

## Frequentist Analysis Strategy for Complex Aliasing

- Some properties of Method II (cont.)
  - The constraint imposed by effect heredity substantially reduces the model space searched in Method II.
  - \* Consider, the 12-run PBD with 11 factors and  $h=4$  in Method II.
  - \* The total number of main effects and two-factor interactions is  $11 + \binom{11}{2} = 66$ .
  - \* The total number of models with five terms (and the intercept) is  $\binom{66}{4} = 720,720$ .
  - \* There are 15,510 models satisfying the effect heredity requirement. This is about 2.2% of the total number of models.
  - As the number of factors and  $h$  increase, the search will become computationally prohibitive. An alternative is to use an efficient stochastic search such as the *Bayesian variable selection*.

❖ Reading: textbook, 9.4

## Analysis of Cast Fatigue Experiment

- The design is an  $OA(12, 2^7)$ .
- Consider the main-effect-only model.
  - The 11 columns in Table 1 are mutually orthogonal.
  - The main effect estimates are uncorrelated and unbiased estimates of the main effects *if there are no interactions*.
  - Half-normal plot:

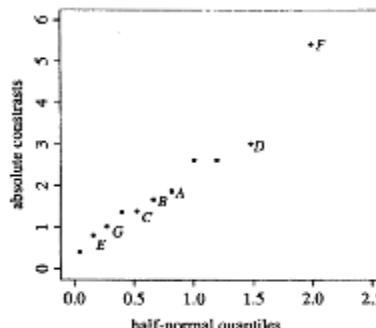


Figure 1: Half-normal plot, cast fatigue experiment

- \* The model with  $F$  alone has an  $R^2 = 0.45$ .
- \* The model  $(F, D)$  has an  $R^2 = 0.59$ , and the fitted model is

$$\hat{y} = 5.73 + 0.458F - 0.258D.$$

## Analysis of Cast Fatigue Experiment

- Entertain a model with ME  $F$  and all the interactions involving  $F$ .
  - Identify a significant  $FG$  interaction.
  - The model  $(F, FG)$  has an  $R^2 = 0.89$
  - The model  $(D, F, FG)$  has an  $R^2 = 0.92 \Rightarrow D$  appears not significant
  - The model for predicted fatigue lifetimes is

$$\hat{y} = 5.7 + 0.458F - 0.459FG,$$

- \* Set  $(F, G) = (+, -)$ . The predicted life is

$$5.7 + 0.458 - 0.459(-1) = 5.7 + 0.92 = 6.62.$$

- \* Compared to the average lifetimes ( $=5.7$ ), it has a 16% increase ( $=0.92/5.7$ )
- \* Compared to  $(F, G) = (-, -)$ , it has a 38% increase ( $=1.84/4.78$ )
- The potential for a dramatic improvement would not have been possible without discovering the  $FG$  interaction.

## Analysis of Cast Fatigue Experiment

- Using Method I, the model  $(F, FG)$  is found in Step 1 and does not change in further steps.
- Method II also identifies model  $(F, FG)$  using  $h = 2$ .
- For larger  $h$  (say, 3 or 4), Method II identifies multiple good fitting models, *each of which contains  $F$  and  $FG$* .
- The original analysis used main-effect-only model and
  - identified the  $F$  and  $D$  as significant,
  - noted a discrepancy between the findings and previous work, i.e., the sign of  $D$  effect was reversed,
  - concluded that the possible cause was an interaction  $DE$ ,
  - claimed that the design did not generate enough information to determine  $DE$ .
- Because the  $DE$  interaction is orthogonal to both  $D$  and  $E$  main effects, these three effect estimates are uncorrelated and therefore the  $DE$  interaction does not affect the sign of the factor  $D$  main effect.

## Analysis of Cast Fatigue Experiment

- The design's aliasing patterns can be used to explain this apparent reversal.

Table 9: Estimates and Alias Patterns, Cast Fatigue Experiment

Effect	Estimated Effect	Alias Pattern
<i>A</i>	0.326	$A - \frac{1}{3}FG$
<i>B</i>	0.294	$B - \frac{1}{3}FG$
<i>C</i>	-0.246	$C + \frac{1}{3}FG$
<i>D</i>	-0.516	$D + \frac{1}{3}FG$
<i>E</i>	0.150	$E - \frac{1}{3}FG$
<i>F</i>	0.915	<i>F</i>
<i>G</i>	0.183	<i>G</i>
8	0.446	$- \frac{1}{3}FG$
9	0.453	$- \frac{1}{3}FG$
10	0.081	$- \frac{1}{3}FG$
11	-0.242	$+ \frac{1}{3}FG$

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## Analysis of Cast Fatigue Experiment

- The *D* estimate can be adjusted as follows:

$$\hat{D} - \frac{1}{3}\widehat{FG} = -0.516 - (-0.306) = -0.210.$$

- A 95% confidence interval for *D*,  $(-0.526, 0.106) = (-0.210 \pm 0.316)$ , shows that a positive *D* main effect is possible.
- The magnitude of the estimated effects of *A*, *B* and *C* can be explained from Table 9 by an *FG* interaction; i.e.,

$$(-\frac{1}{3}\widehat{FG}, -\frac{1}{3}\widehat{FG}, \frac{1}{3}\widehat{FG}) = (0.306, 0.306, -0.306)$$

are close to their estimates, 0.326, 0.294 and -0.246, respectively.

- The effects *A-E* and 8-11 have the same sign as their *FG* aliases, which lend further support to the existence of a significant *FG* interaction.

## Bayesian Analysis Strategy for Complex Aliasing

- The Methods I and II work well if
  - the models under search are restricted to have a small number of terms, and
  - the number of candidated terms to choose from is moderate.
- As the number of candidate terms to choose from and/or the number of terms allowed in the model increases,
  - the search over the model space is too incomplete for Method I to be effective, and
  - the required computations for Method II become prohibitively expensive even with the help of effect heredity to reduce the number of models.

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JOHNSON (DODGE) (JHU, WILMINGTON, USA) and S. W. CHONG (NTHU, Taiwan)

## Bayesian Analysis Strategy for Complex Aliasing

- Bayesian framework:
  - Let  $\mathbf{y}$  denote the data, and  $\boldsymbol{\theta}$  denote the the vector of parameters. Both are random variables.
  - $f(\mathbf{y}|\boldsymbol{\theta})$ : the likelihood of  $\mathbf{y}$  given  $\boldsymbol{\theta}$
  - $\pi(\boldsymbol{\theta})$ : the prior distribution for  $\boldsymbol{\theta}$ .
  - The posterior distribution for  $\boldsymbol{\theta}$  is:

$$\pi(\boldsymbol{\theta}|\mathbf{y}) = f(\mathbf{y}|\boldsymbol{\theta})\pi(\boldsymbol{\theta}) / \int f(\mathbf{y}|\boldsymbol{\theta})\pi(\boldsymbol{\theta})d\boldsymbol{\theta}$$

- Using the posterior distribution, inference about  $\boldsymbol{\theta}$  can then be made.
- Parameters  $\boldsymbol{\theta} = (\boldsymbol{\beta}, \boldsymbol{\delta}, \sigma^2)$ 
  - $\boldsymbol{\beta}$ : a  $(k+1)$  vector containing the intercept and factorial effects,
  - $\boldsymbol{\delta}$ : a  $(k+1)$  vector of 0's and 1's indicating the significance of the effects
  - $\sigma^2$ : error variance in the linear model

# Bayesian Analysis Strategy for Complex Aliasing

- Assignment of distributions

- $\mathbf{y} | \beta, \delta, \sigma^2$ : a linear model structure

$$\mathbf{y} = \mathbf{X}\beta + \boldsymbol{\epsilon},$$

where  $\mathbf{X}$  is the  $N \times (k + 1)$  model matrix, and  $\boldsymbol{\epsilon} \sim MN(\mathbf{0}, \sigma^2 \mathbf{I}_{N \times N})$ .

- $\beta | \delta, \sigma^2$ : normal mixture prior

$$\pi(\beta_i | \delta_i, \sigma^2) = \begin{cases} N(0, \sigma^2 \tau_i^2) & \text{if } \delta_i = 0 \\ N(0, \sigma^2 (c_i \tau_i)^2) & \text{if } \delta_i = 1 \end{cases}.$$

- \* When  $\delta_i = 0$ , the constant  $\tau_i$  needs to be specified so that  $\beta_i$  is tightly centered around 0 and therefore does not have a large effect.
- \* The constant  $c_i$  needs to be chosen with  $c_i \gg 1$  to indicate the possibility of a large  $\beta_i$  when  $\delta_i = 1$ .
- \* The constants  $\tau_i$  and  $c_i$  should be chosen to represent respectively a “small” effect, and how many times larger a “large” effect should be.

# Bayesian Analysis Strategy for Complex Aliasing

- Assignment of distributions (cont.)

- The  $\delta$  and  $\sigma^2$  are independent.

- $\delta$

- \* Independence prior:  $\pi(\delta) = \prod_{i=1}^{p+1} p_i^{\delta_i} (1 - p_i)^{1 - \delta_i}$ , where

$$p_i = \text{Prob}(\delta_i = 1).$$

- \* Prior with effect heredity principle

- Consider the example  $\delta = (\delta_A, \delta_B, \delta_C, \delta_{AB}, \delta_{AC}, \delta_{BC})$

- $\text{Prob}(\delta) = \text{Prob}(\delta_A) \text{Prob}(\delta_B) \text{Prob}(\delta_C)$

$$\times \text{Prob}(\delta_{AB} | \delta_A, \delta_B) \text{Prob}(\delta_{AC} | \delta_A, \delta_C) \text{Prob}(\delta_{BC} | \delta_B, \delta_C).$$

$$\text{Prob}(\delta_{AB} = 1 | \delta_A, \delta_B) = \begin{cases} p_{00} & \text{if } (\delta_A, \delta_B) = (0, 0) \\ p_{01} & \text{if } (\delta_A, \delta_B) = (0, 1) \\ p_{10} & \text{if } (\delta_A, \delta_B) = (1, 0) \\ p_{11} & \text{if } (\delta_A, \delta_B) = (1, 1) \end{cases}.$$

## Bayesian Analysis Strategy for Complex Aliasing

- Assignment of distributions (cont.)
  - $\delta$  (cont.)
    - Some notes for  $Prob(\delta)$ :
      - Independence is assumed among the  $\delta_i$ 's for main effects
      - Conditional independence principle: conditional on  $\delta_i$ 's for main effects, the  $\delta_i$ 's for interactions are independent
      - Inheritance principle: the significance of a term depends only on those terms from which it was formed
    - Some notes for  $Prob(\delta_{AB})$ :
      - We should choose  $p_{00}$  small (e.g., 0.01),  $p_{01}$  and  $p_{10}$  larger (e.g., 0.10) and  $p_{11}$  largest (e.g., 0.25)  $\Rightarrow$  **relaxed weak heredity**
      - Strict weak heredity**: setting  $p_{00} = 0$
      - Strong heredity**: setting  $p_{00}=p_{01}=p_{10}=0$
      - Probabilities of less than 0.5 for both main effects and interactions represent the belief that relatively few terms are active, i.e., *effect sparsity* holds.

## Bayesian Analysis Strategy for Complex Aliasing

- Assignment of distributions (cont.)
  - $\sigma^2$ : inverse gamma
 
$$\sigma^2 \sim \text{IG}(\nu/2, \nu\lambda/2),$$
 whose density is
 
$$\pi(\sigma^2) \propto (\sigma^2)^{-(\nu/2+1/2)} \exp\{-\nu\lambda/(2\sigma^2)\}.$$
- The evaluation of the posterior for  $\theta$  can be implemented by using **Gibbs sampling**, a simple *Markov Chain Monte Carlo* (MCMC) technique for drawing samples from a posterior distribution (read textbook, 9.5.2, for details).
- Because  $\delta$  specifies a model by the  $i$ 's with  $\delta_i = 1$ , the posterior for  $\delta$  is of particular interest.
- Choice of prior tuning constants  $\tau, c, \nu, \lambda$ : see textbook, 9.5.3, for details.