80.8	96.43	72.77
66.36	94.86	79.46	92.76	73.66
68.42	93.33	80.66	95.46	74.44
67.27	95.17	84.32	94.89	76.89

RBD XVAR

ANALYSIS OF VARIANCE TABLE
RANDOMIZED BLOCK DESIGN: MIXED MODEL

SOURCE	SS	DF	MS	F
BETWEEN TREATMENTS	3422.16348	4	855.54087	272.920
BETWEEN BLOCKS	16.27739	5	3.25548	1.039
RESIDUAL [BLOCK*TREATMENT]	62.69536	20	3.13477	
TOTAL	3501.13623	29		

ZBAR

67.415 94.47 82.48 94.59 75.16

TABLE 3
POSTHOC

Example 1: a) A posteriori select values, b) Newman-Keuls

POSTHOC

TYPE A POSTERIORI SELECT VALUE [0=LISTING]

□:

0

A POSTERIORI TEST	SELECT VALUE	SAMPLING DISTRIBUTION
NEWMAN-KEULS	1	STUDENTIZED RANGE
DUNCAN'S NEW MULTIPLE RANGE [NMR]	2	DUNCAN'S QR
TUKEY'S HONESTLY SIGNIFICANT DIFFERENCE(A) [HSD(A)]	3	STUDENTIZED RANGE
TUKEY'S HONESTLY SIGNIFICANT DIFFERENCE(B) [HSD(B)]	4	STUDENTIZED RANGE
FISHER'S LEAST SIGNIFICANT DIFFERENCE [LSD]	5	STUDENT'S T
DUNN'S PROCEDURE [BONFERRONI T STATISTIC]	6	DUNN'S T'D
SCHEFFE'S S METHOD	7	F
SCHEFFE'S PROCEDURE [PAIRWISE/NONPAIRWISE]	8	F
DUNN'S PROCEDURE [PAIRWISE/NONPAIRWISE]	9	DUNN'S T'D

TYPE A POSTERIORI SELECT VALUE [0=LISTING]

□:

1

TYPE: 1) MEAN SQUARE ERROR, 2) COMPARISON SAMPLE SIZE

□:

3.135 6

TYPE MEAN VALUES OR VARIABLE MEAN NAME

□:

67.415 94.47 82.48 94.59 75.16

TYPE Q VALUES FOR NEWMAN-KEULS TEST [R=2-5]

□:

4.02 4.64 5.02 5.29

TABLE 3 (continued)

MULTIPLE COMPARISON: NEWMAN-KEULS

SIGNIFICANT CONTRAST	DIFFERENCE
M 1 M 4	27.175
M 1 M 2	27.055
M 1 M 3	15.065
M 1 M 5	7.745
M 5 M 4	19.430
M 5 M 2	19.310
M 5 M 3	7.320
M 3 M 4	12.110
M 3 M 2	11.990

Example 2: a) Variable mean name, b) Duncan's NMR test

```

POSTHOC
TYPE A POSTERIORI SELECT VALUE [0=LISTING]
□:
  2
TYPE: 1) MEAN SQUARE ERROR, 2) COMPARISON SAMPLE SIZE
□:
  3.135 6
TYPE MEAN VALUES OR VARIABLE MEAN NAME
□:
  ZBAR
TYPE Q VALUES FOR DUNCAN'S TEST [R=2-5]
□:
  4.02 4.22 4.33 4.40
    
```

MULTIPLE COMPARISON: DUNCAN'S NMR TEST

SIGNIFICANT CONTRAST	DIFFERENCE
M 1 M 4	27.175
M 1 M 2	27.055
M 1 M 3	15.065
M 1 M 5	7.745
M 5 M 4	19.430
M 5 M 2	19.310
M 5 M 3	7.320
M 3 M 4	12.110
M 3 M 2	11.990

select values and the appropriate table to enter (see Example 1 in Table 3). After the printout, Q-1 is repeated and the desired a posteriori test may be chosen by typing the appropriate select value. Q-2 requests first the appropriate mean square error term (Kirk, 1968), and then the comparison sample size. Only a single value is typed in response to the comparison sample size, either the n corresponding to any given treatment level when ns are equal (a posteriori Tests 1-7) or a zero when the ns are unequal (a posteriori Tests 8-9). A zero response causes the program to generate the necessary Q later. Fns POSTHOC organizes responses Q-1 and Q-2 into vector NP; the first element equals the mean square error term; the second, the comparison sample size; and the third, the a posteriori select value. Vector NP is the left argument of fns APPC and ANPC (below). The final Q (Q-3) generated by fns POSTHOC requests mean values. Either the variable mean name is typed in response to Q-3 (i.e., mean vector has already been formed) or the value of each treatment mean is typed.

The mean vector (ZBAR) is the right argument of fns APPC and ANPC (below). For illustrations of fns POSTHOC, see Table 3.

Function 3: APPC. Only pairwise contrasts on means are calculated within fns APPC. Equal ns are required for all treatment levels. Fns APPC has both left (NP) and right (ZBAR) arguments (above) and can be used alone or in combination with the calling fns POSTHOC. APPC generates a single Q—a request for the value of the test statistic (e.g., Student's t distribution, F distribution, or Dunn's t'D distribution). The degrees of freedom for the statistic is that associated with the appropriate error term. The fns generates as an aid (reminder) in entering the appropriate tables: the values of k, the number of treatments (number of elements in ZBAR); r, the number of steps separating ordered means (range of elements in ZBAR); or c, the number of contrasts $[(k(k-1)) \div 2]$.

Output of fns APPC is a table organized into two columns. The first column gives all the significant

TABLE 4
APPC

Example 1: a) left and right fns arguments, b) Tukey's HSD(A) test

3.135 6 3 APPC 67.415 94.47 82.48 94.59 75.16
TYPE Q/K VALUE FOR TUKEY'S HSD TEST [K=5]

□: 5.29

MULTIPLE COMPARISON: TUKEY'S HSD(A) TEST

SIGNIFICANT CONTRAST	DIFFERENCE
M 1 M 4	27.175
M 1 M 2	27.055
M 1 M 3	15.065
M 1 M 5	7.745
M 5 M 4	19.430
M 5 M 2	19.310
M 5 M 3	7.320
M 3 M 4	12.110
M 3 M 2	11.990

Example 2: a) left and right arguments, b) Tukey's HSD(B) test

3.135 6 4 APPC ZBAR
TYPE Q/K VALUE FOR TUKEY'S HSD TEST [K=5]

□: 5.29
TYPE Q/R VALUES FOR TUKEY'S HSD TEST [R=2-5]
□: 4.02 4.64 5.02 5.29

MULTIPLE COMPARISON: TUKEY'S HSD(B) TEST

SIGNIFICANT CONTRAST	DIFFERENCE
M 1 M 4	27.175
M 1 M 2	27.055
M 1 M 3	15.065
M 1 M 5	7.745
M 5 M 4	19.430
M 5 M 2	19.310
M 5 M 3	7.320
M 3 M 4	12.110
M 3 M 2	11.990

Example 3: a) Fisher's LSD test

3.135 6 5 APPC ZBAR
TYPE T VALUE FOR LSD TEST [$\alpha/2$]

□: 2.845

MULTIPLE COMPARISON: FISHER'S LSD TEST

SIGNIFICANT CONTRAST	DIFFERENCE
M 1 M 4	27.175
M 1 M 2	27.055
M 1 M 3	15.065
M 1 M 5	7.745
M 5 M 4	19.430
M 5 M 2	19.310
M 5 M 3	7.320
M 3 M 4	12.110
M 3 M 2	11.990

Example 4: a) Dunn's procedure

3.135 6 6 APPC ZBAR
TYPE T VALUE FOR DUNN'S TEST [C=10]

□: 3.85

MULTIPLE COMPARISON: DUNN'S PROCEDURE

SIGNIFICANT CONTRAST	DIFFERENCE
M 1 M 4	27.175
M 1 M 2	27.055
M 1 M 3	15.065
M 1 M 5	7.745
M 5 M 4	19.430
M 5 M 2	19.310
M 5 M 3	7.320
M 3 M 4	12.110
M 3 M 2	11.990

Example 5: a) Scheffé's S method

3.135 6 7 APPC ZBAR
TYPE F VALUE FOR SCHEFFE'S TEST [K=5]

□: 4.43

MULTIPLE COMPARISON: SCHEFFE'S S METHOD

SIGNIFICANT CONTRAST	DIFFERENCE
M 1 M 4	27.175
M 1 M 2	27.055
M 1 M 3	15.065
M 1 M 5	7.745
M 5 M 4	19.430
M 5 M 2	19.310
M 5 M 3	7.320
M 3 M 4	12.110
M 3 M 2	11.990

TABLE 5
ANPC

Example 1: a) POSTHOC, b) Scheffe's Procedure

POSTHOC

TYPE A POSTERIORI SELECT VALUE [0=LISTING]

□: 8
TYPE: 1) MEAN SQUARE ERROR, 2) COMPARISON SAMPLE SIZE

□: 3.135 6
TYPE MEAN VALUES OR VARIABLE MEAN NAME

□: ZBAR
HOW MANY CONTRAST? [TYPE LONGEST CONTRAST FIRST]

□: 3
TYPE F VALUE FOR SCHEFFÉ'S PROCEDURE [K=5]

□: 4.43
TYPE MEAN INDEXES [C-1]

□: 1 2 3 4 5
TYPE CONTRAST COEFFICIENTS [C-1]

1.00 .25 .25 .25 .25

TYPE MEAN INDEXES [C-2]

□: 2 3 4 5
TYPE CONTRAST COEFFICIENTS [C-2]

-1.00 .333 .333 .333

TYPE MEAN INDEXES [C-3]

□: 2 4
TYPE CONTRAST COEFFICIENTS [C-3]

-1.00 1.00

MULTIPLE COMPARISON: SCHEFFÉ'S PROCEDURE

CONTRASTED MEANS	CONTRAST COEFFICIENTS	DIFFERENCE MEANS	SIGNIFICANT CONTRASTS[*]
M 2 M 4	-1.00 1.00	.120	
M 2 M 3 M 4 M 5	-1.00 .333 .333 .333	-10.477	*
M 1 M 2 M 3 M 4 M 5	1.00 .25 .25 .25 .25	-19.260	*

Example 2: a) unequal n's, b) Scheffé's Procedure

3.135 0 8 APPC ZBAR

HOW MANY CONTRAST? [TYPE LONGEST CONTRAST FIRST]

□: 3
TYPE F VALUE FOR SCHEFFÉ'S PROCEDURE [K=5]

□: 4.43
TYPE SAMPLE SIZE FOR EACH TREATMENT

□: 5 6 4 5 3
TYPE MEAN INDEXES [C-1]

□: 1 2 3 4 5
TYPE CONTRAST COEFFICIENTS [C-1]

-1.00 .25 .25 .25 .25

TYPE MEAN INDEXES [C-2]

□: 2 3 4 5
TYPE CONTRAST COEFFICIENTS [C-2]

1.00 .333 .333 .333

TYPE MEAN INDEXES [C-3]

□: 2 4
TYPE CONTRAST COEFFICIENTS [C-3]

-1.00 1.00

TABLE 5 (continued)

MULTIPLE COMPARISON: SCHEFFE'S PROCEDURE

CONTRASTED MEANS	CONTRAST COEFFICIENTS	DIFFERENCE MEANS	SIGNIFICANT CONTRASTS[*]
M 2 M 4	1.00 1.00	.120	
M 2 M 3 M 4 M 5	1.00 .333 .333 .333	10.477	*
M 1 M 2 M 3 M 4 M 5	1.00 .25 .25 .25 .25	19.260	*

Example 3: a) Dunn's Procedure

```

3.135 6 9 APPC ZBAR
HOW MANY CONTRAST? [TYPE LONGEST CONTRAST FIRST]
[]:
3
TYPE T VALUE FOR DUNN'S PROCEDURE [C=3]
[]:
2.61
TYPE MEAN INDEXES [C-1]
[]:
1 2 3 4 5
TYPE CONTRAST COEFFICIENTS [C-1]
1.00 .25 .25 .25 .25
TYPE MEAN INDEXES [C-2]
[]:
2 3 4 5
TYPE CONTRAST COEFFICIENTS [C-2]
1.00 .333 .333 .333
TYPE MEAN INDEXES [C-3]
[]:
2 4
TYPE CONTRAST COEFFICIENTS [C-3]
1.00 1.00
    
```

MULTIPLE COMPARISON: DUNN'S PROCEDURE

CONTRASTED MEANS	CONTRAST COEFFICIENTS	DIFFERENCE MEANS	SIGNIFICANT CONTRASTS[*]
M 2 M 4	1.00 1.00	.120	
M 2 M 3 M 4 M 5	1.00 .333 .333 .333	10.477	*
M 1 M 2 M 3 M 4 M 5	1.00 .25 .25 .25 .25	19.260	*

pairwise mean contrasts within vector ZBAR; the second provides the difference between each pair of means. If there are no significant contrasts among all possible paired means, the table is empty. For illustrations of fns APPC, see Table 4.

Function 4: ANPC. Fns ANPC is used when; (1) the a posteriori comparisons are nonpairwise [Scheffé's procedure (Test 8) is most appropriate]; (2) the number of contrasts is small compared to the number of means in the experiment [Dunn's procedure (Test 9) is most appropriate]; and (3) the ns are unequal across treatment levels (either Scheffé's or Dunn's procedure may be used) (Kirk, 1968). Only computational differences exist between the a posteriori tests in fns ANPC and similar tests in fns APPC. These differences exist in order to "handle" unequal ns and user-determined contrast coefficients.

Fns ANPC has the same left (NP) and right (ZBAR) arguments as fns APPC. The only difference is that the select value is not the same (i.e., Scheffé = 8, Dunn = 9). All analyses and outputs are done within fns ANPC.

Thus, fns ANPC can be used independently or in combination with fns POSTHOC.

There are four or five Qs in fns ANPC: (1) Q-1 requests the total number of contrasts desired; (2) Q-2a requests the n for each treatment level given that NP(2) = 0 (user-determined sample size), Q-2b the F (Scheffé's procedure) or t'D (Dunn's procedure) value; (3) Q-3, the indices of the means involved in the contrast; (4) Q-4, the appropriate contrast coefficients. Tables are entered (F or t'D) with degrees of freedom for the appropriate error term and either the number of treatment levels, k (k = total number of means in ZBAR) - 1 for Scheffé's, or the number of contrasts, c (c = number of contrasts specified in Q-1), for Dunn's procedure.

Fns ANPC loops iteratively through Qs, Q-3, and Q-4 until the number of user-selected contrasts is reached. Output, upon loop termination, is immediate. Output is in the form of a table which gives for each contrast: (1) the means involved in the contrast, (2) the respective contrast coefficients, (3) the differences between all means contrasted, and (4) the significance of the con-

trast, if any. Significance for a particular contrast is indicated by an asterisk (*). For illustrations of fns ANPC, see Table 5.

Computer. All functions were developed on an IBM 370/168 under APL/SV.

Note Added in Proof. After submission of the present manuscript, we discovered that the computer facilities at a number of institutions (including our own) have been updated (VS APL). To make the program we have described compatible with VS APL, monadic format and catenation must be used rather than semicolons to intermix character and numeric parts of expressions. A revised and extended version of the program is available from William T. McGowan, Behavioral Pharmacology Laboratory, Department of Psychology, University of South Carolina, Columbia, South Carolina 29208.

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