Group Sequential Tests with Data-Dependent Treatment Allocation

Chris Jennison,

Department of Mathematical Sciences, University of Bath, UK

and

Bruce Turnbull,

Department of Statistical Science, Cornell University, Ithaca, NY

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http://www.bath.ac.uk/~mascj for slides and full references

The Selection Problem

For each "population" or "treatment" $i = 1, \ldots, k$,

$$X_{i1}, X_{i2}, \ldots \sim N(\theta_i, \sigma^2),$$
 i.i.d.

Aim: To select the population i with the largest mean θ_i .

Method to include:

- Group sequential comparisons
 - early elimination of weak treatments.
- Response-dependent treatment allocation
 - fewer observations on inferior treatments,
 - lower total sample size.

Earlier work

Paulson (Ann. Math. Statist., 1964)

Elimination procedures based on *continuous* sequential comparisons

of 2 populations at a time.

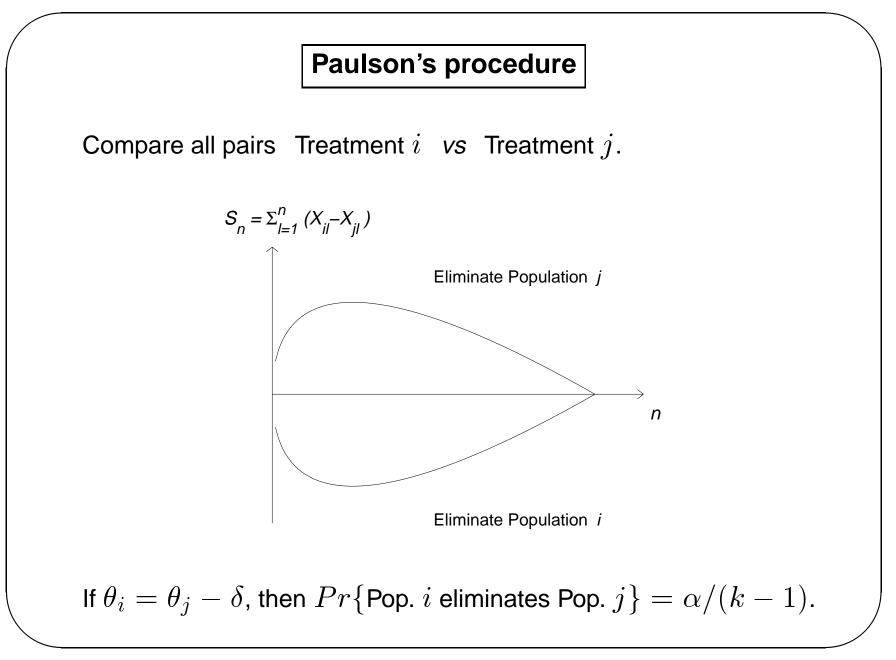
Robbins and Siegmund (JASA, 1974)

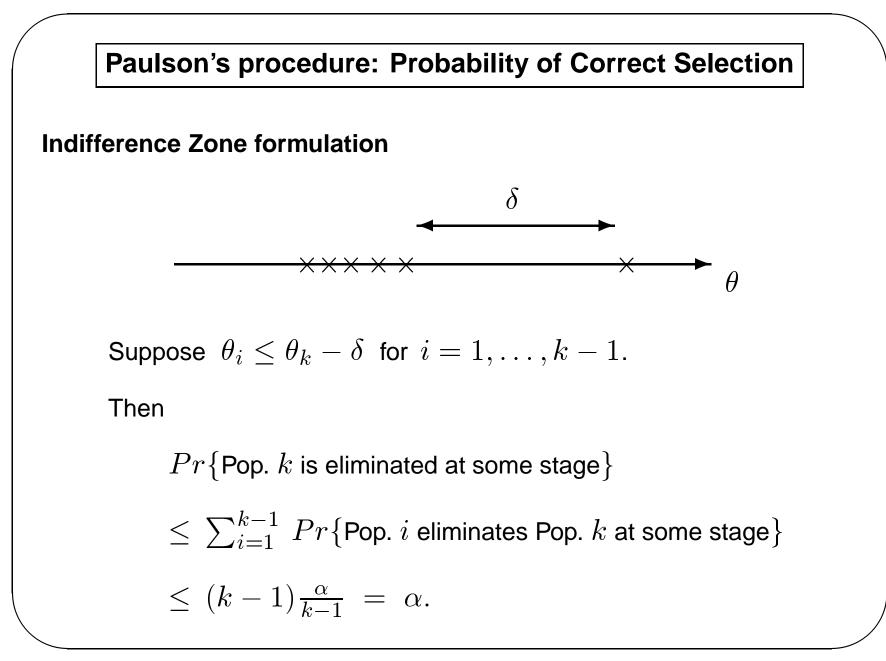
Adaptive sampling for a 2 population comparison with continuous monitoring.

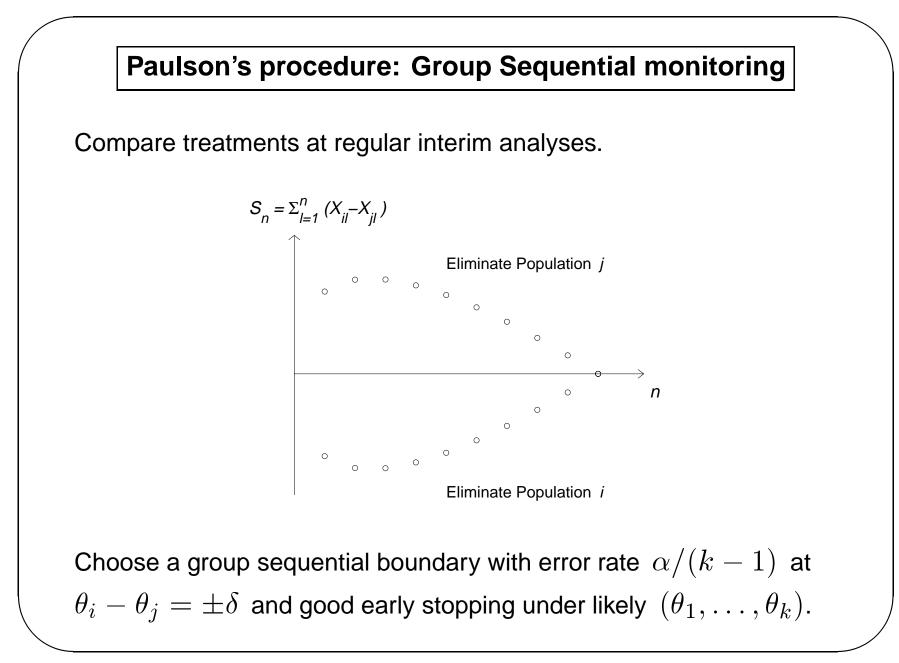
Jennison, Johnstone and Turnbull (*Purdue Symposium*, 1982)

Combining the above.

Update: To take advantage of group sequential tests, error spending, modern computation.







Adaptive Sampling in Paulson's procedure

Motivation

Observations on the leading population are used in k-1 comparisons.

Allocating more observations to the leader can

- Reduce total sample size
- Reduce observations on inferior treatments
 - ethical for medical studies
 - we learn more about better treatments.

Need:

Theory to support adaptive sampling in each pair-wise comparison.

Adaptive Sampling in a Group Sequential Test

Jennison and Turnbull (*Sequential Analysis*, 2001)

For a 2-treatment comparison with

$$X_{1i} \sim N(\theta_1, \, \sigma^2) \quad i = 1, 2, \, \dots \, ,$$

$$X_{2i} \sim N(\theta_2, \, \sigma^2) \quad i = 1, 2, \, \dots$$

At analysis m out of M, with n_{1m} observations on population 1 and n_{2m} on population 2,

$$\begin{split} \widehat{\theta}_1(m) - \widehat{\theta}_2(m) &= \bar{X}_1(m) - \bar{X}_2(m) \ \sim \ N(\theta_1 - \theta_2, \ \sigma^2(\frac{1}{n_{1m}} + \frac{1}{n_{2m}})) \\ &\sim \ N(\theta_1 - \theta_2, \ \mathcal{I}^{-1}(m)), \quad \text{say.} \end{split}$$

Adaptive Sampling continued

The score statistic

$$S(m) = \mathcal{I}(m)(\widehat{\theta}_1(m) - \widehat{\theta}_2(m)) \sim N((\theta_1 - \theta_2)\mathcal{I}(m), \mathcal{I}(m)).$$

Without adaptive sampling, $\{S(1), S(2), \ldots\}$ is distributed as a Brownian motion with drift $\theta_1 - \theta_2$ observed at $\mathcal{I}(1), \mathcal{I}(2), \ldots$

This remains true if group sizes $n_{1m} - n_{1,m-1}$ and $n_{2m} - n_{2,m-1}$ depend on $\hat{\theta}_1(m-1) - \hat{\theta}_2(m-1)$ — but sampling *cannot* depend more generally on $(\hat{\theta}_1(m-1), \hat{\theta}_2(m-1))$.

Theory generalises to normal linear models containing θ_1 and θ_2 .

This extends Robbins and Siegmund (1974) to the group sequential case.

Problem 1

With $k \geq 3$, interesting sampling rules do not satisfy

"m th group sizes for populations 1 and 2 depend only on $\widehat{ heta}_1(m-1) - \widehat{ heta}_2(m-1)$ ".

Solution

- Fix sampling ratios at the start of each group,
- estimate $\theta_i \theta_j$ within each group of data,
- combine estimates with weights $\,\propto\,$ variance $^{-1}$.

This equates to fitting a linear model with additive "stage" effects — recommended in medical studies to avoid bias from time trends.

JT (2001) assess performance of 2-treatment tests:

With stage effects in the model, one cannot compensate later on for sub-optimal sampling ratios in early stages. Savings in Inferior Treatment Numbers are reduced by about a half.

- Fitting stage effects to avoid bias from a time trend is reasonable.
- If modelling such a trend is not really necessary, data are being used inefficiently
 - ethically dubious for medical studies
 - at best pragmatic in other applications.

Problem 2

Information levels for comparing populations i and j

 $\mathcal{I}_{ij}(1), \mathcal{I}_{ij}(2), \mathcal{I}_{ij}(3), \ldots,$

depend on the sampling rule, which involves $S_{ij}(1), S_{ij}(2), \ldots$

Standard group sequential designs, including error spending tests, do not allow such a dependence.

Solution A

Reported studies of such "data-dependent analysis times" show only minor effects on error probabilities — trust these studies and ignore the problem!

Solution B

Recent designs which "adapt" to observed data offer a precise solution:

Denne (Statistics in Medicine, 2001),

Müller and Schäfer (*Biometrics*, 2001).

Procedure

- Set up an error spending test for anticipated $\{\mathcal{I}_1, \mathcal{I}_2, \dots\}$
- \bullet Recursively for $\,m=1,2,\ldots$,
 - At analysis m, compute conditional error probabilities given S(m)
 - Run stages m+1 to M as an error spending test with this conditional error.

A sampling rule (JJT, 1982)

In comparing N-1 populations with a control, the most efficient allocation is

 $\sqrt{N-1}$ observations on the control to

 $1\,$ observation on each other population.

Adaptive rule:

At stage m, with N_m non-eliminated populations, sample

 $\sqrt{N_m-1}$ observations on the leading population to

 $1\,$ observation on each other population.

An updated procedure

Eliminate populations using Paulson's pair-wise comparisons.

Run these comparisons as error spending group sequential tests.

a) Base tests on overall population means (cf JJT, 1982)

Sample in stage m to achieve ratios $\sqrt{N_m}$: 1 : ... : 1 of total observations on the N_m surviving populations.

b) Combine stage-wise estimates of each $\theta_i - \theta_j$

Sample in ratios $\sqrt{N_m}$: 1 : ... : 1 within stage m.

Problem 1 is dealt with properly in (b); Problem 2 is ignored (Solution A!)

Modern applications: Medical

Aim: Combine phase II and phase III clinical trials by running a single study to select a treatment (e.g., dose level) and compare this treatment with a control.

References:

Thall, Simon and Ellenberg (*Bmka*, 1988)
Schaid, Wieand and Therneau (*Bmka*, 1990)
Proschan, Follmann and Geller (*Statist. in Med.*, 1994)
Stallard and Todd (*Univ. Reading Technical Report*, 1999)

The need to compare with a control treatment changes the problem.

Sequential elimination offers scope for improvement on proposed designs

— but there are computational complications.

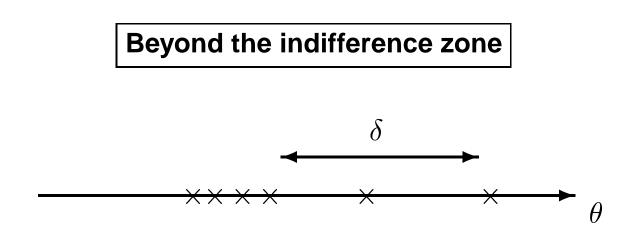
Modern applications: Computer learning

Aim: To select and rank algorithms for their performance on randomly chosen problems.

Experimentation is on a computer, hence any well-defined procedure is easily implemented.

In selecting the best s out of k competitors, k can be large, e.g., 10,000. Clearly, sequential methods are desirable — or, with parallel processing, group sequential methods.

Computer scientists are discovering the selection and ranking literature!



What if there is a θ_i within δ of the highest θ_j ?

It should be OK to select a population within δ of the best. But can a non-optimal population eliminate the best, then be eliminated itself?

Kao and Lai (*Comm. Statist. Th. Meth.*, 1980) provide a solution, raising the boundary for any pair-wise elimination before the final decision.

This method works for Paulson's procedure with adaptive sampling and can be extended to choosing the best s populations out of k.

Conclusions

- Solutions found in 1982 are still very appropriate.
- Applications for selection procedures are appearing in important research areas.
- Interesting challenges remain.

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