

Multiple Testing of General Contrasts:
Truncated Closure and the Extended
Shaffer-Royen Method

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Pairwise Comparisons

- ANOVA, $g = 10$ groups, $n = 2$ per group
- Averages: $-1.9695, 2.1520, 5.5915, 0.8695, 4.6315, 0.2780, -1.3210, 0.3490, -7.1530,$ and 3.6220 ; $MSE=1.051976$
- $k = 45$ pairwise comparisons
- Bonferroni-Based
 - Bonferroni
 - Holm
 - Shaffer Method 2
- MaxT-Based
 - Tukey
 - Stepdown MaxT-based
 - Royen

SAS Code

```
proc glimmix data=allpairs;  
  class g; model y=g;  
  lsmeans g/pdiff adjust=bonferroni;  
  lsmeans g/pdiff adjust=bonferroni stepdown(type=free);  
  lsmeans g/pdiff adjust=bonferroni stepdown(type=logical);  
  lsmeans g/pdiff adjust=tukey;  
  lsmeans g/pdiff adjust=simulate stepdown(type=free);  
  lsmeans g/pdiff adjust=simulate stepdown(type=logical);  
run;
```

Assumptions

- $\hat{\boldsymbol{\mu}} \sim N_g(\boldsymbol{\mu}, \sigma^2 \mathbf{V})$, $\nu \hat{\sigma}^2 / \sigma^2 \sim \chi_\nu^2$, independent of $\hat{\boldsymbol{\mu}}$
- \mathbf{V} is known, positive definite
- $H_i : \mathbf{c}'_i \boldsymbol{\mu} = 0$ vs. $H'_i : \mathbf{c}'_i \boldsymbol{\mu} \neq 0$, $i = 1, \dots, k$, for contrast vectors \mathbf{c}_i
- $T_i = |\mathbf{c}'_i \hat{\boldsymbol{\mu}}| / (\hat{\sigma}^2 \mathbf{c}'_i \mathbf{V} \mathbf{c}_i)^{1/2}$

Shaffer 2 Method (JASA 81, 826-831)

- Let $H_{(i)}$ correspond with $T_{(i)} = T_{a_i}$.
- Let $S = \{1, \dots, k\}$. Define
$$\mathbf{S}_i = \{K \subset S \mid a_i \in K \text{ and } H_K \cap H'_{(k)} \cap \dots \cap H'_{(k-i+2)} \neq \emptyset\}.$$
- S2: starting with $i = k$, $H_{(i)}$ is rejected iff $H_{(k)}, \dots, H_{(i+1)}$ are rejected and $p_{(i)} \leq \alpha/k_i$, where
$$k_i = \max\{|K|; K \in \mathbf{S}_i\}.$$
- Adjusted p -values are $\tilde{p}_{(i)}^{S2} = \min\{\max_{j \geq i}(k_j p_{(j)}), 1\}$
- S2 is uniformly more powerful than Holm.

Extended Shaffer 2 Method (JASA 92, 299-306)

- Define $t_{ES}(\alpha, i) = \max\{t(\alpha, K) \mid K \in \mathbf{S}_i\}$, where

$$P_0\{\max_{j \in K} T_j \geq t(\alpha, K)\} = \alpha$$

- Starting with $i = k$, $H_{(i)}$ is rejected iff $H_{(k)}, \dots, H_{(i+1)}$ are rejected and $T_{(i)} \geq t_{ES}(\alpha, i)$.

- Adjusted p -values are

$$\tilde{p}_{(i)}^{ES2} = \max_{j \geq i} \left\{ \max_{K \in \mathbf{S}_j} P_0(\max_{l \in K} T_l \geq t_{(j)}) \right\}$$

- Uniformly more powerful than S2.

Example: All Pairs, $g=5$

i	1	2	3	4	5	6	7	8	9	10
$p(i)$.001	.001	.006	.010	.015	.020	.039	.060	.266	.392
H	H_{13}	H_{15}	H_{34}	H_{12}	H_{45}	H_{23}	H_{14}	H_{25}	H_{24}	H_{35}

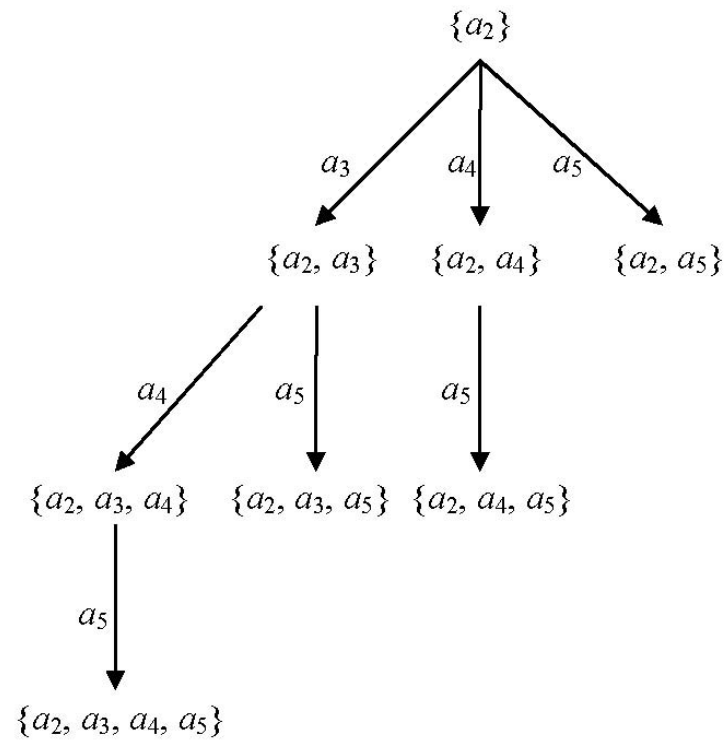
- $\mathbf{S}_1 = \{\{1, 2, 3, 4, 5, 6, 7, 8, 9, 10\}\}$ ($k_1 = 10$)
- $\mathbf{S}_2 = \{\{2, 4, 5, 7, 8, 9\}, \{2, 3, 4, 8\}, \{2, 5, 6, 7\}, \{2, 3, 6, 9\}\}$
($k_2 = 6$)
- $\mathbf{S}_3 = \{\{3, 4, 5, 10\}, \{3, 5, 6, 8, 9, 10\}\}$ ($k_3 = 6$)
- $\mathbf{S}_4 = \{\{4, 7, 9, 10\}, \{4, 5\}\}$ ($k_4 = 4$)
- $\mathbf{S}_5 = \{\{5, 6\}, \{5, 8, 9\}\}$ ($k_5 = 3$)
- $\mathbf{S}_6 = \{\{6, 7, 8, 10\}\}$ ($k_6 = 4$)
- $\mathbf{S}_7 = \{\{7, 8\}, \{7, 10\}\}$ ($k_7 = 2$)
- $\mathbf{S}_8 = \{\{8\}\}$, $\mathbf{S}_9 = \{\{9, 10\}\}$, $\mathbf{S}_{10} = \{\{10\}\}$

S2 and Closure Connection

- Westfall proved FWE control for ES2 directly, not using closure
- Rom and Holland, Hommel and Bernhard showed that S2 is more conservative than full closure with Bonferroni tests for all pairwise comparisons
- Royen noted closed methods are sometimes “nonmonotonic in p -values,” and developed a “truncated closed” procedure: closed testing follows the ordered p -values, stopping as soon as there is insignificance, with application to all pairwise comparisons in ANOVA.

New Results

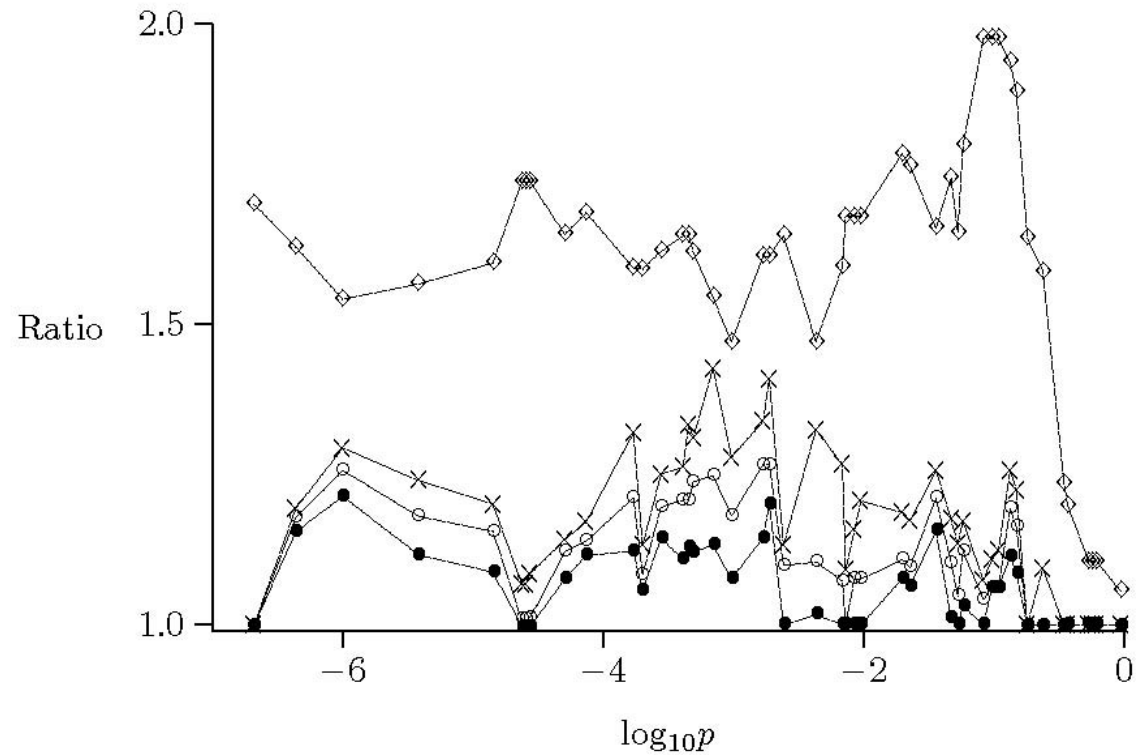
- Westfall and Tobias (a) extend Royen's truncated closure to general contrasts, (b) prove equivalence with ES2, and (c) develop a tree-based branch-and-bound algorithm to
 - prune the all-subsets tree, reducing $O(2^k)$ computational complexity
 - stop the search at early stages if the problem is computationally infeasible, with increasing power for increased search depth.
 - call the method ESR (extended Shaffer-Royen)



Tree representation of the search space

$S_{i+}^{(0)}$ Covering Sets

- $\mathbf{S}_{1+}^{(0)} = \{\{1, 2, 3, 4, 5, 6, 7, 8, 9, 10\}\}$ ($k_{1+}^{(0)} = 10$)
- $\mathbf{S}_{2+}^{(0)} = \{\{2, 3, 4, 5, 6, 7, 8, 9\}\}$ ($k_{2+}^{(0)} = 8$) (9 for free stepdown)
- $\mathbf{S}_{3+}^{(0)} = \{\{3, 4, 5, 6, 8, 9, 10\}\}$ ($k_{3+}^{(0)} = 7$) (8 " " " ")
- $\mathbf{S}_{4+}^{(0)} = \{\{4, 5, 7, 9, 10\}\}$ ($k_{4+}^{(0)} = 5$) (7 " " " ")
- $\mathbf{S}_{5+}^{(0)} = \{\{5, 6, 8, 9\}\}$ ($k_{5+}^{(0)} = 4$)
- $\mathbf{S}_{6+}^{(0)} = \{\{6, 7, 8, 10\}\}$ ($k_{6+}^{(0)} = 4$)
- $\mathbf{S}_{7+}^{(0)} = \{\{7, 8, 10\}\}$ ($k_{7+}^{(0)} = 3$)
- $\mathbf{S}_{8+}^{(0)} = \{\{8\}\}$, $\mathbf{S}_{1+}^{(0)} = \{\{9, 10\}\}$, $\mathbf{S}_{1+}^{(0)} = \{\{10\}\}$



Ratios of adjusted p -values of S_2 (\diamond), $S_{i+}^{(0)}$ (\times), $S_{i+}^{(1)}$ (\circ), and $S_{i+}^{(2)}$ (\bullet) to truncated closure for $g = 10$, $k = 45$ example.

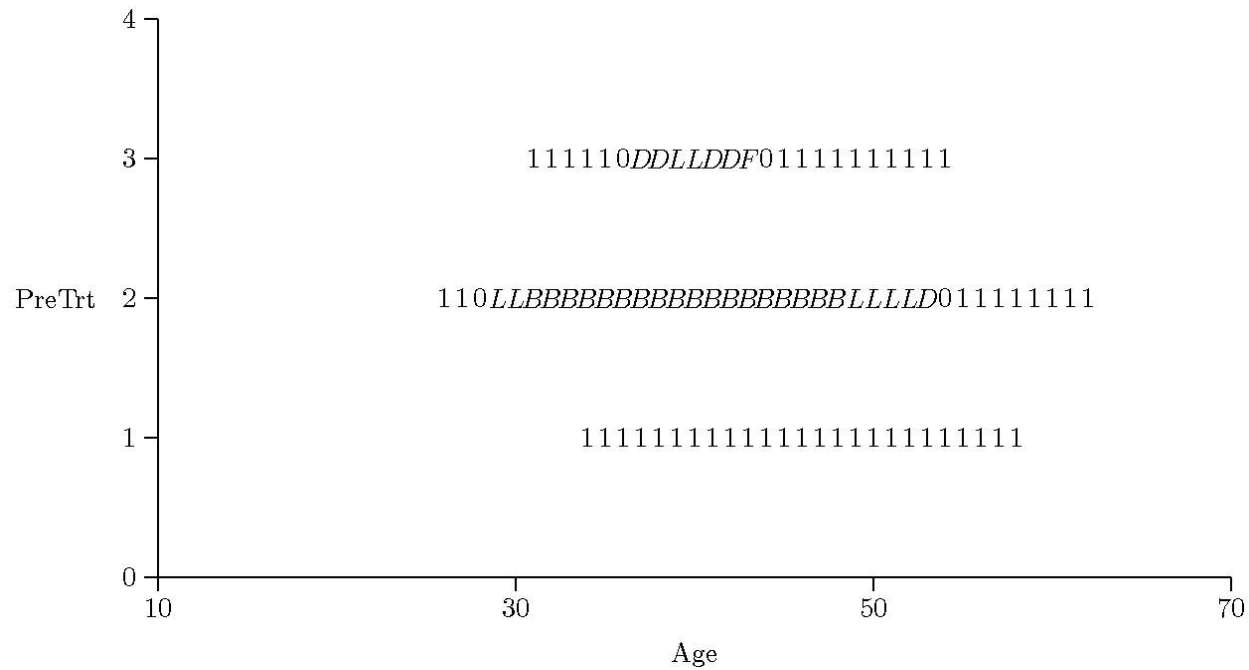
Response Surface Comparisons

- Compare health for Active and Placebo at various initial health (PreTrt) and Age values. Estimates:

$$\hat{y}_A = 1.96204 + 0.42977 \cdot \text{PreTrt} - 0.05410 \cdot \text{Age}/100,$$

$$\hat{y}_P = 1.54004 + 0.53557 \cdot \text{PreTrt} - 1.73130 \cdot \text{Age}/100.$$

- Comparisons made at $(\text{PreTrt}, \text{Age}) \in \{\{1, 2, 3, 4, 5\} \times \{10, 11, \dots, 70\}\}$ ($k = 5 \times 61 = 305$).
- B & B search finds every element of \mathbf{S}_i despite $k = 305$.
- SAS GLIMMIX with CONTRAST statement used for analysis



Significant (0.05 level simultaneous) differences using Benjamini-Hochberg FDR-controlling method (*B*), Liu et al.'s method (*L*), the discrete grid approximation to Liu et al. (*D*), free combination covering sets (*F*), $S_{i+}^{(0)}$, and $S_{i+}^{(1)}$ ($= S_i$ here).

Timing Issues

- 26 minutes to find all relevant subsets in the response surface example (not $O(2^{305})$)
- Full method feasible for all pairwise comparisons with $g = 11$ ($k = 55$), $\mathbf{S}_{i+}^{(2)}$ is feasible with as many as 19 groups ($k = 171$), $\mathbf{S}_{i+}^{(1)}$ is feasible with as many as 32 groups ($k = 496$), and $\mathbf{S}_{i+}^{(0)}$ is feasible with as many as 80 groups ($k = 3160$).
- Donoghue developed an algorithm for S2 that is faster, but limited.

Conclusions

- ESR is a truncated closed testing procedure
- Branch-and-bound provides conservative, but computable approximations, that are usually much more powerful than standard alternatives.
- The method is fully computable in response surface comparisons, even with large numbers of comparisons.
- Available in PROC GLIMMIX for arbitrary contrasts and general model structures (approximate analyses used for more general models)
- General strategy: choose depth as large as compute resources allow and use the resulting multiplicity adjustment.

References

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