

Screening Strategies in the Presence of Interactions – comparison of strategies –

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Joint work with

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- Screening Strategies in the Presence of Interactions
- Screening designs
 - Two-stage group screening
 - Single stage supersaturated design
- Analysis methods
 - Dantzig Selector, LASSO, SCAD,
 - Bayesian Model Selection, Bayesian MAP estimation
- Simulation study comparing methods
- Issues for simulations

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has a *substantial effect* on the response average or response variability through main effects or interactions

- It does not mean “has a non-zero effect”
- There needs to be a cost-effective benefit to changing current product design

Screening

- Assumes factor sparsity

Screening

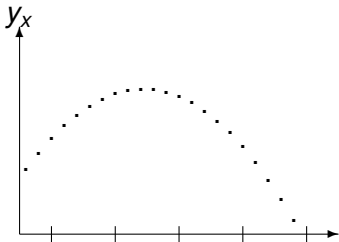
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Screening Experiment: Cold Start Optimization

- Vine, Lewis, Dean, and Brunson, (2008), *Technometrics*.



Aim of experiment:

- To find which factors most affect the cold start performance of the new generation Jaguar engine

Screening Experiment: Cold Start Optimization

Factor List

40 factors

- Afr
- Air Assisted Injection
- Altitude
- Ambient Temperature
- Back Pressure
- Calibration
- Carbon Deposits
- Combustion Chamber Contamination
- Combustion/start History
- Cranking Speed
- Early Entry Into Fuel Cutoff
- Effective Compression Ratio
- Engine Age

- 40 factors identified and assessed
- 10 factors in a screening experiment
- Investigated 10 main effects, 45 2-factor interactions

Two-stage approach

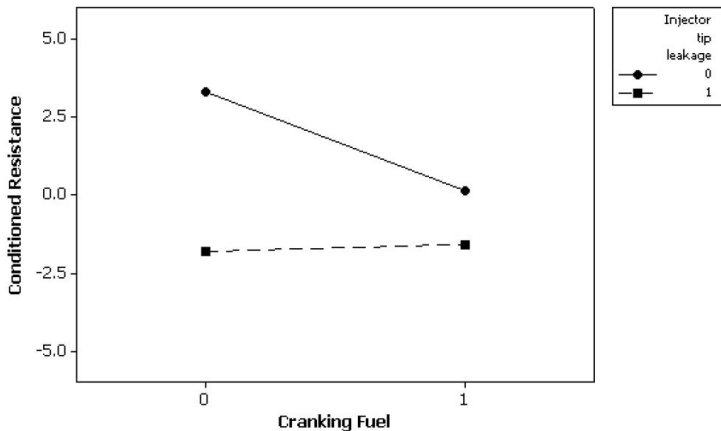
- Run a small experiment to estimate only main effects
- Select factors with large (active) main effects
- Use follow-up experimentation to investigate interactions
- Strategy works well if interactions are small or occur only between factors with large main effects
 - (*strong effect heredity*);
 - Hamada & Wu (1992); Chipman (1996)

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 - (*strong effect heredity*);
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- In the previous example, this approach would not have worked

Interaction Illustration: Jaguar experiment

(e)



- Main effect of cranking fuel is small
- Would be screened out at stage 1

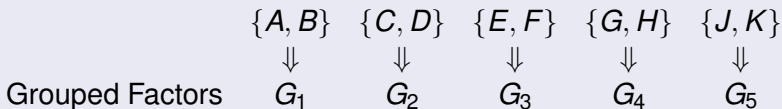
- ⇒ Important to measure interactions at stage 1
- Other studies stressing importance of screening interactions
 - Phoa, Wong, and Xu, 2009
 - Scinto, Wilkinson, and Lin, 2011

Screening Strategies

for investigating interactions

- Two-stage group screening
- Single-stage supersaturated design

I. Two-stage Group Screening



Stage 1: perform an experiment on the **grouped factors**

- Factors within a group always have matching levels

Stage 2: dismantle active groups and experiment on their **individual factors**

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Stage 1: grouped factors

- Estimate main effects and interactions
- Grouped factors with “large” effects are declared active

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Stage 2: individual factors within a group

If group factor

- was inactive, hold individual factors at nominal levels
- was active, measure main effects and two-factor interactions among factors in the group

If two grouped factors

- had active interaction, measure interactions between individual factors in these groups

I. Two-stage Group Screening

- Aliasing among main effects and two-factor interactions is induced by the grouping
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$$\text{mean} = AB = CD = EF = GH = JK$$

$G_i \Rightarrow$	$A = B$	$C = D$	$E = F$	$G = H$	$J = K$		
$G_1 G_2 \Rightarrow$	AC	$=$	BD	$=$	AD	$=$	BC
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- each set of aliased two-factor interactions involves factors from within only two groups
- This remains true even if hundreds of factors
- and no matter the group size

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In a **supersaturated design**:

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- there are fewer runs than effects to be estimated
- no main effects or interactions are completely aliased, but estimates of some main effects and interactions are correlated
- For good results, need design column correlations ≤ 0.33
- There are many different construction methods in the literature
- Or can use algorithm of Jones, Lin, & Nachtsheim (2008) to find best (*D*-optimal) SSD for estimating a given set of factorial effects

Analysis methods

- Single-stage Supersaturated Design (SSD)
 - **SCAD**: Smoothly Clipped Absolute Deviation
 - **Lasso**: Least Absolute Shrinkage and Selection Operator
 - **DS**: Gauss-Dantzig Selector
 - **Bayes**: Bayesian Model Selection
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- Two-stage Group Screening
 - **GS5**: Group Screening with R^2
 - **GS5-DS**: Group Screening with Dantzig Selector
 - **GS5-B**: Group Screening with Bayesian Model Selection
 - **GS5-B-MAP**: Grp Screening with Bayesian MAP estimation

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Small effects less than a threshold t screened out

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- 2 Select effects whose regression coefficients have *maximum a posteriori* (MAP) estimates $> t$
 - tuning parameters in the prior distributions chosen via graphical methods
 - for group screening, we developed prior distributions at stage 2 based on the posterior distributions at stage 1

Simulation Study Details

to compare designs and analysis methods

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Number of factors f equal to 10 or 15 or 20

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I. Group screening used

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II. Single stage supersaturated design

- generated from alg. of Jones, Lin, and Nachtsheim (2008)
- size = median number of runs required by Group Screening
 - For $f = 10, p = 56$ effects, $n = 32$.
 - For $f = 15, p = 121$ effects, $n = 58$.
 - For $f = 20, p = 211$ effects, $n = 94$.

Simulation: effect size generation

- Every **main effect** is active with probability q_{me}
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Simulation: effect size generation

- Every **main effect** is active with probability q_{me}
 - ($q_{me} \in \{.05, .1, .15, .2\}$)
- A two-factor **interaction** is active with conditional probability
 - $w_{00} = 0.005$; $w_{10} = w_{01} = 0.125$; $w_{11} = 0.25$
- Generated **active** effect sizes from $N(6, 1)$, $N(12, 4)$, $N(24, 4)$
- Generated **inactive** effect sizes from $N(0, 1)$ or $N(0, 16)$

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- Given the generated effect sizes β
- and the model $y = X\beta + \epsilon$
- generated **data** y for given design X with error ϵ from $N(0, 1)$
- Analyzed the data using each of the specified methods
- Repeated a total of 1,000 times for each setting

Simulation: performance measures

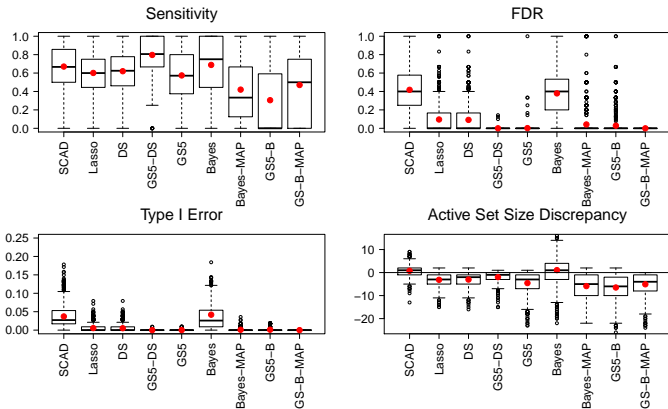
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- **Sensitivity**: proportion of active effects identified as active
- **Type I error**: proportion of inactive effects identified as active
- **False discovery rate**: proportion of effects identified as active that are truly inactive
- **Active Set Size Discrepancy**:
true # active effects – declared # active effects

Comparisons of designs and analyses; $f=15$ factors

[$q_{me} = 0.20$, Active $N(24, 4)$, Inactive $N(0, 16)$]



3b.pdf

Comparisons of designs and analyses—summary

In general:

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Comparisons of designs and analyses—summary

In general:

- **GS5-DS** seems to perform the **best overall**
- followed by **Lasso and DS** in the supersaturated design
- **SCAD** in the ssd tended to have **high FDR**
- **GS5** (using R^2) tended to have **low sensitivity**
- Bayesian model selection in ssd sometimes had very high Type I error rate
- Bayesian methods had a greater drop in performance for the difficult cases than the frequentist methods

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- In a simulation, assign active effects at random to columns, and compute summary measures over a large number of such assignments.

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 - We used n for SSD size = median number of runs for group screening procedure
 - Possible to match more closely and compare performance on each individual run of the simulation.

- For comparison, the sizes of group screening and supersaturated designs need to match
 - We used n for SSD size = median number of runs for group screening procedure
 - Possible to match more closely and compare performance on each individual run of the simulation.
- In a simulation study, automatic decisions have to be made at stage 1 of group screening about active groups.
 - More control can be exercised in a practical experiment.

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- In group screening, aliased small effects may amalgamate, resulting in a non-active group appearing to be active at stage 1. However, a high FDR can be corrected at stage 2
- Such amalgamation can also happen in any fractional factorial experiment

- Active main effects and interaction effects could be drawn from different distributions.

Future extensions

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- To mimic a situation where effects can be classified in terms of their likelihood of being active, the active effect distributions could be set differently
- We used the same threshold t for screening out small effects regardless of whether the effect was a main effect or interaction. Could be set differently.
- The Bayesian approaches are more computationally intensive than the frequentist methods. Need to refine the procedures so that larger numbers of factors can be handled
- Need to tune Bayes procedures to be more successful overall

Thank you!