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MORS Symposium  
 16-19 June 2014, Hilton Mark Center, Alexandria, VA  
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*Abstract  
 595*

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Title of Presentation:  
**Efficient Modeling and Simulation (M&S) Using Design of Experiments (DOE) Methods**

This presentation is:  SECRET  SECRET//REL TO FVEY  CONFIDENTIAL  CONFIDENTIAL//REL TO FVEY  
 UNCLASSIFIED  Other \_\_\_\_\_ and will be presented in:

Tutorial  List all WG(s) #:

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This presentation was NOT done under a government contract, contains no government information, is my own work and is approved for public release.  YES (Complete Part I only)

# Efficient Simulation Using Design of Experiments (DOE)

**June 16, 2014**  
**82<sup>nd</sup> MORSS Tutorial**  
**Alexandria, VA**

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**Tom Donnelly, PhD**  
SAS Institute Inc.  
JMP Systems Engineer & Co-insurrectionist

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# Outline

- **Overview of Design of Experiments (DOE)**
  - Slides 1 to 27 – background – many may be skipped if most people have an understanding of DOE
- **Efficient M&S Using DOE**
  - Sequential traditional DOE
  - Sequential space-filling DOE

## Design of Experiments (DOE) for 29 Years

- '83-'87 **Honeywell, Inc., Engineer**  
 First saw the power of DOE in 1984 – career changing event
- '87-'99 **ECHIP, Inc., Partner & Technical Director**  
 200+ DOE courses, on-site at 40+ companies - many  
 chemical/food/pharma - requiring mixture/formulation DOE
- '99-'05 **Peak Process, LLC, Consultant**
- '05-'08 **US Army, Edgewood Chemical Biological Center (ECBC),  
 Physicist, Analyst, & Co-insurrectionist**  
**DOE with Real data and Modeling & Simulation data**
- Dec. '08 **Joined the SAS Institute Inc., Customer Advocate**  
 Work in DOE and Federal Government domains
 
  - Data Visualization, Data Mining and their synergy with DOE
  - Support DoD sites, National Labs, & Gov't Contractors

## Detection, Decontamination & Protection

- JPM Nuclear Biological Chemical Contamination Avoidance (NBCCA) - Whole Systems Live Agent Test (WSLAT) Team support to the Joint Biological Point Detection System (JBPDS)
- Agent Fate wind tunnel experiments
- Decontamination Sciences Team
  - Contact Hazard Residual Hazard Efficacy Agent T&E Integrated Variable Environment (CREATIVE) - real and simulation data
  - Modified vaporous hydrogen peroxide (mVHP) decontamination – real data
- Smoke and Target Defeat Team
  - Pepper spray characterization – real data
  - Obscurant material evaluation (with OptiMetrics, Inc.) – simulation data
- U.S. Army Independent Laboratory In-house Research (ILIR) on novel experimental designs used with simulations
  - **Re-analysis of U.S. Air Force Kunsan Focused Effort BWA simulation data**
  - CB Sim Suite used for sensitivity analysis of atmospheric stability
- U.S. Marine Corps Expeditionary Biological Detection (EBD) Advanced Technology Demonstration (ATD)
  - Chamber testing of detectors – real data
  - CB Sim Suite sensor deployment studies – simulation data
- U.S. Navy lead on Joint Expeditionary Collective Protection (JECP)
  - Swatch and chamber testing – real data
  - **Computational Fluid Dynamics (CFD) – simulation data**

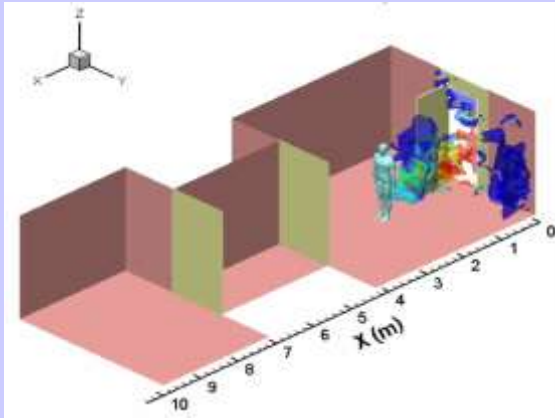
- **PDFs available**
  - Today's Slides
  - White Paper - “Efficient Modeling & Simulation of Biological Warfare Using Innovative Design of Experiments Methods”
  - MORSS Tutorial Summary 11 X 17 Handout  
“DOE for Real-World Problems”

*Quicker answers, lower costs, solve bigger problems*

- Obtain a fast surrogate model of the simulation
  - Individual simulations can run for hours, days, weeks
    - Computational Fluid Dynamics (CFD)
    - Simulation runs in real-time
  - Numbers of factors can be very large (40+)
  - Numbers of simulations needed can be large (thousands in many cases)
  - Simulations can be stochastic requiring many replications
- Surrogate model yields a fast approximation of the simulation
  - more rapidly answer “what if?” questions
  - do sensitivity analysis of the control factors
  - optimize multiple responses and make trade-offs
- By running efficient subsets of all possible combinations, one can – for the same resources and constraints – *solve bigger problems*
- By running sequences of designs one can be as *cost effective as possible & run no more trials than are needed to get a useful answer*

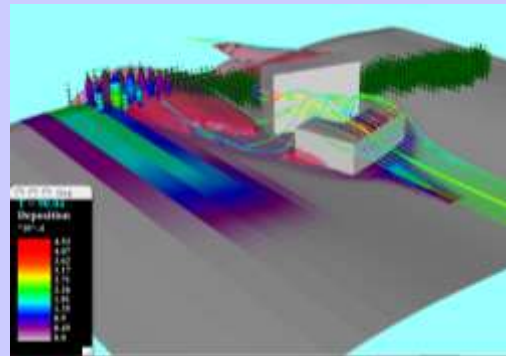
Detailed Physics Models can require a great deal of runtime to generate a short period of simulation time.

## Computational Fluid Dynamics (CFD) Models



Developed for Interior  
**Moving Man in Simulation**  
 8M cells  
**10 Seconds of Simulation**  
**64 CPUs – 4K slower**  
 12 Hours of Runtime

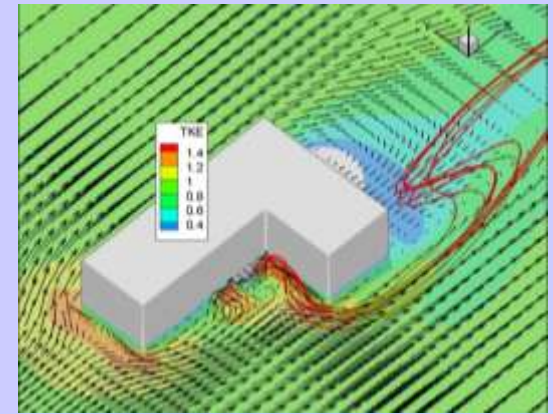
**Detailed Ingress/Egress,  
 Internal Airflow and  
 Convection**



Developed for Exterior  
**Stationary Grids**  
 1.5M Cells  
**30 Seconds of Simulation**  
**Single CPU – 20K slower**  
 7 Days of Runtime

**External CW Deposition/  
 Evaporation, Vegetation,  
 Solar Heating**

## Lagrangian-Particle

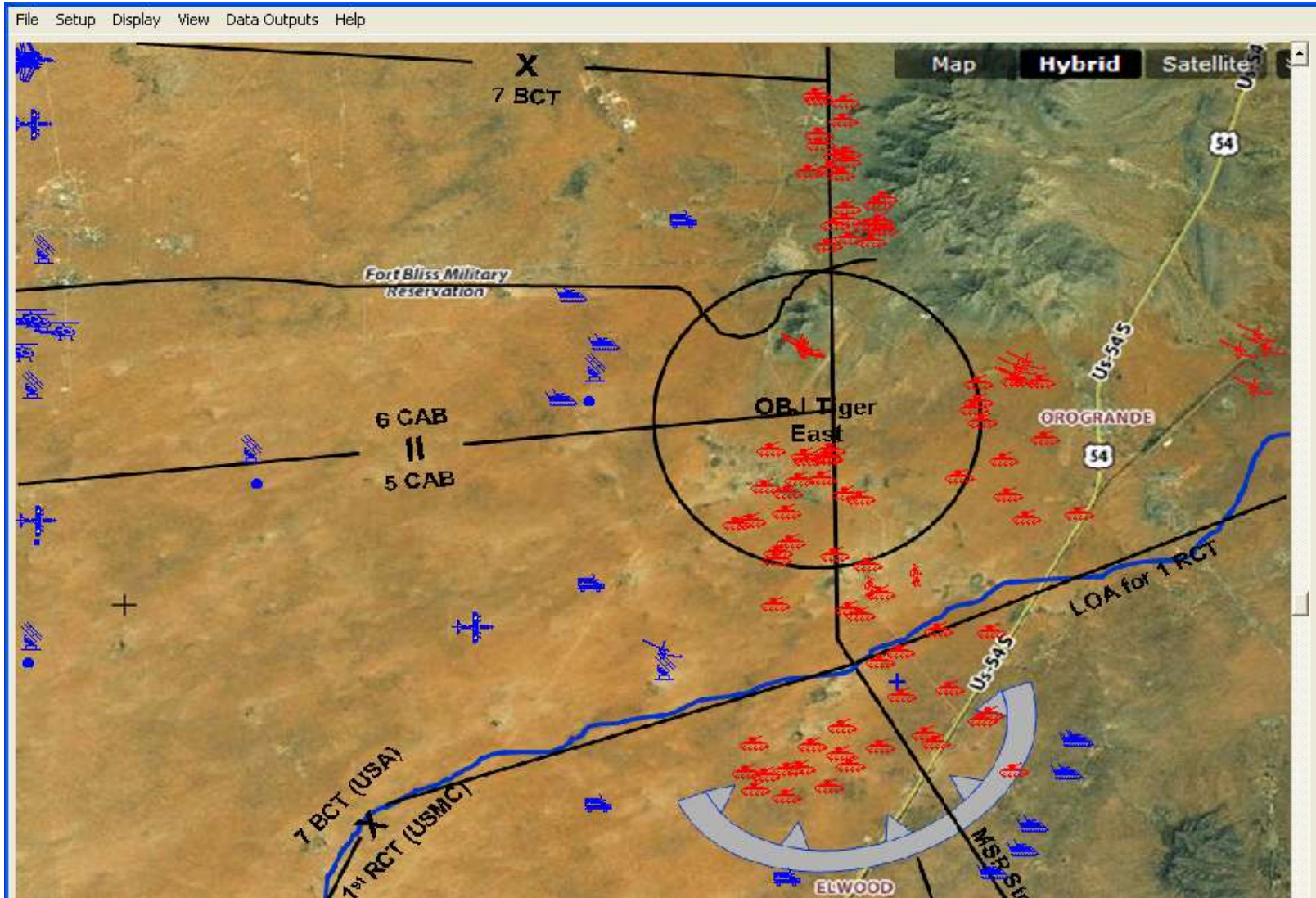


Developed for Exterior  
**Stationary Grids**  
 TBD Cells  
**Min-Hours of Simulation**  
**Single CPU**  
 Minutes-Days of Runtime

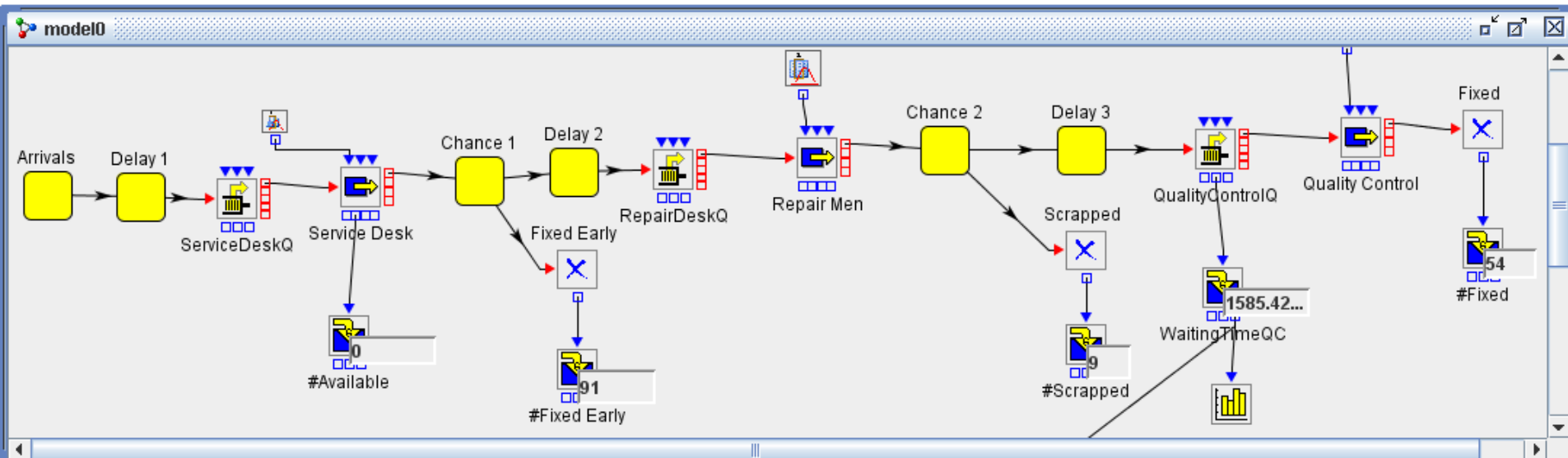
**Speed, Flexibility, More User  
 Friendly, V&V**



## Agent-Based Simulations



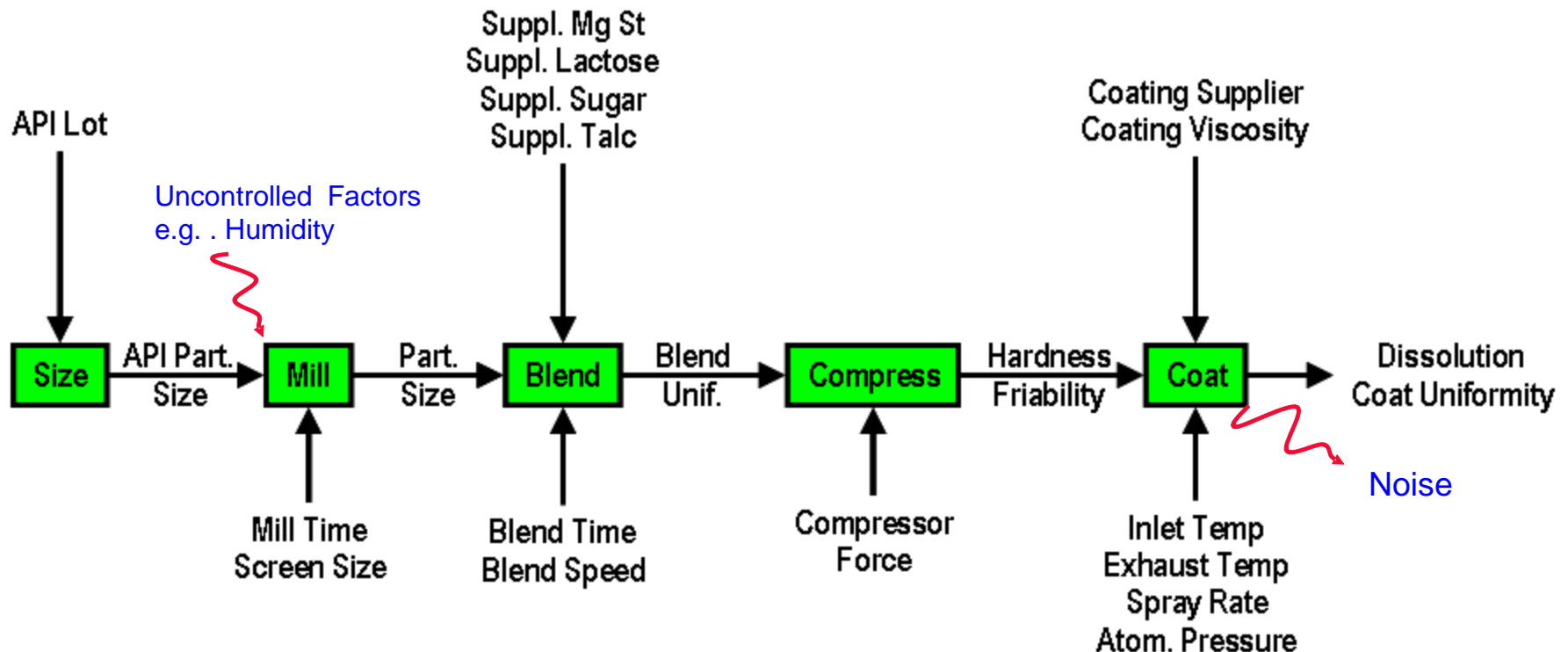
## Discrete Event Simulations



Point...	StartT...	EndTI...	Num...	Num...	Num...	Repli...	Num...	AvgUt...	AvgW...	AvgUti...	AvgUt...	AvgW...	AvgW...
point 1	0	2,700	1	1	3	▶ 5	55.8	98.1...	640...	32.90...	22.7...	1.681...	0.11...
point 2	0	2,700	3	2	1	▶ 5	105.6	61.8...	3.65...	16.39...	67.8...	0.0	16.7...
point 3	0	2,700	2	3	1	▶ 5	100.2	88.3...	84.8...	10.92...	67.8...	0.0	16.7...
point 4	0	2,700	2	1	3	▶ 5	100.6	88.5...	97.3...	32.90...	22.7...	1.681...	0.11...
point 5	0	2,700	2	1	1	▶ 5	100.2	88.3...	84.6...	32.78...	67.8...	0.233...	16.7...
point 6	0	2,700	3	1	2	▶ 5	105.8	61.9...	8.69...	32.90...	34.1...	1.382...	0.83...
point 7	0	2,700	2	2	2	▶ 5	100.4	88.4...	97.9...	16.47...	34.1...	0.020...	0.83...
point 8	0	2,700	2	2	3	▶ 5	100.6	88.5...	98.4...	16.46...	22.7...	0.094...	0.11...
point 9	0	2,700	1	1	1	▶ 5	55.8	98.1...	621...	32.78...	67.8...	0.233...	16.7...
point ...	0	2,700	3	3	3	▶ 5	105.8	61.9...	9.32...	10.97...	22.7...	0.001...	0.11...
point ...	0	2,700	1	3	2	▶ 5	55.8	98.1...	641...	10.98...	34.1...	4.305...	0.83...
point ...	0	2,700	1	2	1	▶ 5	55.8	98.1...	621...	16.39...	67.8...	0.0	16.7...

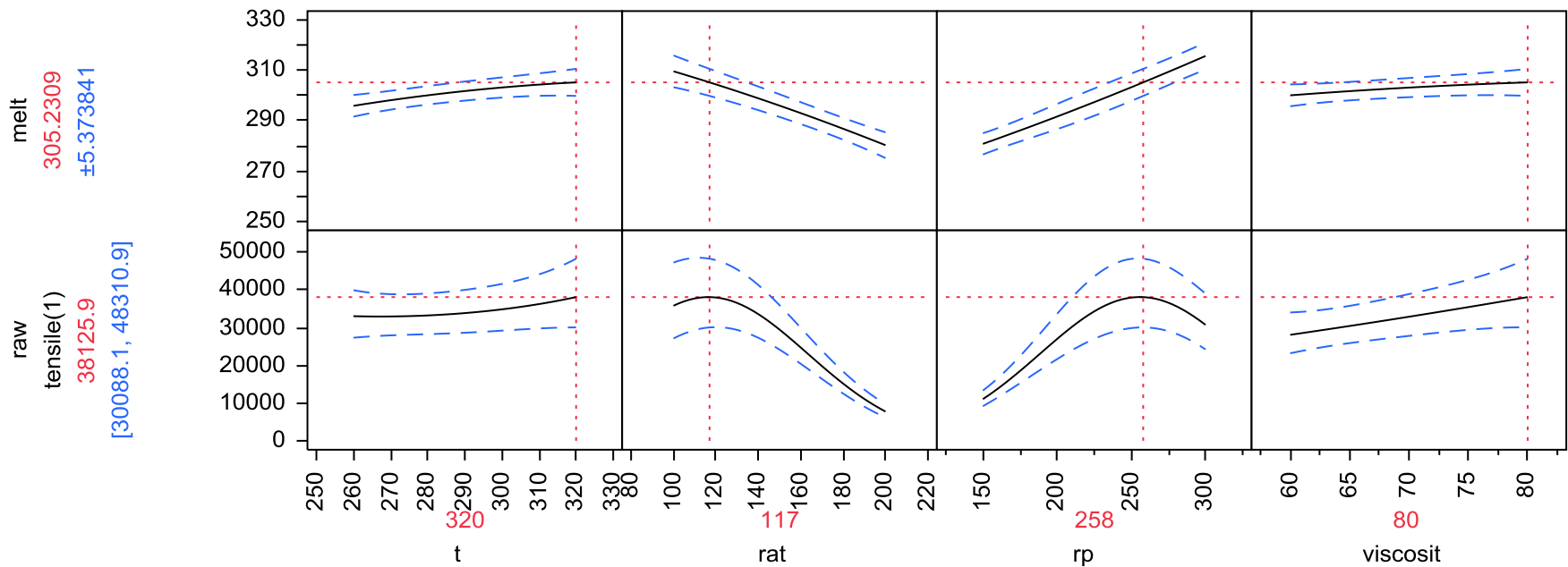
## Classic Definition of DOE

- Purposeful control of the inputs (factors) in such a way as to deduce their relationships (if any) with the output (responses).



# Here are 4 Controls (inputs) & 2 Responses (outputs) and their empirical relationships (model)

Prediction Profiler

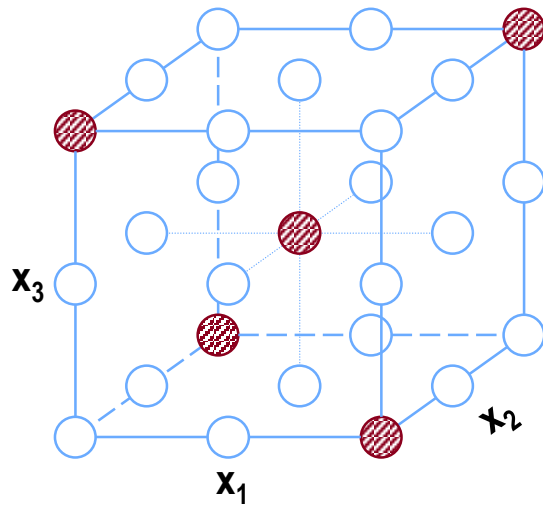


Get this Prediction Profiler as result of analyzing data collected for a DOE

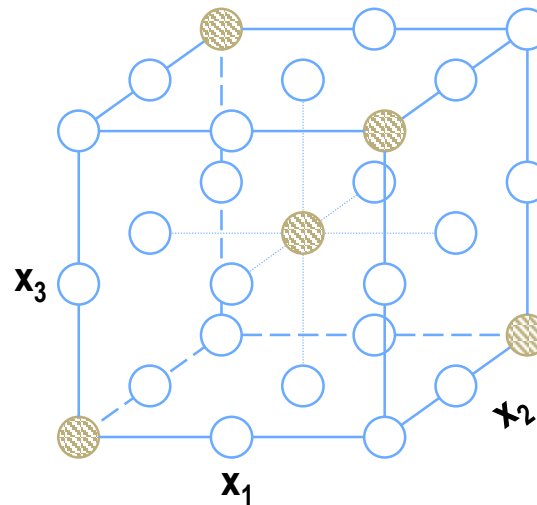
## Alternative Definition

- A DOE is the specific collection of trials run to support a proposed model.
  - If proposed model is **simple**, e.g. just main or 1<sup>st</sup> order effects ( $x_1, x_2, x_3$ , etc.), the design is called a **screening** DOE
    - Goals include **rank factor importance** or find a “winner” quickly
    - Used with many (> 6?) factors at start of process characterization
  - If the proposed model is **more complex**, e.g. the model is 2<sup>nd</sup> order so that it includes two-way interaction terms ( $x_1x_2, x_1x_3, x_2x_3$ , etc.) and in the case of continuous factors, squared terms ( $x_1^2, x_2^2, x_3^2$ , etc.), the design is called a **response-surface** DOE
    - Goal is generally to develop a **predictive model** of the process
    - Used with a few (< 6?) factors after a screening DOE

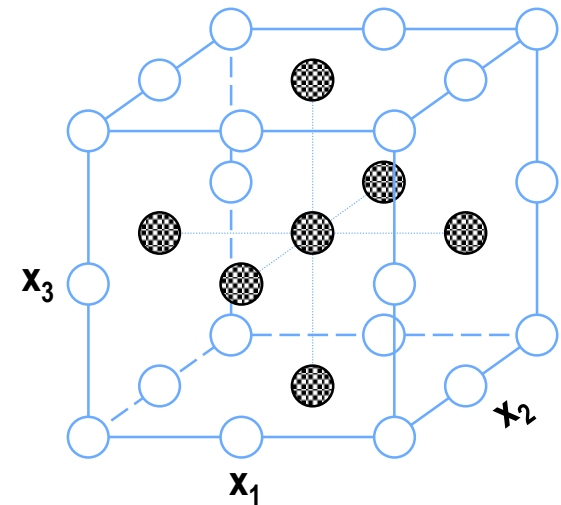
### Block 1



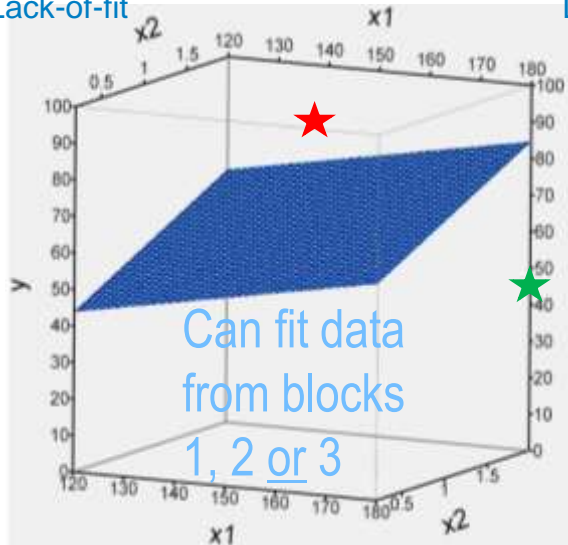
### Block 2



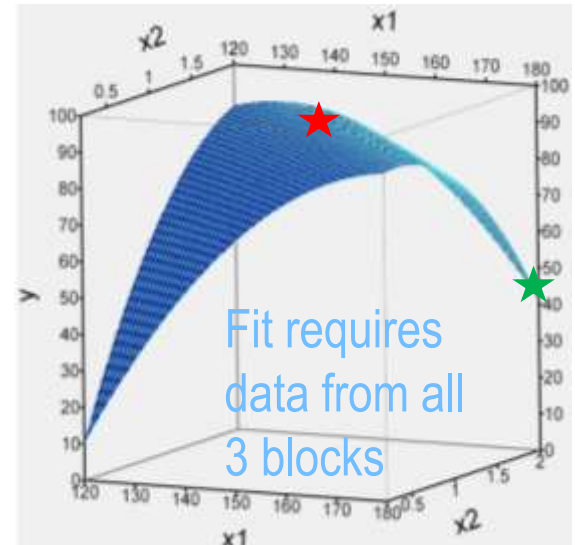
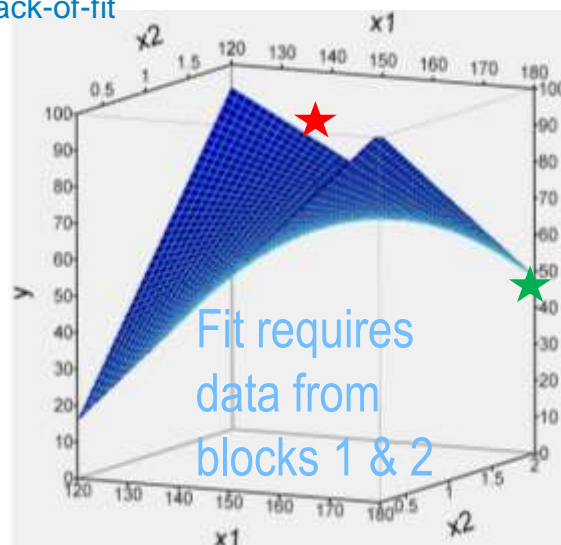
### Block 3



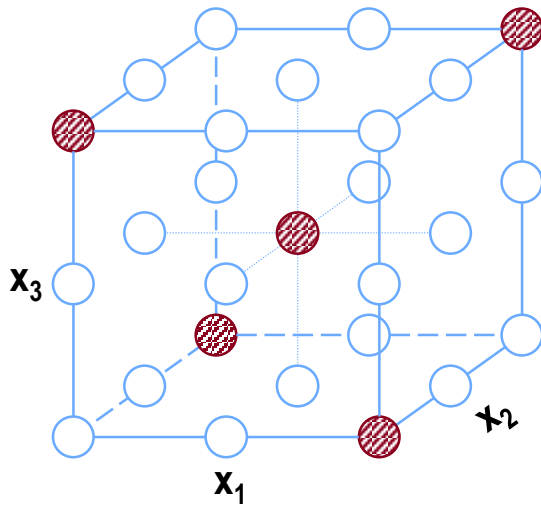
Lack-of-fit



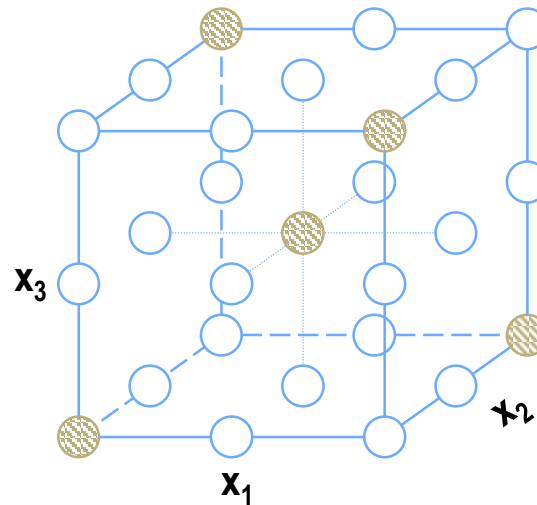
Lack-of-fit



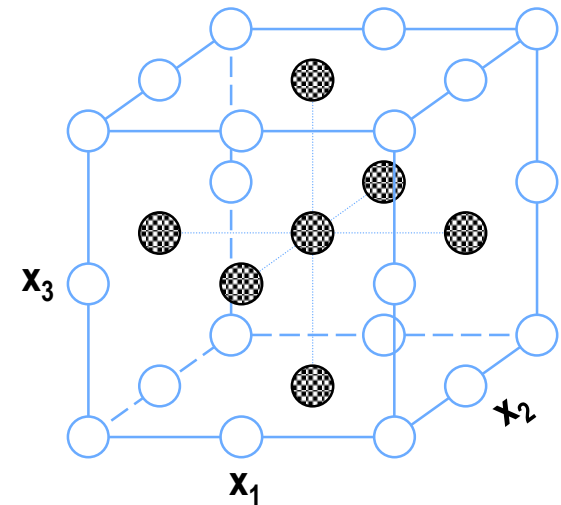
### Block 1



### Block 2



### Block 3



$$y = a_0 + a_1x_1 + a_2x_2 + a_3x_3$$

Run this block 1st to:

- (i) estimate the main effects\*
- (ii) use center point to check for curvature.

\*May be all that are needed with appropriate physics-based scaling

Also called non-linear modeling

$$y = a_0 + a_1x_1 + a_2x_2 + a_3x_3$$

$$+ a_{12}x_1x_2 + a_{13}x_1x_3 + a_{23}x_2x_3$$

Run this block 2nd to:

- (i) repeat main effects estimate,
- (ii) check if process has shifted
- (iii) add interaction effects to model if needed.

$$y = a_0 + a_1x_1 + a_2x_2 + a_3x_3$$

$$+ a_{12}x_1x_2 + a_{13}x_1x_3 + a_{23}x_2x_3$$

$$+ a_{11}x_1^2 + a_{22}x_2^2 + a_{33}x_3^2$$

Run this block 3rd to:

- (i) repeat main effects estimate,
- (ii) check if process has shifted
- (iii) add curvature effects to model if needed.

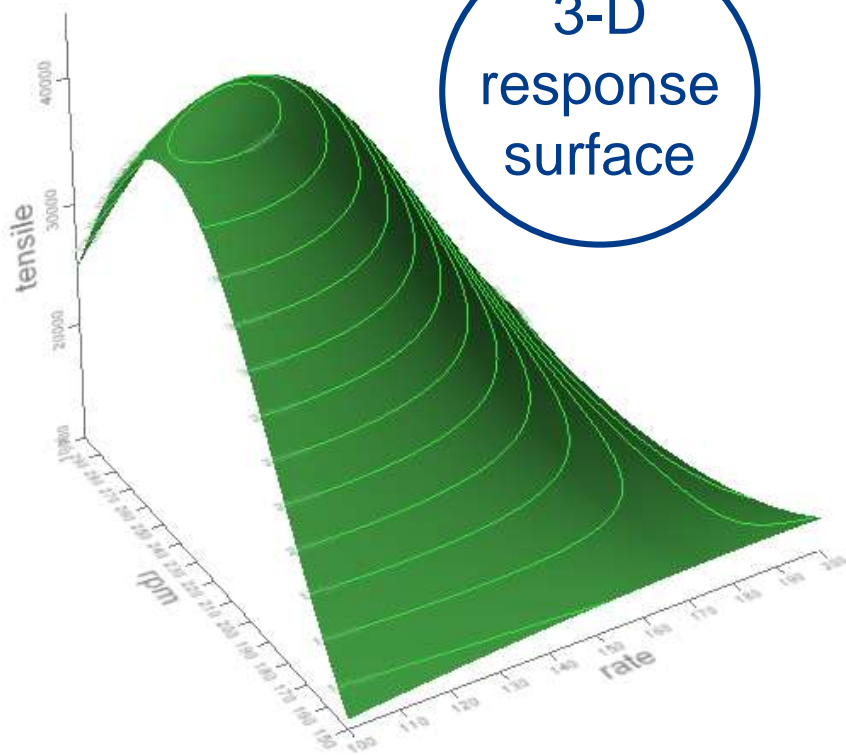
## Why is Using DOE Important?

- *“One thing we have known for many months is that the spigot of defense funding opened by 9/11 is closing.”*
- *“In the past, modernization programs have sought a 99 percent solution over a period of years, rather than a 75 percent solution over a period of weeks or months.”*
  - Two quotes from the January 27, 2009 submitted statement of Secretary of Defense Robert M. Gates to the Senate Armed Services Committee.
- DOE is one of the more powerful tools we can use to efficiently accomplish our goals.
  - DOE yields the maximum information from the fewest experiments.
  - DOE often yields an 80% solution in less than 20% of the work.



# Response Surface & Contour Plot (four control variables)

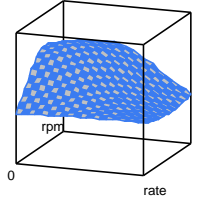
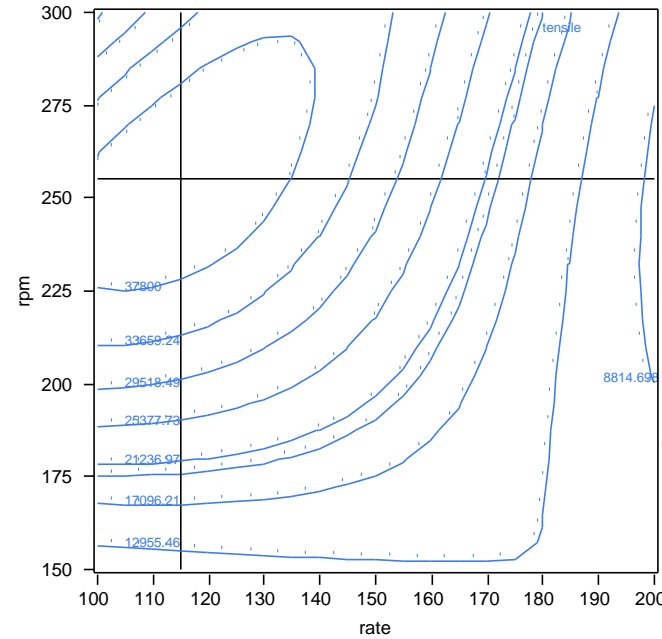
3-D  
response  
surface



Horiz	Vert	Factor	Current X
<input type="radio"/>	<input type="radio"/>	t4	320
<input checked="" type="radio"/>	<input type="radio"/>	rate	115
<input type="radio"/>	<input checked="" type="radio"/>	rpm	255
<input type="radio"/>	<input type="radio"/>	viscosity	80

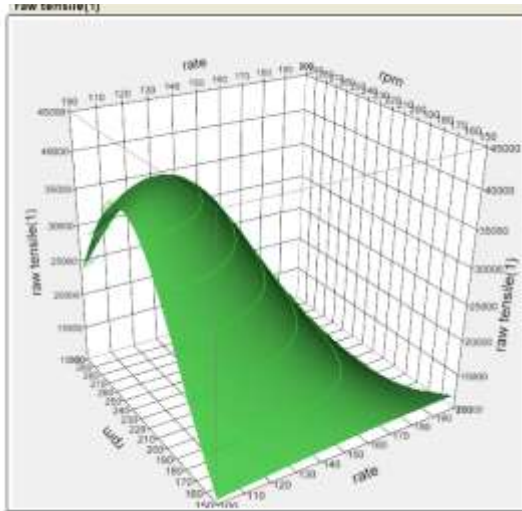
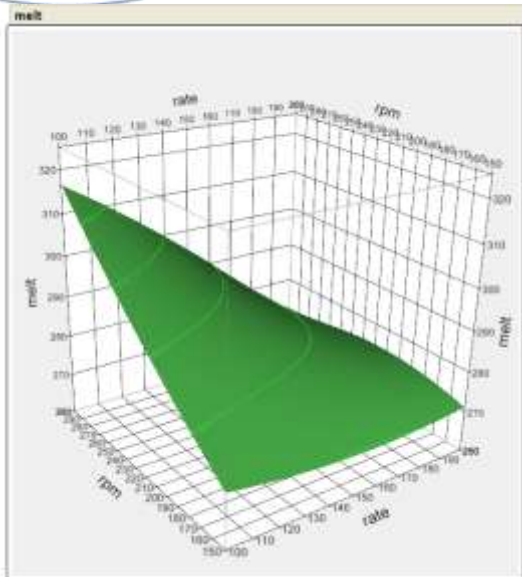
  

Response	Contour	Current Y	Lo Limit	Hi Limit
melt	250	305.35337	.	.
tensile	20000	41081.766	.	.



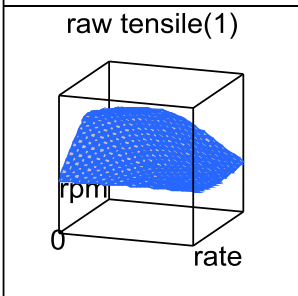
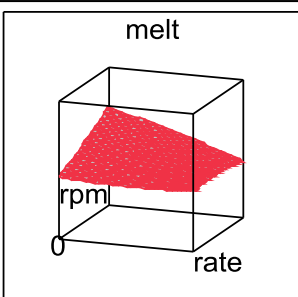
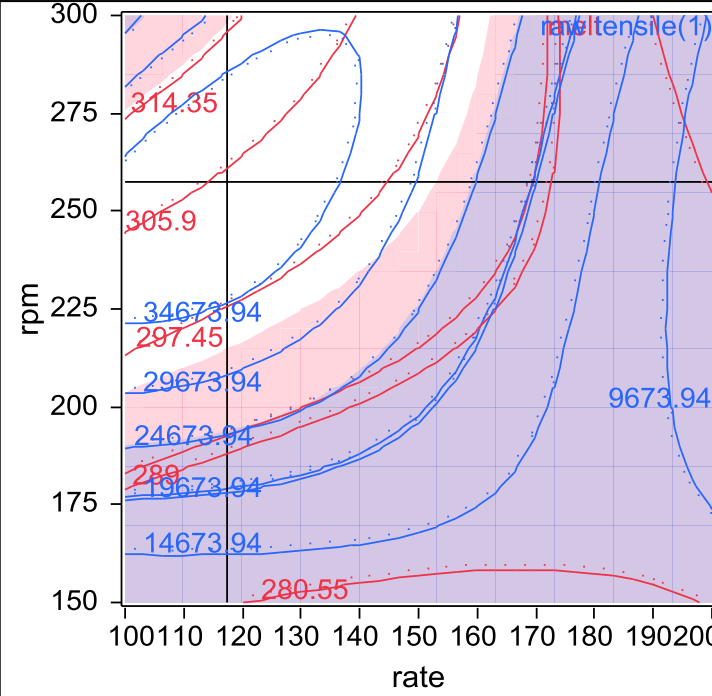
2-D  
contour  
plot

*(four control variables & two responses)*



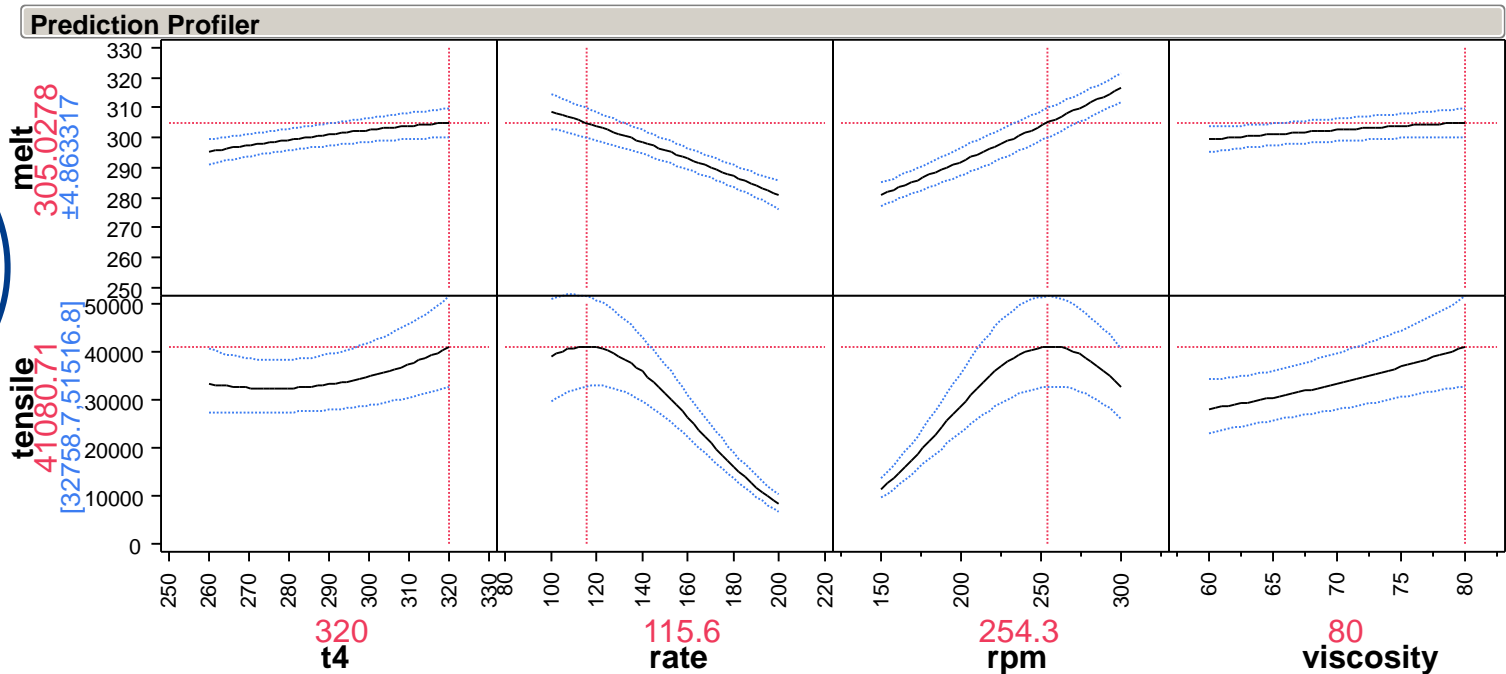
## Contour Profiler

HorizVert Factor		Current X			
t4			320		
rate			117.29498		
rpm			257.64505		
viscosity			80		
Response	Contour	Current Y	Lo Limit	Hi Limit	
melt	290	305.06654	295	315	
raw tensile(1)	20000	38127.616	25000	40000	

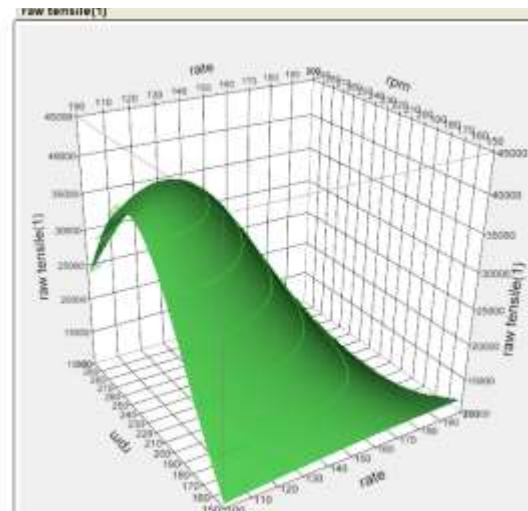
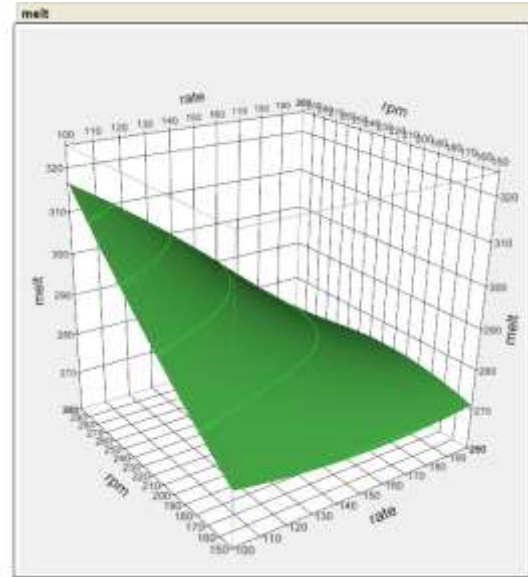


# 1-D Prediction Profiles are a Way to View Higher Dimensionality as “Interactive Small Multiples” - Here 4 Controls & 2 Responses

1-D  
profiler  
plots



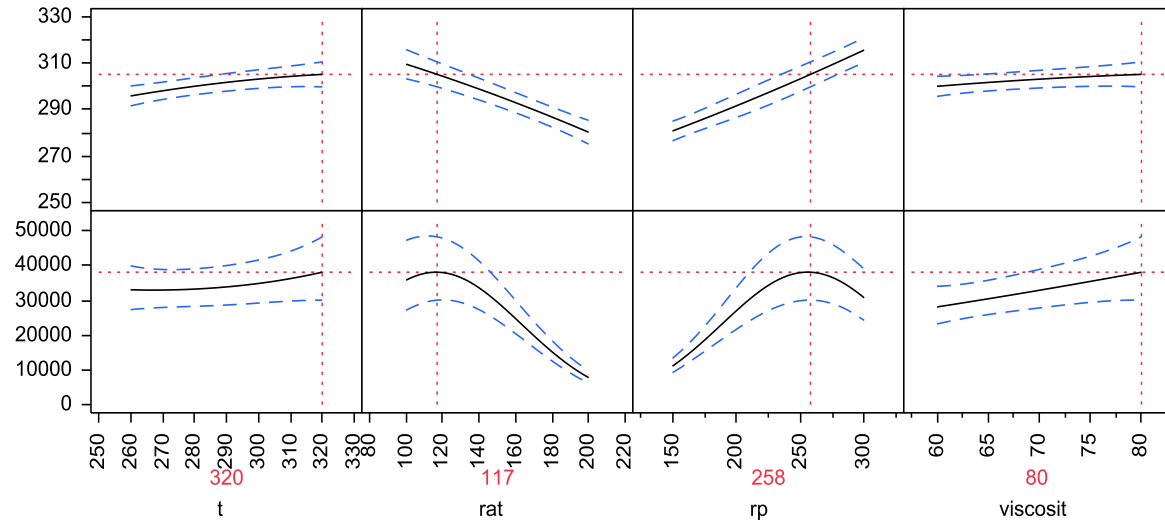
# 1-D Prediction Profiles are a Way to View Higher Dimensionality as “Interactive Small Multiples” - Here 4 Controls & 2 Responses



Prediction Profiler

melt  
305.2309  
±5.373841

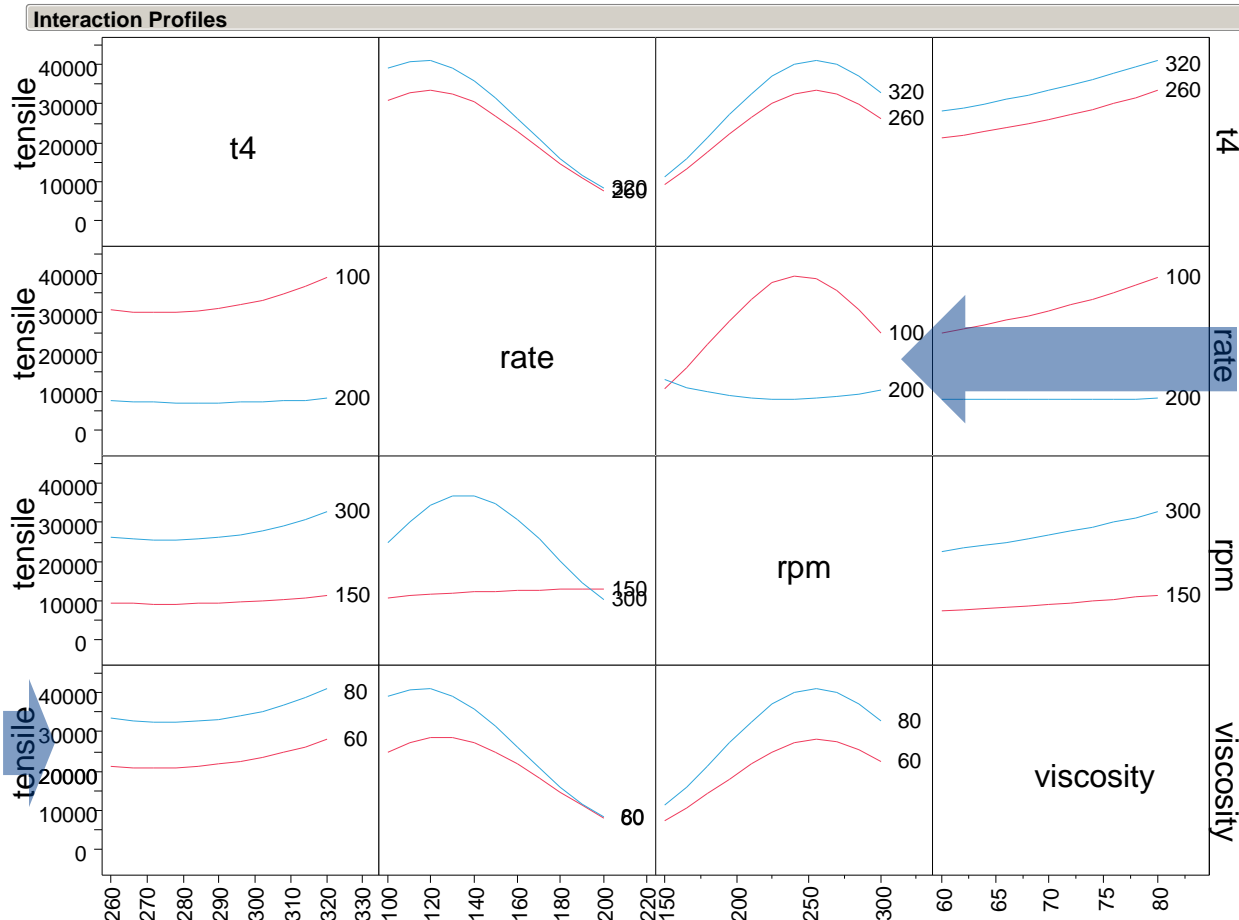
raw  
tensile(1)  
38125.9  
[30088.1, 48310.9]



# Interaction Profiles are Another Way to View Higher Dimensionality - Here 4 Controls and 1 Response

1-D plots at high & low of other factors

Parallel indicates NO interaction



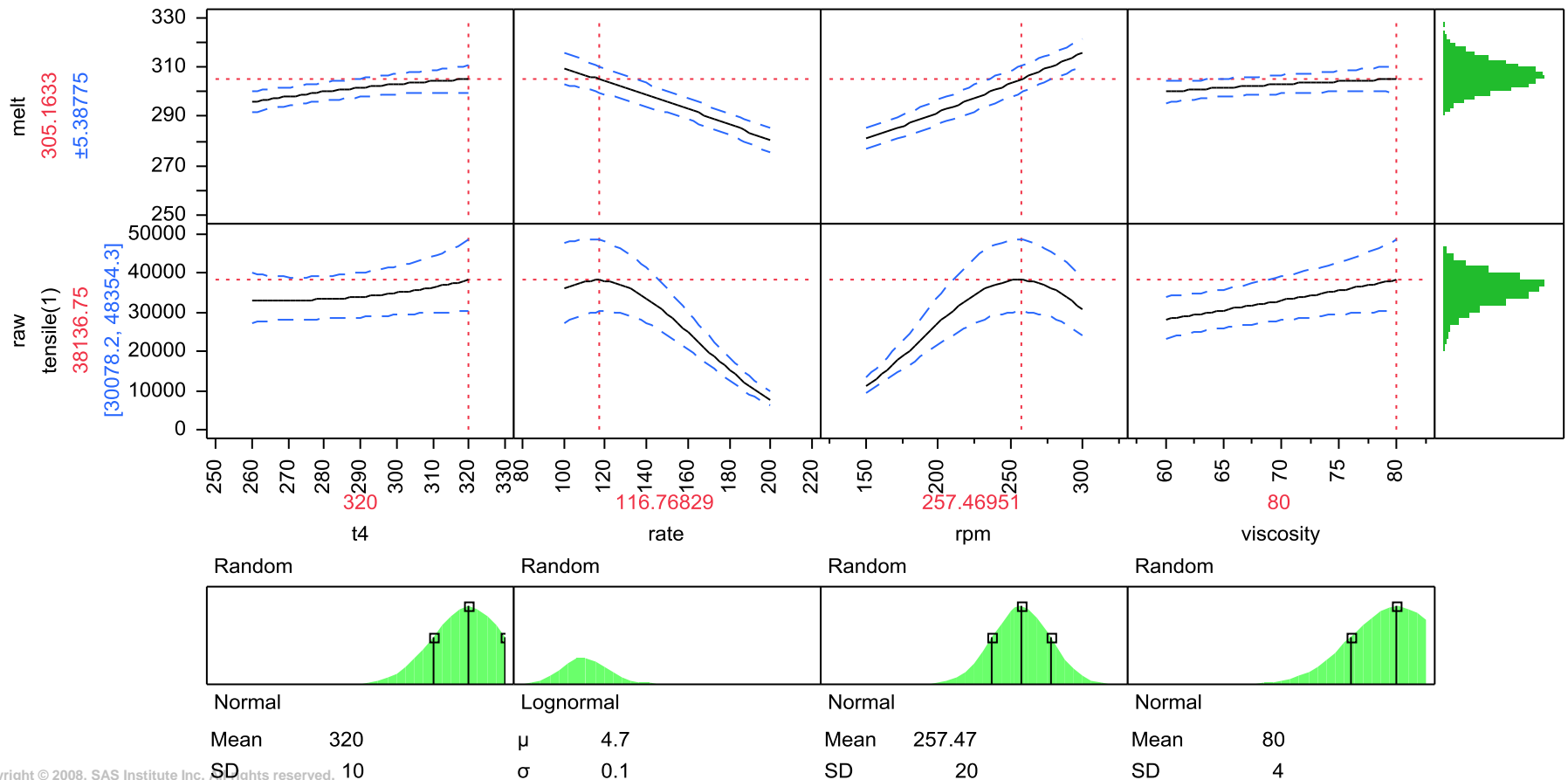
NOT Parallel indicates interaction



# Assess Uncertainty in Surrogate Model Predictions Even for a Non-Stochastic Simulation with No Replications

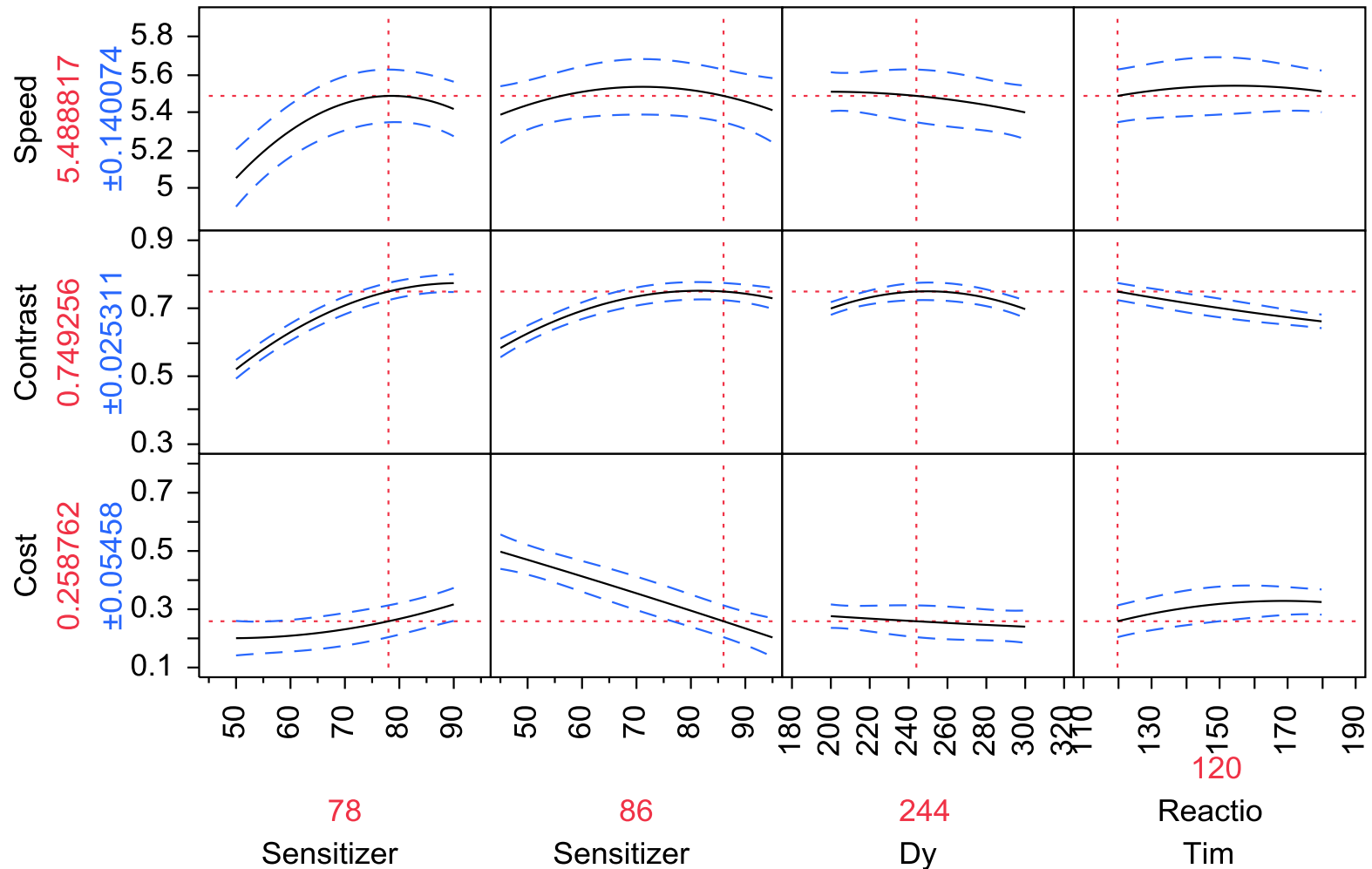
For non-stochastic simulations for which a surrogate model has been created, Monte Carlo simulations can be run using assumed distributions for inputs to better assess transmitted variation about the model point estimate.

Prediction Profiler



- System with 2 primary MoPs and Cost
- Management wants to lower cost but maintain performance
- Multi-response and multi-factor process characterized using a JMP Custom DOE that supports interactive optimization – the trading off of performance and cost
- Management provided with visually interactive process knowledge that makes their decision making easier

## 3 responses and 4 control factors





Box, G. E. P., Hunter, W. G., and Hunter, J. S. (2005), *Statistics for Experimenters*, 2<sup>nd</sup> ed., Wiley, New York

- **The classic 1978 text recently revised**

Wu, C. F. J. and Hamada, M. (2009), *Experiments, Planning, Analysis and Parameter Design Optimization*, , 2<sup>nd</sup> ed., Wiley, New York

- **Both classic DOE approaches and orthogonal arrays & orthogonal main effects plans**

Montgomery, D. C. (2009), *Design and Analysis of Experiments*, 7<sup>th</sup> ed., Wiley, New York

- **Popular text, solution book available, examples illustrated with DOE software.**

- **8<sup>th</sup> Edition includes newly developed screening approaches**

- **All problems worked in JMP software due out this year**

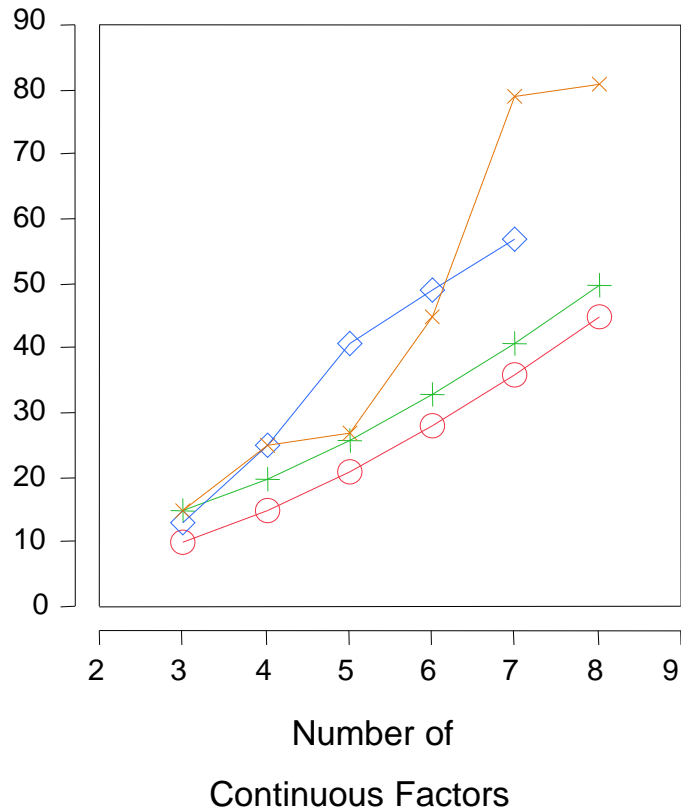
Texts Specifically on DOE for Computer Experiments:

Kleijnen, J. P. C. (2008), *DASE: design and analysis of simulation experiments*. Springer, New York.

Santner, T. J., Williams, B. J., and Notz, W. I. (2003), *The Design and Analysis of Computer Experiments*, Springer, New York

Fang, K. T., Li, R. Z., and Sudjianto, A. (2005), *Design and Modeling for Computer Experiments*, Chapman & Hall/CRC Press, New York

# # Unique Trials for 3 Response-Surface Designs and # Quadratic Model Terms VS. # Continuous Factors



- Y
- × — Unique Trials in Central Composite Design
  - ◇ — Unique Trials in Box-Behnken Design
  - + — Unique Trials in Custom Design with 5 df for Model Error
  - — Terms in Quadratic Model = Minimum # of Trials

If generally running 3, 4 or 5-factor fractional-factorial designs...

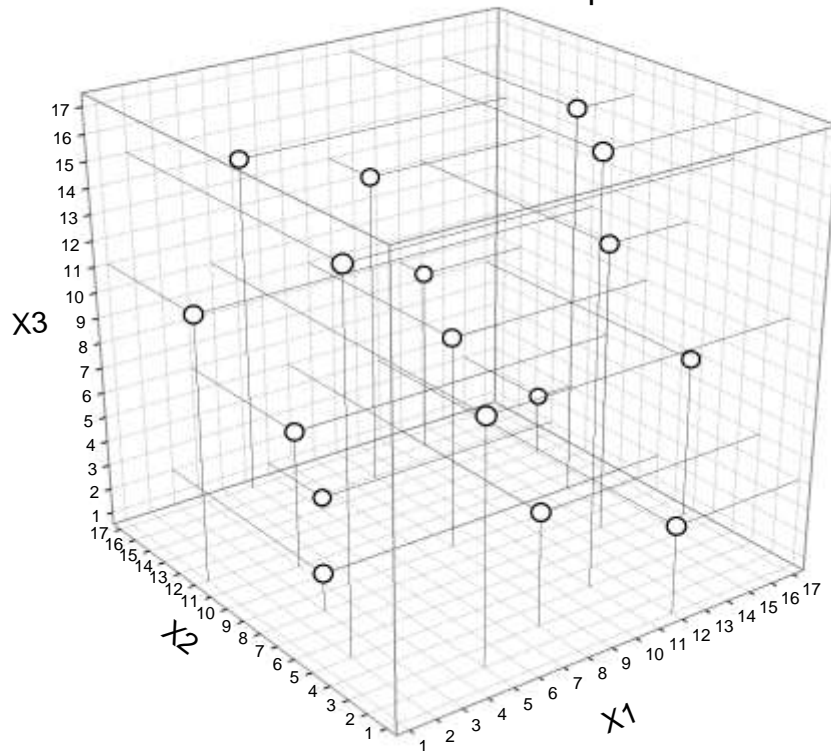
1. How many interactions are you NOT investigating?
2. How many more trials needed to fit curvature?
3. Consider two stages – Definitive Screening + Augmentation

## How many folks have any of these issues?

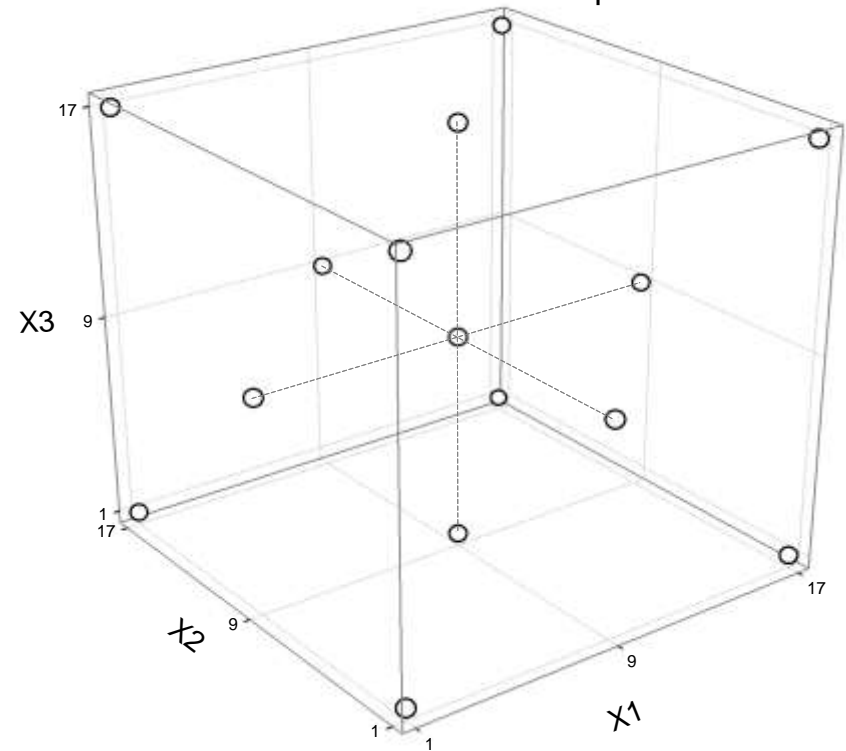
- Work with these different kinds of control variables/factors:
  - **Continuous/quantitative?** (Finely adjustable like *temperature, speed, force*)
  - **Categorical/qualitative?** (Comes in types, like material = *rubber, polycarbonate, steel* with mixed # of levels; 3 chemical agents, 4 decontaminants, 8 coupon materials...)
  - **Mixture/formulation?** (Blend different amounts of *ingredients* and the process performance is dependent on the *proportions* more than on the amounts)
  - **Blocking?** (e.g. “lots” of the same raw materials, multiple “same” machines, samples get processed in “groups” – like “eight in a tray,” run tests over multiple days – i.e. variables for which there *shouldn’t* be a causal effect)
- Work with **combinations of these four kinds** of variables?
- Certain **combinations cannot be run?** (too costly, unsafe, breaks the process)
- Certain factors are **hard-to-change** (temperature takes a day to stabilize)
- Would like to **add onto existing trials?** (really expensive/time consuming to run)
- Characterize process or **run experiments using computer simulations?** (war gaming, agent-based, discrete event, computational fluid dynamics (CFD))
- Measure response **data in vicinity of physical limits?** (counts, hardness, resistivity can’t fall below zero, or percentage yield or killed can’t exceed 100%)

- “Traditional factorial/response surface” designs for polynomial modeling with categorical (qualitative) and continuous (quantitative) variables
  - Designs can be sequentially constructed to support increasingly complex models
  - Example featured here reanalyzes a simulation case matrix in which all combinations of 6 variable settings were originally run- a total of  $648 = 6 \times 3 \times 3 \times 3 \times 2 \times 2$
  - References on Resolution V, Fractional-Factorial Designs for many (40+) factors
    - Mee, R. W. (2004), **Efficient Two-Level Designs for Estimating Main Effects and Two-Factor Interactions**, *Journal of Quality Technology*, 36, 400-412.
    - Sanchez, S.M. and Sanchez, P.J. (2005), **Very Large Fractional Factorial and Central Composite Designs**, *ACM Transactions on Modeling and Computer Simulation*, Vol. 15, No. 4, October 2005, Pages 362–377.
    - Xu, H. (2009), **Algorithmic Construction of Efficient Fractional Factorial Designs with Large Run Sizes**, *Technometrics*, (in press) <http://www.stat.ucla.edu/~hqxu/pub/ffd2r3.pdf>
  
- “Space-filling” designs primarily for use with continuous variables AND non-stochastic/deterministic responses
  - These designs can support “Gaussian Process” or “Kriging” spatial regression analysis – an interpolation technique, as well as linear regression – an approximation method

Space-Filling Design  
for 3 Variables with 17 Unique Trials



Response-Surface Design  
for 3-Variables with 15 Unique Trials



Rather than emphasizing high leverage trials (“corners”) for a simple polynomial model, space-filling designs “spread” their trials more uniformly through the space to better capture the local complexities of the simulation model.

- I used to say “If a “textbook” fractional-factorial, orthogonal array or response-surface design is available, then use it.”  
 Now I say, “If Definitive Screening or Minimal Alias design is available, then use it.”
- Textbooks and web site catalogs do not always contain designs for categorical variables with:
  - all combinations of mixed numbers of levels (e.g. 3, 4, 5, and 21)
  - large numbers of levels for variables (e.g. 5+)
- Algebraic (Orthogonal Array) and algorithmic (D-optimal) computer generated designs can often be used
  - Orthogonal Arrays are good at yielding analysis with unconfounded estimates of the “main effects” when variables have many different levels
  - D-optimal designs are good for adding on the fewest additional trials to support higher order “interaction” terms in the model

- Simulation experiments – Sequential designs are easily employed because “restricted randomization” is not an issue
  - Many simulations are deterministic
  - Even if stochastic (random), correlation with unknown factors is not possible
  - All factors are generally just as easy to change
  - Can still inexpensively add a blocking variable to test if “the code has been changed!”
  
- Real experiments – The issue of “restricted randomization” does arise making sequential experimentation a bit more complicated – but still possible to employ
  - Groups of trials run at different (even widely spaced) periods of time
    - Addressed using a *blocking* factor
  - Sometimes there are factors that are harder to change than others, e.g. *Oven Temperature*
    - Addressed using *split-plot* designs

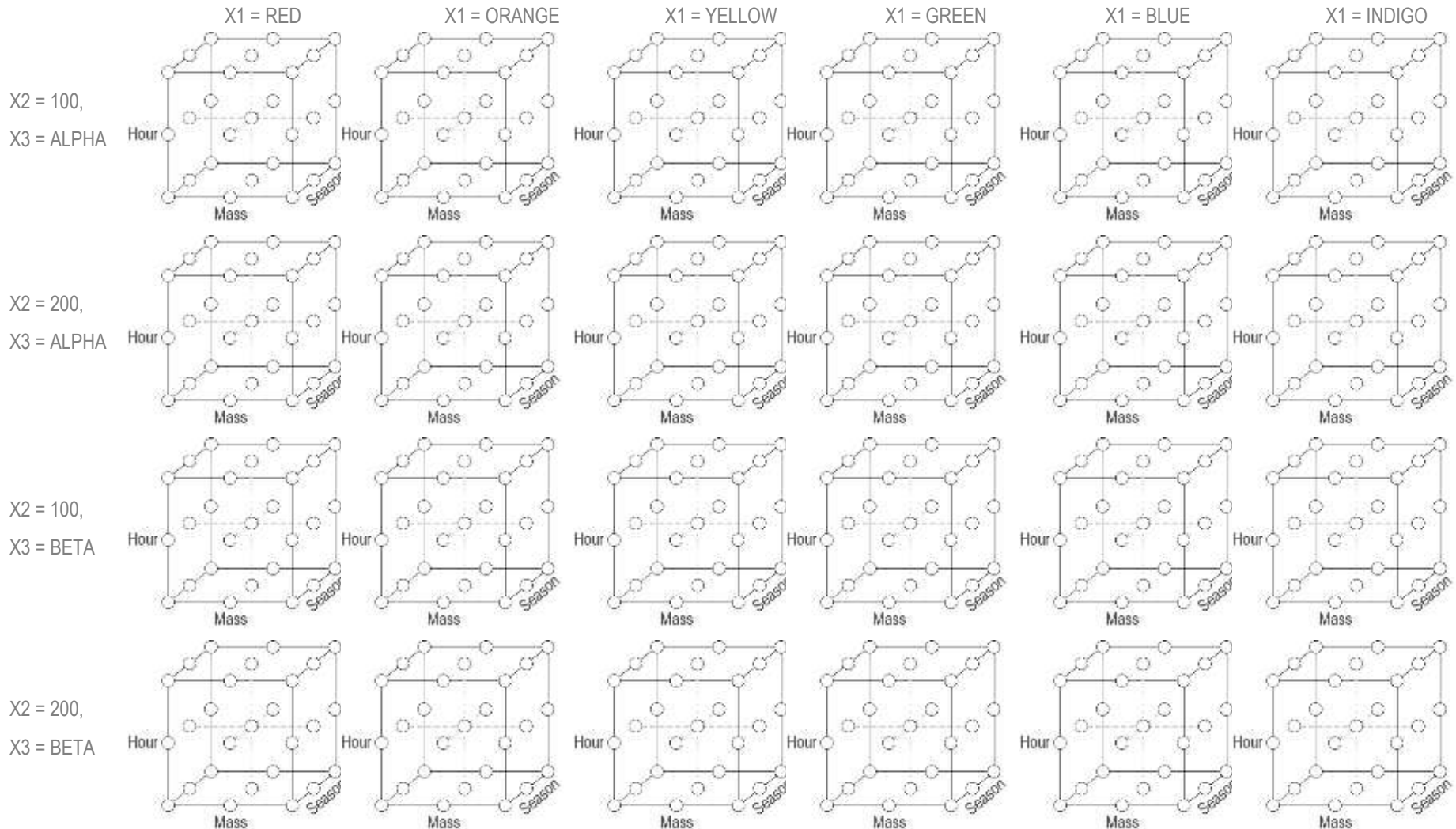


# Case Matrix as Used in Study of the Observed Response “Probability of Casualty” (PCAS)

Variable	# Levels	Levels
Agent Codes (X1)	6	A, N, T, H, R, Y (categorical)
Season	3	Winter, Summer, Spring/Fall (categorical)
Time of Attack (Hour)	3	0500, 1200, 2200 Local Time (continuous)
No. of TBMs & Spread Radius (X2)	2	1 TBM & 1 m, 2 TBMs & 1000 m (categorical)
Mass (relative)	3	1.00, 1.57, 2.00 (continuous)
Height of Burst (X3)	2	0, 10 m (continuous)
Total Cases	648	



# All 648 Possible Combinations of Settings for 6 Variables (6 X 2 X 2 X 3 X 3 X 3)



# Four Stage Design Sequence

## Stage 1

36 Total Simulations

Design 1, 36 trials

Main effects only for ALL variables  
+ some 2-way interactions

5.6% of 648

## Stage 2

108 Total Simulations

Design 1, 36 trials

Design 2, 72 trials

Stage 1 effects plus all 2-way interactions  
+ some 3-way interactions

16.7% of 648

## Stage 3

324 Total Simulations

Design 1, 36 trials

Design 2, 72 trials

Design 3, 216 trials

Stage 2 effects plus all 3-way interactions

50% of 648

## Stage 4

ALL 648 Simulations

Design 1, 36 trials

Design 2, 72 trials

Design 3, 216 trials

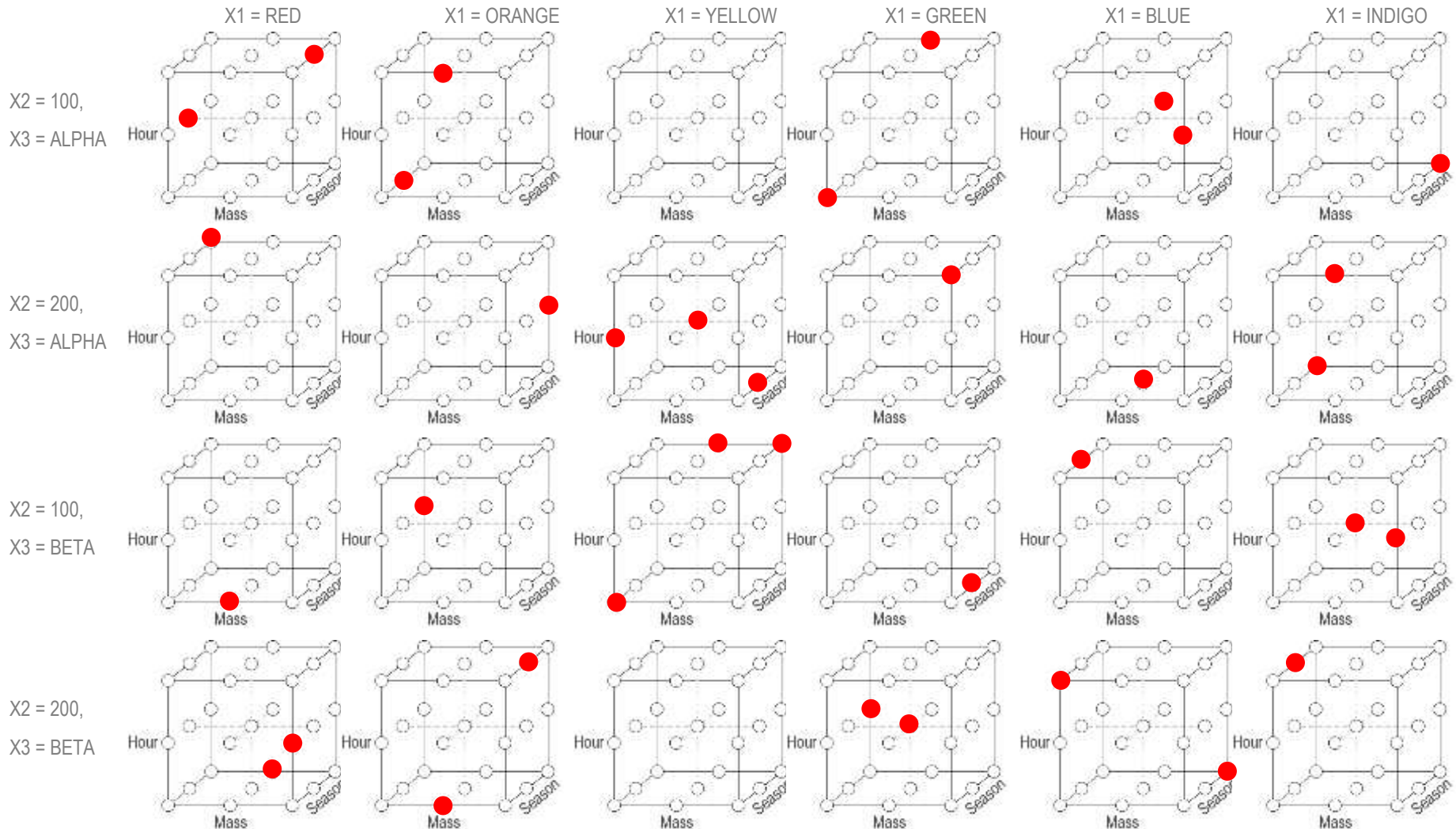
Stage 3 effects plus ALL remaining 4-way, 5-way and 6-way interactions

Design 4, 324 trials

NOTE: Length of this green box should be longer than shown

324 trials in Design 4 used as checkpoints for Designs 1, 2 & 3 →

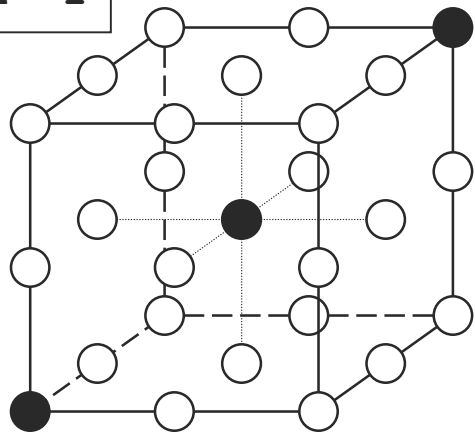
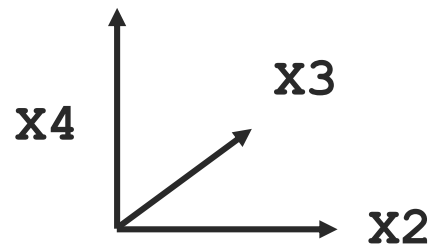
# 36 of All 648 Possible Combinations of Settings for 6 Variables (6 X 2 X 2 X 3 X 3 X 3)



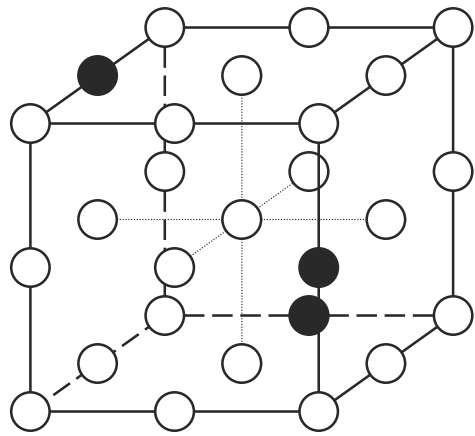
Red Dots Mark the 36 Trials (an Orthogonal Array) Analyzed for Stage 1

## Locations of Trials for a 4-variable, 9-trial Orthogonal Array Design

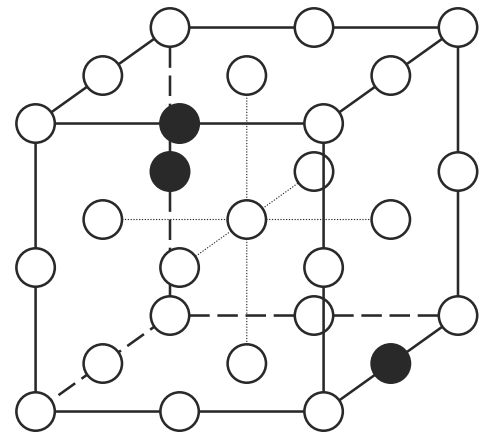
x1	x2	x3	x4
1	1	1	1
1	2	2	2
1	3	3	3
2	1	2	3
2	2	3	1
2	3	1	2
3	1	3	2
3	2	1	3
3	3	2	1



**x1 = 1**



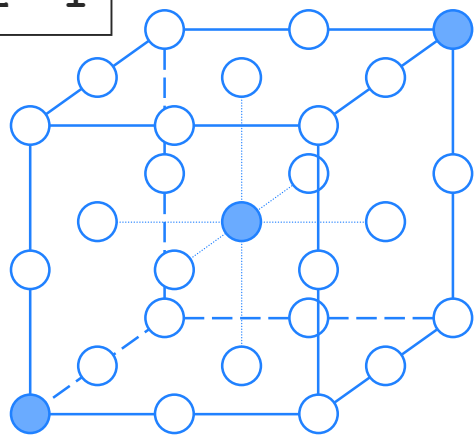
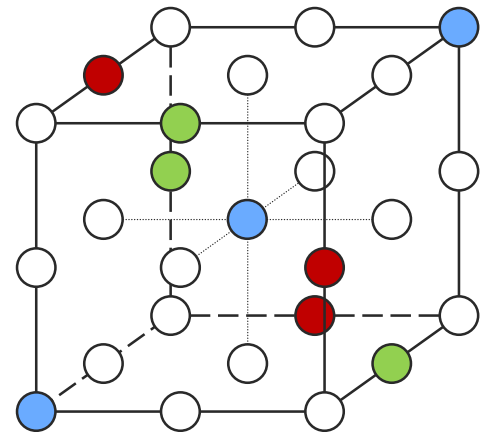
**x1 = 2**



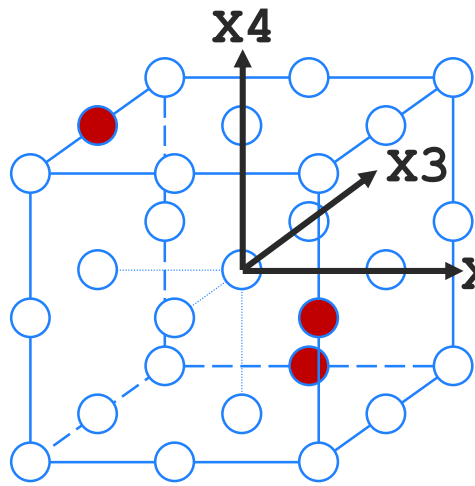
**x1 = 3**

Delete **x1** and View  
Locations of Trials for a  
3-Variable OA9 Design

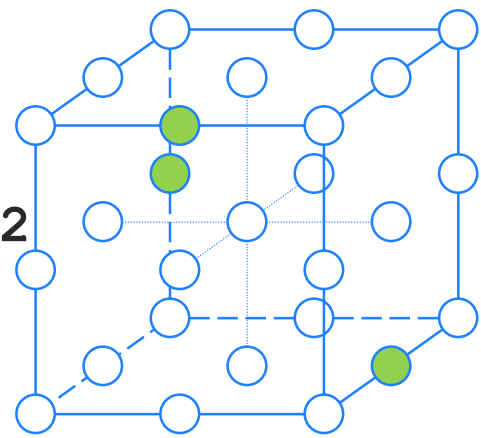
x1	x2	x3	x4
1	1	1	1
1	2	2	2
1	3	3	3
2	1	2	3
2	2	3	1
2	3	1	2
3	1	3	2
3	2	1	3
3	3	2	1



x1 = 1

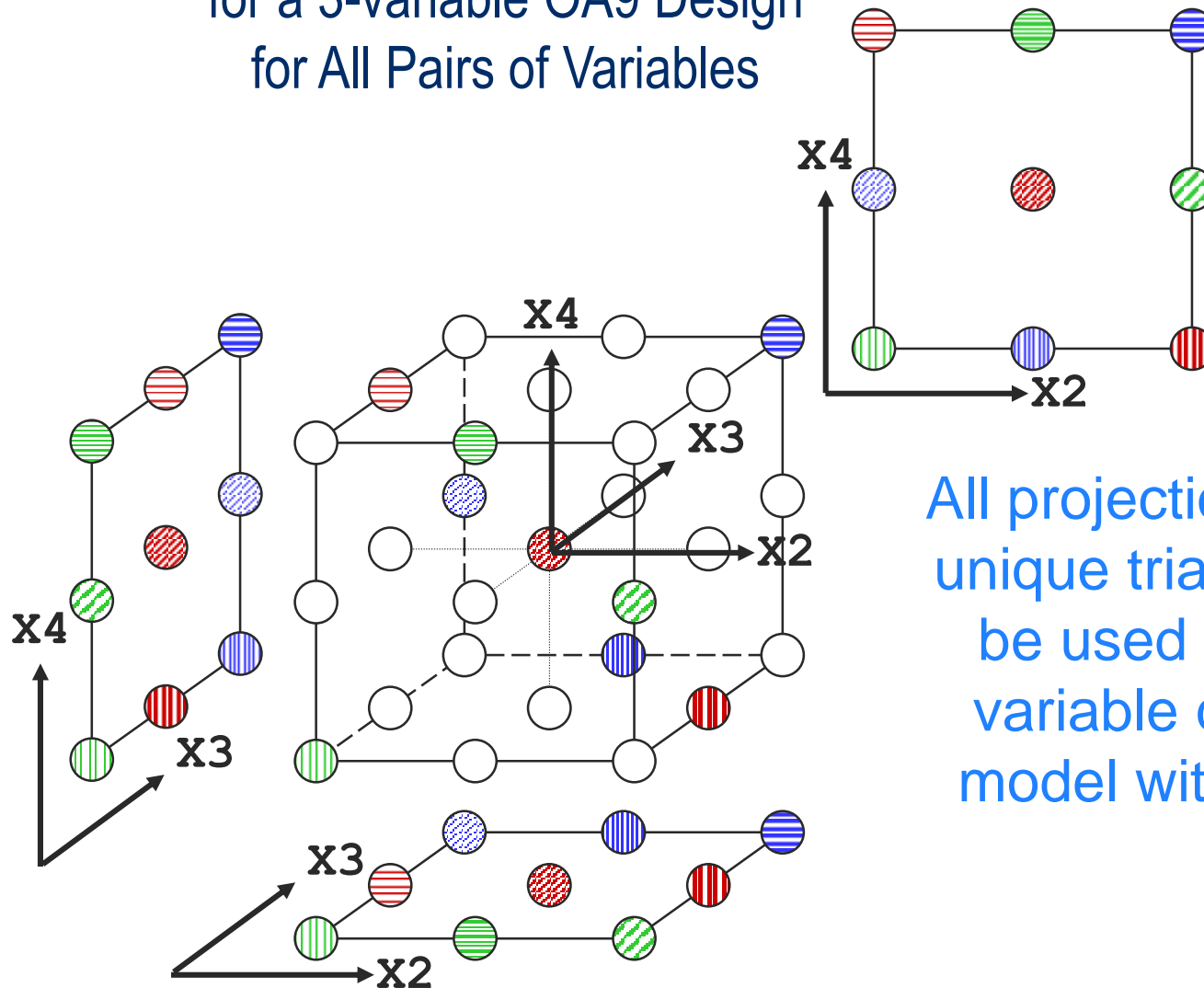


x1 = 2



x1 = 3

# Projection of Trial Locations for a 3-variable OA9 Design for All Pairs of Variables



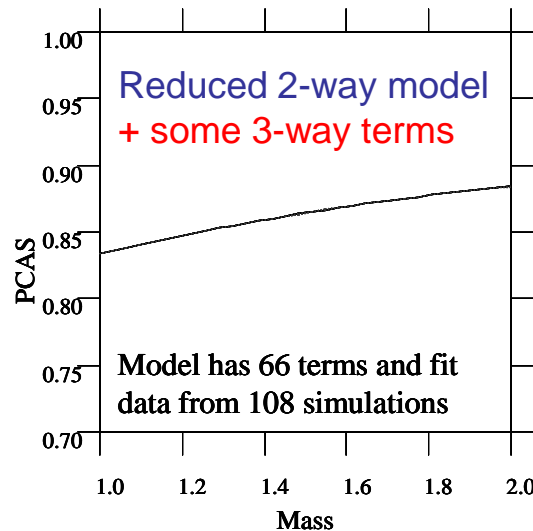
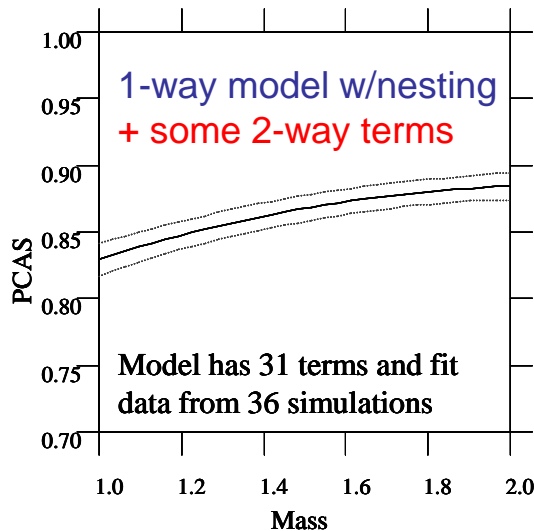
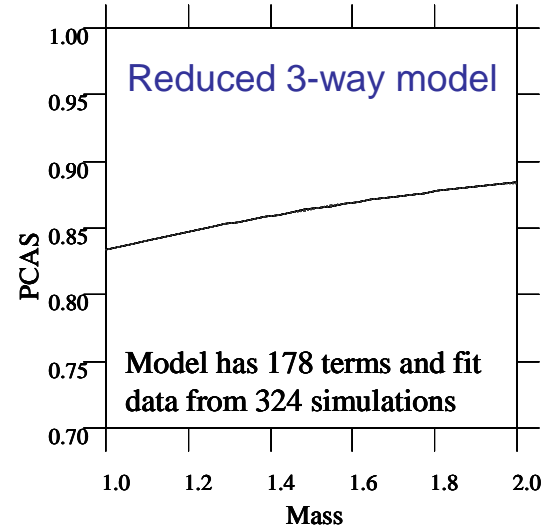
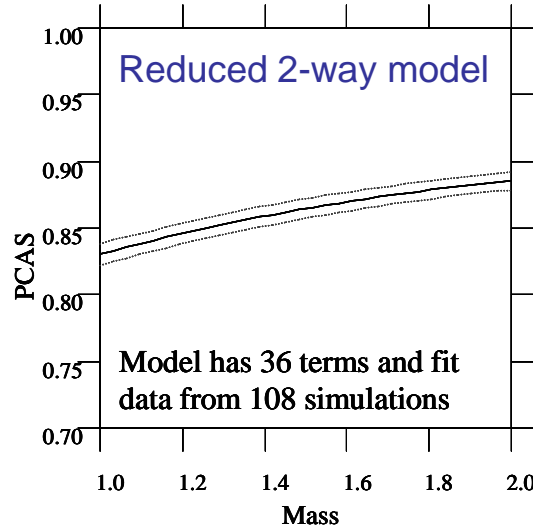
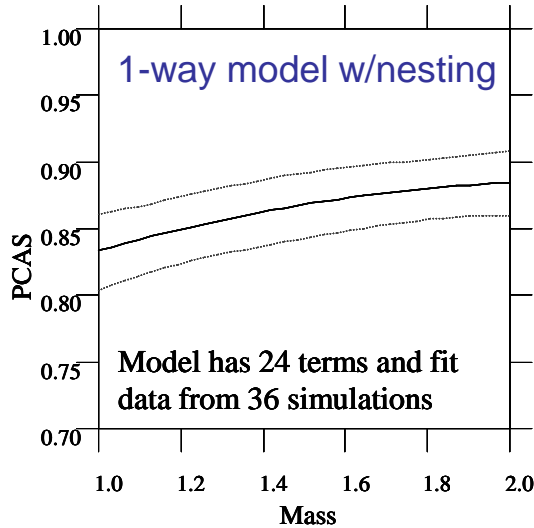
All projections have 9 unique trials that can be used to fit a 2-variable quadratic model with 6 terms

- **Textbook**
  - Limited number of catalogued solutions – experimenters frequently change their problem to match available designs
  - Variable settings are in coded units
- **Web sites of designs**
  - Greater number of catalogued solutions – but never all
  - Variable settings are in coded units
- **Custom computer code**
  - Can find solutions for previously un-catalogued cases
  - Variable settings are in coded units (-1, 0, 1)
- **COTS Solution**
  - Textbook and algorithmic code for generating custom designs
  - Variable settings in natural or laboratory units (120, 150, 180)



# Predictions (w/95% Pred. Limits) of PCAS vs. Nested Mass and MunCnt\_Spread for 1-way, reduced 2-way and reduced 3-way models

## Predicted Probability of Casualty (PCAS) vs. Mass – with Mass Treated as a Continuous Variable – for 5 Different Models Fit to 3 Sets of Simulation Data



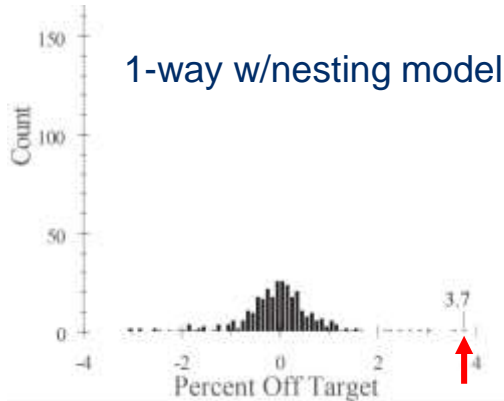
Five other variables were held constant at these settings:

Agent = R  
Season = F  
Time = 12  
HOR = 0  
#TBM &  
Spread Radius = 1

— Predicted Mass  
..... 95% Prediction Limits

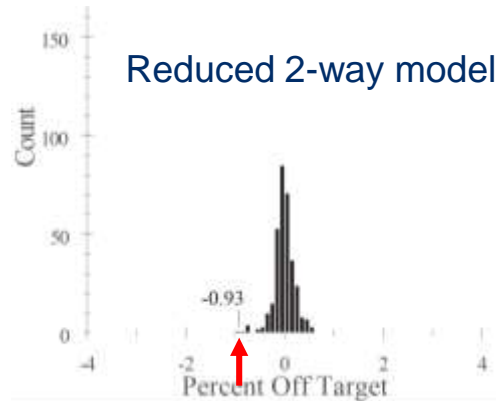


36 trials



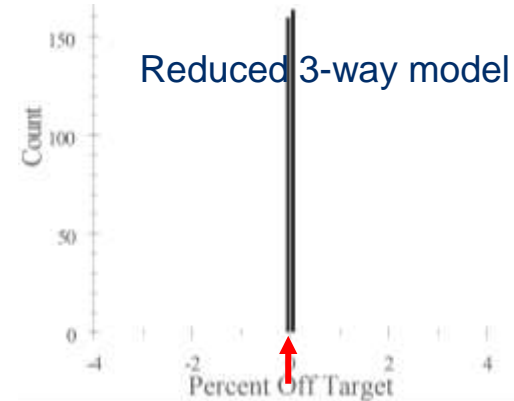
Worst Case = 3.7%  
Half of Cases < 0.37%

108 trials

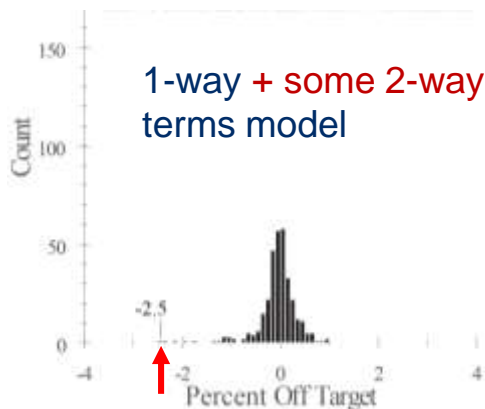


Worst Case = -0.93%  
Half of Cases < 0.11%

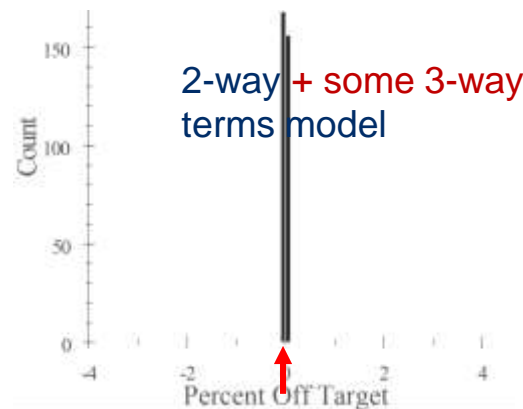
324 trials



Worst Case = -0.0081%  
Half of Cases < 0.0007%



Worst Case = -2.5%  
Half of Cases < 0.16%



Worst Case = -0.0251%  
Half of Cases < 0.0010%

Factor Sparsity states only a few variables will be active in a factorial DOE

Effect Heredity states significant interactions will only occur if at least one parent is active

See Wu & Hamada, p. 112

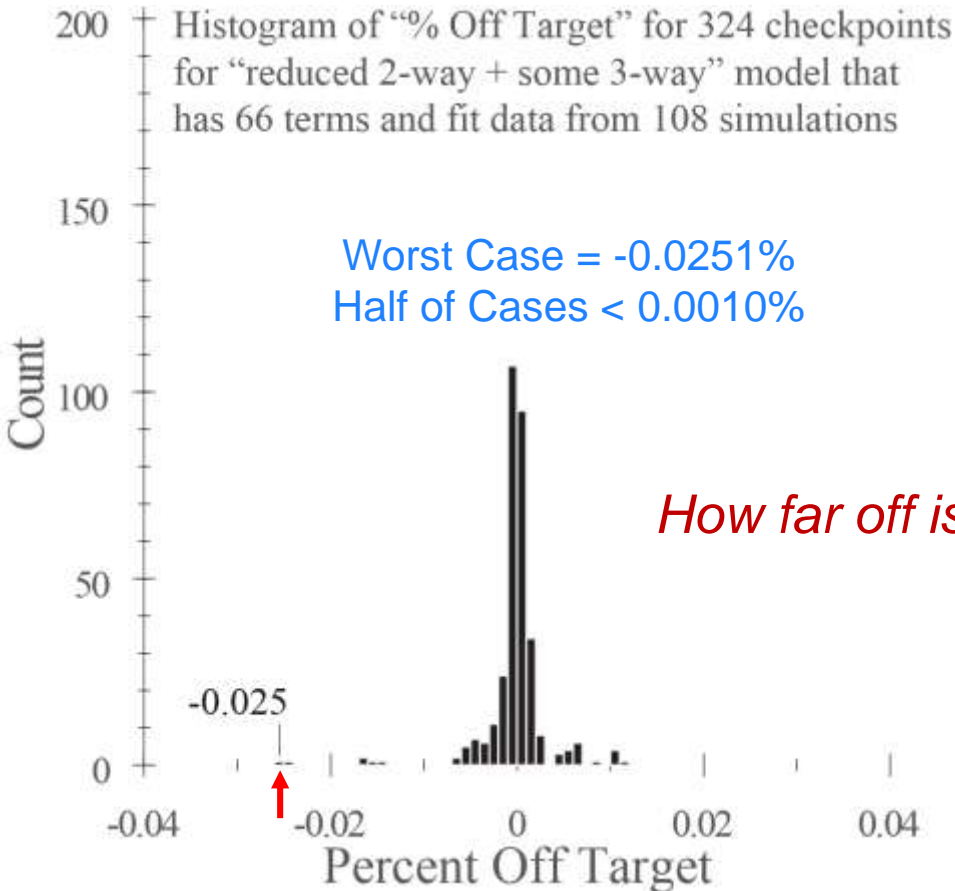
108 trials

324 trials

Higher Resolution (100X) Histograms of the “Percent Off Target” that Response Predictions Fell Relative to 324 Checkpoint Observations

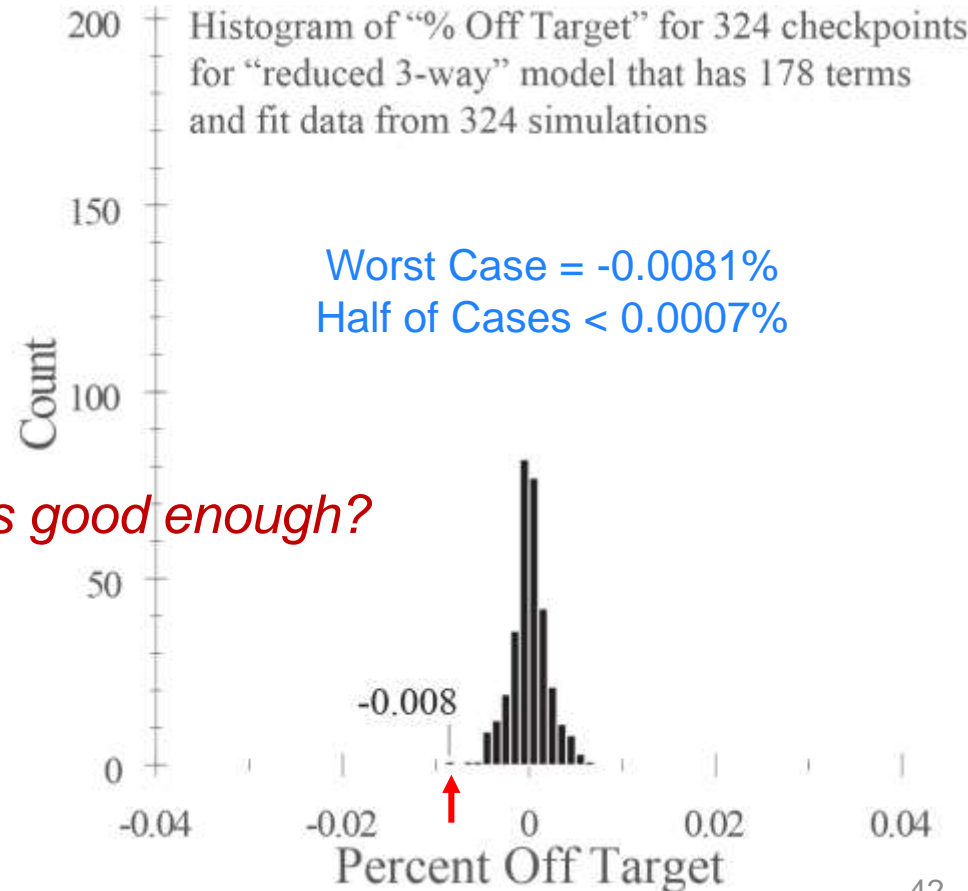
Histogram of “% Off Target” for 324 checkpoints for “reduced 2-way + some 3-way” model that has 66 terms and fit data from 108 simulations

Worst Case = -0.0251%  
Half of Cases < 0.0010%



Histogram of “% Off Target” for 324 checkpoints for “reduced 3-way” model that has 178 terms and fit data from 324 simulations

Worst Case = -0.0081%  
Half of Cases < 0.0007%



*How far off is good enough?*



# More Complexity - Better Fit Fewer "Extraneous" Terms - Better Fit

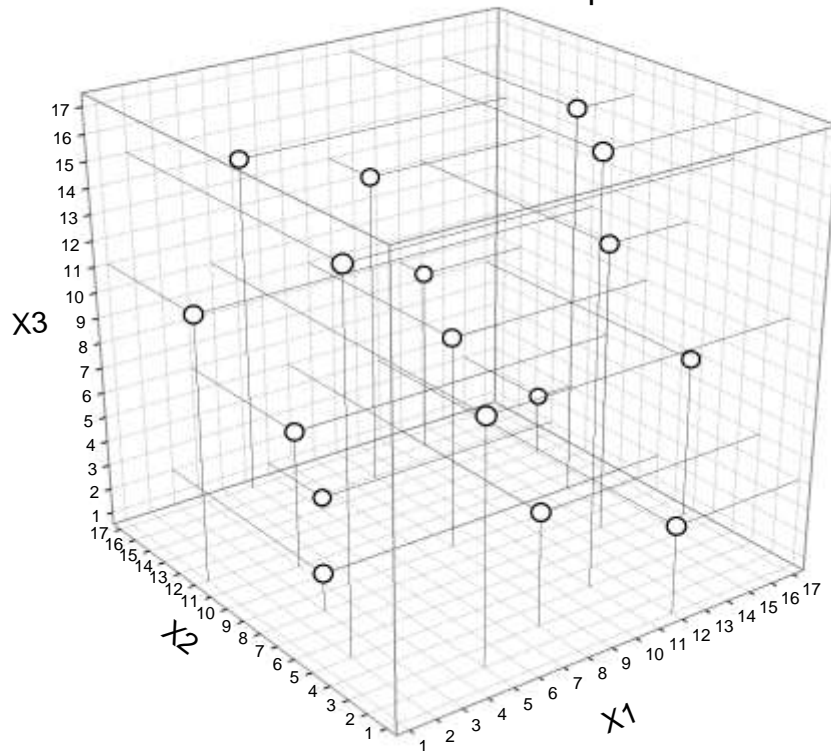
## Error Estimates for 8 Models Fit to Data Sets of 36, 108, & 324 Observations

<b>Number of Trials</b>	<b>Model Used to Fit Data</b>	<b>Number of Model Terms</b>	<b>Residual SD (model error from data used in fit)</b>	<b>Cross-Validation RMS ("one-left out" error from data used in fit)</b>	<b>Checkpoint RMS (model error from 324 data values NOT used in fit)</b>	<b>Adjusted R-squared</b>
36	1-way	14	0.043623	0.055802	0.037217	0.977
36	1-way w/nesting	24	0.026557	0.047269	0.035424	0.992
36	1-way w/nesting + some 2-way	31	0.008212	0.025188	0.016153	0.999
108	2-way	79	0.011197	0.022207	0.010772	0.998
108	reduced 2-way	36	0.008469	0.010933	0.008612	0.999
108	reduced 2-way + some 3-way	66	0.000045	0.000132	0.000179	1.000
324	3-way	242	0.000039	0.000078	0.000083	1.000
324	reduced 3-way	178	0.000037	0.000058	0.000064	1.000

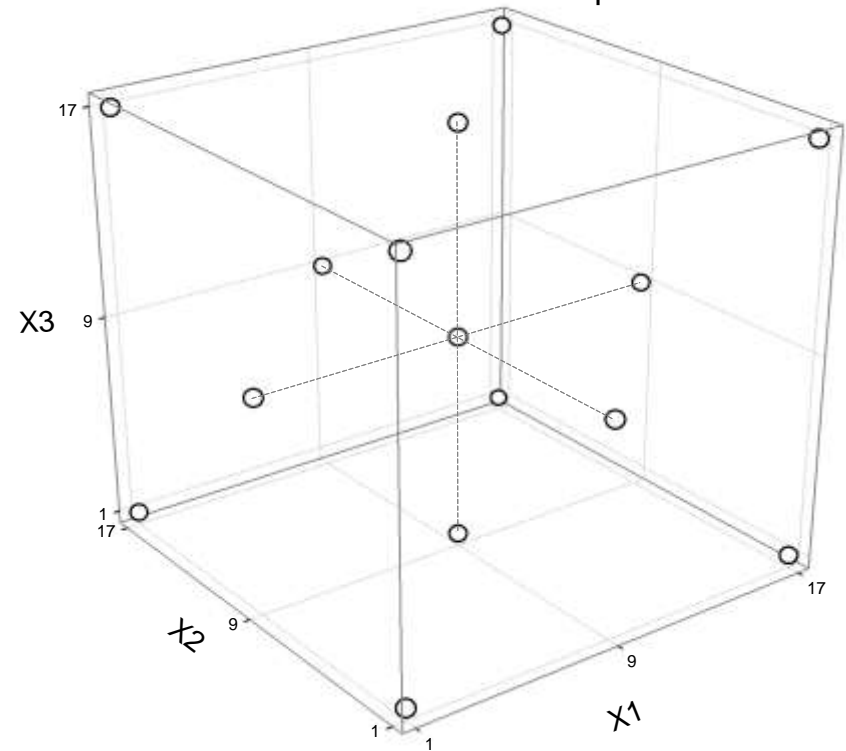
Higher Cross-Validation RMS May Be an Indicator of "Over Fitting"

- Possible to get the 80% to 95% solution with less than 20% of the brute force running of all factor combinations
- Use of “factor sparsity” and “effect heredity” principles can help to get more information than the design was originally built to support
- Next stage trials can first be used as checkpoints for previous stages
- With improved efficiency over running all combinations, more factors can be studied with the same resources

Space-Filling Design  
for 3 Variables with 17 Unique Trials



Response-Surface Design  
for 3-Variables with 15 Unique Trials



Rather than emphasizing high leverage trials (“corners”) for a simple polynomial model, space-filling designs “spread” their trials more uniformly through the space to better capture the local complexities of the simulation model.

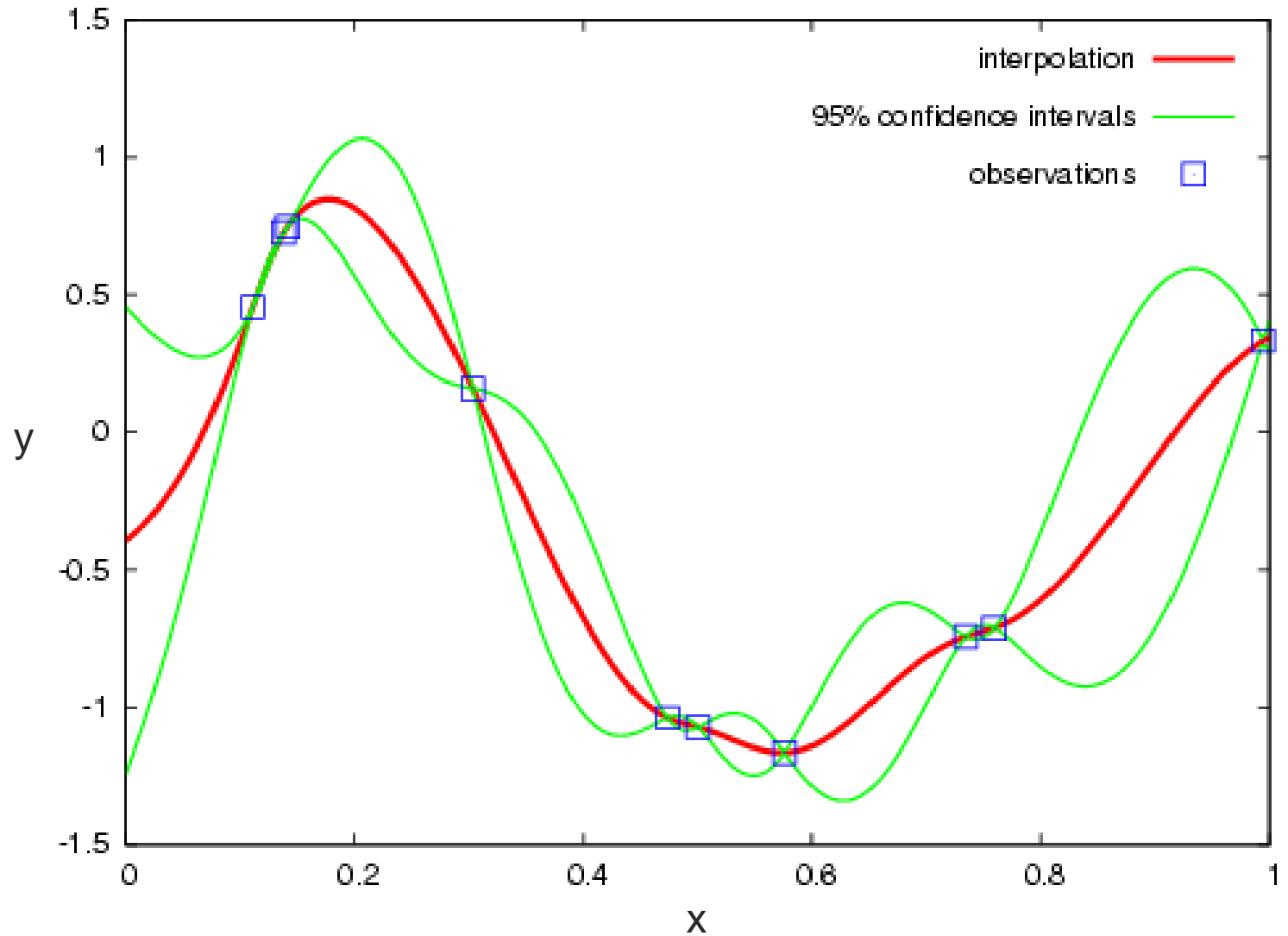


# 29 CFD Simulations Run – 17 Used to Metamodel & 12 Used as Checkpoints

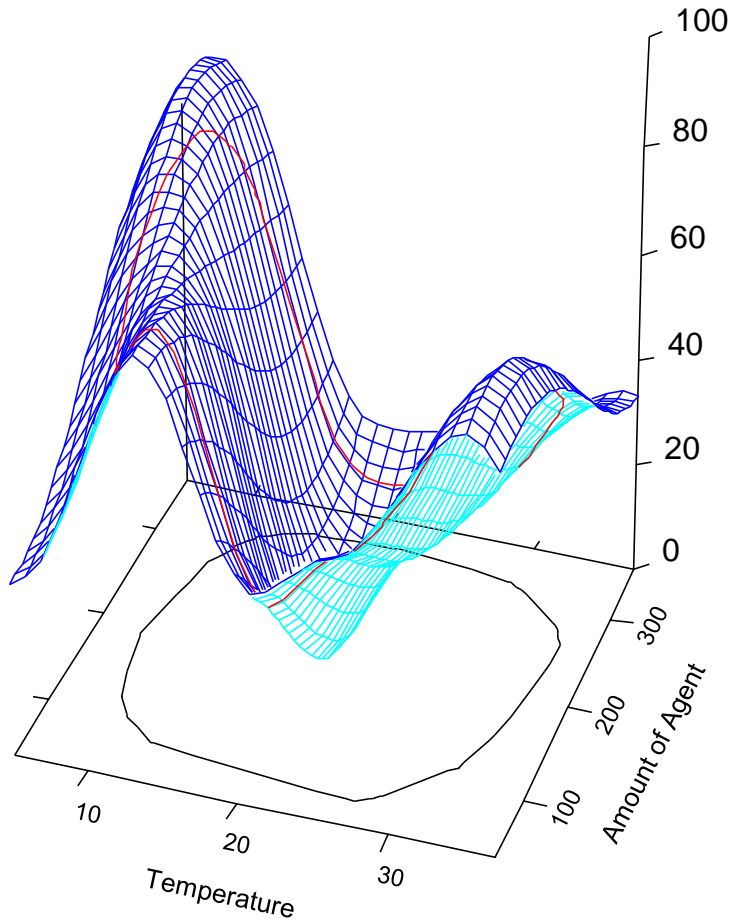
17-trial Orthogonal Latin Hypercube (OLH) space-filling design settings used for creating the metamodel

12-trial Plackett-Burman screening design settings used as checkpoints – half just inside and half just outside design boundary (convex hull)

Trial	Time of Day	Temperature	Wind Speed	Wind Direction	Relative Humidity	Cloud Cover	
1	505	37	5.3	247.5	30	0.92	
2	165	13	5.6	281.25	10	0.32	
3	250	19	1.7	225	60	0.8	
4	335	25	2.9	360	55	0.14	
5	1100	35	3.5	202.5	35	0.02	- Min
6	1440	15	3.2	326.25	15	0.74	
7	930	11	6.2	236.25	80	0.44	
8	845	33	5	348.75	75	0.62	
9	760	21	3.8	270	50	0.5	- Mid
10	1015	5	2.3	292.5	70	0.08	
11	1355	29	2	258.75	90	0.68	
12	1270	23	5.9	315	40	0.2	
13	1185	17	4.7	180	45	0.86	
14	420	7	4.1	337.5	65	0.98	- Max
15	80	27	4.4	213.75	85	0.26	
16	590	31	1.4	303.75	20	0.56	
17	675	9	2.6	191.25	25	0.38	
18	972.5	26	3.05	298.125	62.5	0.65	Inside
19	547.5	16	4.55	241.875	62.5	0.65	Outside
20	972.5	26	3.05	241.875	37.5	0.65	Outside
21	547.5	26	4.55	298.125	37.5	0.35	Outside
22	972.5	16	4.55	298.125	62.5	0.35	Inside
23	547.5	16	3.05	241.875	37.5	0.35	Inside
24	547.5	26	4.55	241.875	62.5	0.65	Outside
25	972.5	16	4.55	298.125	37.5	0.65	Inside
26	547.5	26	3.05	298.125	62.5	0.35	Inside
27	547.5	16	3.05	298.125	37.5	0.65	Outside
28	972.5	16	3.05	241.875	62.5	0.35	Outside
29	972.5	26	4.55	241.875	37.5	0.35	Inside



## 10-Variable Metamodel Prediction



## Off-Axis Variable Settings

Time wrt Sunset = 360  
 Wind Speed = 3.8  
 Wind Direction = 270  
 Humidity = 50  
 Cloud Cover = 0.50  
 $\text{Log}_{10}(\text{Duration}) = 1.0$   
 Latitude (coded) = 17  
 Longitude (coded) = 17

NOTE: This is a plot of Kriging analysis of the 100 integers between 0 and 99 randomly assigned to 100 space-filling design trials.

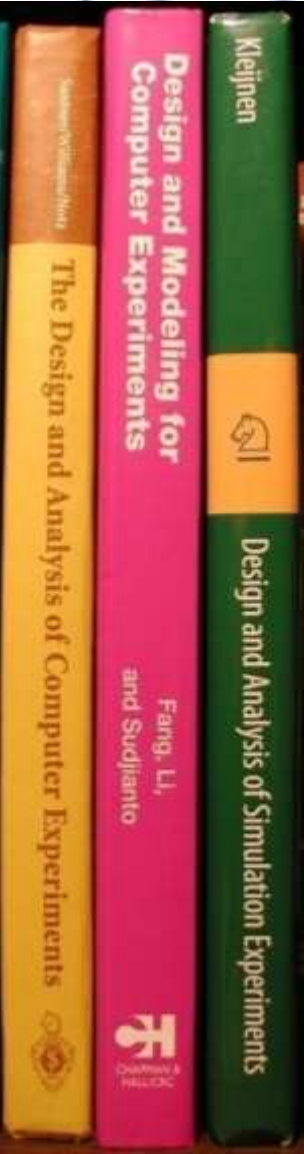
The “noise” has been fit perfectly!  
 This is why one should only use this technique with non-stochastic or nearly non-stochastic data!



- **Design and Analysis of Computer Experiments**  
 Sacks, J., Welch, W.J., Mitchell, T.J. and Wynn, H.P.  
*Statistical Science* 4. 409-423, 1989

- Textbooks on this topic include:

- Santner, T. J., Williams, B. J., and Notz, W. I. (2003), *The Design and Analysis of Computer Experiments*, Springer, New York
- Fang, K. T., Li, R. Z., and Sudjianto, A. (2005), *Design and Modeling for Computer Experiments*, Chapman & Hall/CRC Press, New York
- Kleijnen, J. P. C. (2008), *DASE: design and analysis of simulation experiments*. Springer, New York.



- JMP® (called Gaussian Process modeling)
- ECHIP® (called Smoothing analysis)
- SYSTAT® (called Kriging analysis)
- Matlab® Toolbox Modules
  - Design and Analysis of Computer Experiments (DACE)
  - SURrogate MOdeling (SUMO)
    - Contains DACE as well as another Kriging tool and many other surrogate modeling methods
- PErK (code available from authors of 2003 text by Santner, et. al.)
- “Blind” Kriging – R code potentially available from GA Tech
- The Gaussian Processes Website: <http://www.gaussianprocess.org>
- Code to do Bayesian Hierarchical Gaussian Process (BHGP) modeling by combining simulation and real experimental data is available from Prof. Peter Qian of the University of Wisconsin
- Code for Nested and Sliced Latin Hypercube Designs also available from Prof. Qian..

- <http://harvest.nps.edu/>  
 The Simulation Experiments & Efficient Design (SEED) Center for Data Farming at Naval Postgraduate School
  - Designs
    - Nearly Orthogonal Latin Hypercubes (NOLH) and
    - Resolution V, Fractional Factorials for many factors
  - Agent-Based Simulation Software
    - Pythagoras
    - MANA (Map Aware Non-uniform Automata)
  - Many Papers for Download and Links to INFORMS and WSC
- <http://www.research.att.com/~njas/oadir/index.html>  
 Library of Orthogonal Arrays maintained by Neil J.A. Sloane
- <http://support.sas.com/techsup/technote/ts723.html>  
 Library of Orthogonal Arrays maintained by Warren F. Kuhfield

- **Blind Kriging: A New Method for Developing Metamodels,**  
Joseph, V.R., Hung, Y., and Sudjianto, A.,  
*ASME Journal of Mechanical Design*, 130, 031102-1-8, 2008
- **Gaussian Process Models for Computer Experiments  
With Qualitative and Quantitative Factors,**  
Qian, P.Z.G., Wu, H., and Wu, C.F.J.,  
*Technometrics*, 50 (4), 383-396, 2008
- **Bayesian Hierarchical Modeling for Integrating Low-Accuracy  
and High-Accuracy Experiments,**  
Qian, P. Z. G. and Wu, C. F. J.,  
*Technometrics*, 50 (2), 192-204, 2008
- **Regression-Based Inverse Distance Weighting for Multivariate  
Interpolation,**  
Joseph, V.R., and Kang, L.,  
(submitted) Preprint May 2009
- **Nested Latin Hypecube Designs,**  
Qian, P. Z. G.  
*Biometrika*, 96, 957-970, 2008

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- 19. Zhou, Q., Qian, P. Z. G. and Zhou, S (2010), [A Simple Approach to Emulation for Computer Models With Qualitative and Quantitative Factors](#), *Technometrics*, Accepted.
- 18. He, X. and Qian, P. Z. G. (2010), [Nested orthogonal array based Latin hypercube designs](#), *Biometrika*, Accepted.
- 17. Xu, X., Haaland, B. and Qian, P. Z. G. (2010), [Sudoku based space-filling designs](#), *Biometrika*, Accepted.

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- 16. Tang, Q. and Qian, P. Z. G. (2010), ``Enhancing the Sample Average Approximation Method with U Designs," *Biometrika*, 97, 947-960.
- 15. Haaland, B., Min, W., Qian, P. Z. G. and Amemiya, Y. (2010), ``A Statistical Approach to Thermal Management of Data Centers under Steady State and System Perturbations," *Journal of the American Statistical Association*, 105, 1030-1041.
- 14. Qian, P. Z. G. and Ai, M. (2010), ``Nested Lattice Sampling: A New Sampling Scheme Derived by Randomizing Nested Orthogonal Arrays," *Journal of the American Statistical Association*, 105, 1147-1155.
- 13. Haaland, B. and Qian, P. Z. G. (2010), "An Approach to Constructing Nested Space-Filling Designs for Multi-Fidelity Computer Experiments," *Statistica Sinica*, 20, 1063-1075.
- 12. Wang, F., Hwang, Y., Qian, P. Z. G. and Wang, X. (2010), ``A Statistics-Guided Approach to Precise Characterization of Nanowire Morphology," *American Chemical Society Nano* , 4, 855-862.

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8. Qian, P. Z. G., Tang, B. and Wu, C. F. J. (2009), "Nested Space-Filling Designs for Experiments with Two Levels of Accuracy," *Statistica Sinica*, 19, 287-300.

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4. Negrut, D., Qian, P. Z. G., and Khude, N. (2007), "Building Gaussian Process Based Metamodels Using Variable-fidelity Experiments for Dynamic Analysis of Mechanical Systems," *Proceedings of the 2007 ASME International Mechanical Engineering Congress and Exposition*, Seattle, WA.
3. Qian, Z., Seepersad, C., Joseph, R., Allen, J. and Wu, C. F. J. (2006), "[Building Surrogate Models with Detailed and Approximate Simulations](#)," *ASME Journal of Mechanical Design*, 128, 668-677.
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Deng, X. and Qian, P. Z. G. (2010), Sliced Cross-validation for Efficient Estimation of the Error Rate of a Classification Rule.

Qian, P. Z. G. and Amemiya, Y. (2010), A Structural Equation Method for Modeling Multivariate Temperature Data from Data Center Computer Experiments.

Qian, P. Z. G. (2010), Sliced Latin Hypercube Designs.

Li, J. and Qian, P. Z. G. (2010), [Construction of Nested \(Nearly\) Orthogonal Designs for Computer Experiments.](#)

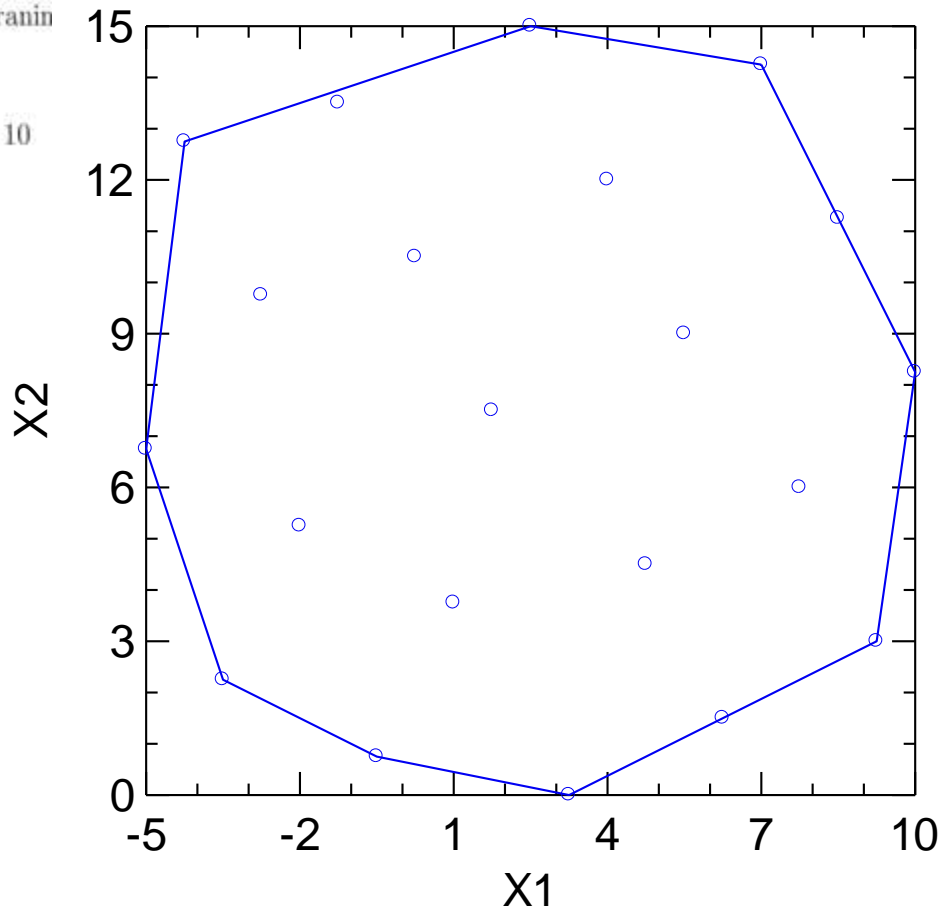
Qian, P. Z. G. and Ai, M. (2010), [Nested Asymmetric Lattice Samples.](#)

## C.3 Examples

The following examples demonstrate many possible uses of PErK. The responses for these examples are based on the *Branin function*. The Branin function is the real-valued function of two variables

$$y_B(x_1, x_2) = \left(x_2 - \frac{5.1}{4\pi^2} x_1^2 + \frac{5}{\pi} x_1 - 6\right)^2 + 10 \left(1 - \frac{1}{8\pi}\right) \cos(x_1) + 10$$

Trial	x1	x2	Y_B
1	7.75	6	35.80951
2	1	3.75	14.86287
3	10	8.25	31.41880
4	4.75	4.5	19.87899
5	2.5	15	141.88566
6	-3.5	2.25	99.43335
7	3.25	0	3.88973
8	-5	6.75	97.47380
9	-4.25	12.75	6.27060
10	6.25	1.5	19.85914
11	8.5	11.25	95.50587
12	7	14.25	181.74214
13	-0.5	0.75	49.39445
14	-2	5.25	23.13762
15	0.25	10.5	43.09524
16	9.25	3	2.82392
17	-2.75	9.75	3.61474
18	5.5	9	75.79100
19	4	12	104.11175
20	-1.25	13.5	43.33586
21	1.75	7.5	23.39797





## C.3 Examples

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Plot from textbook of Branin Function

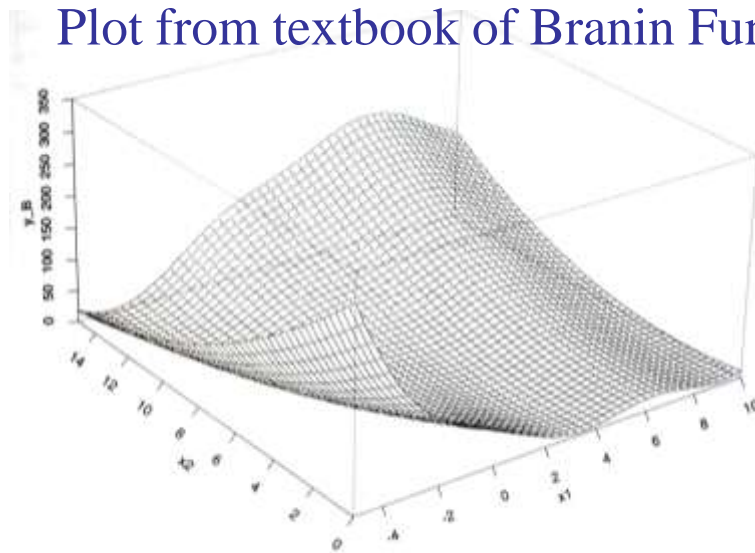
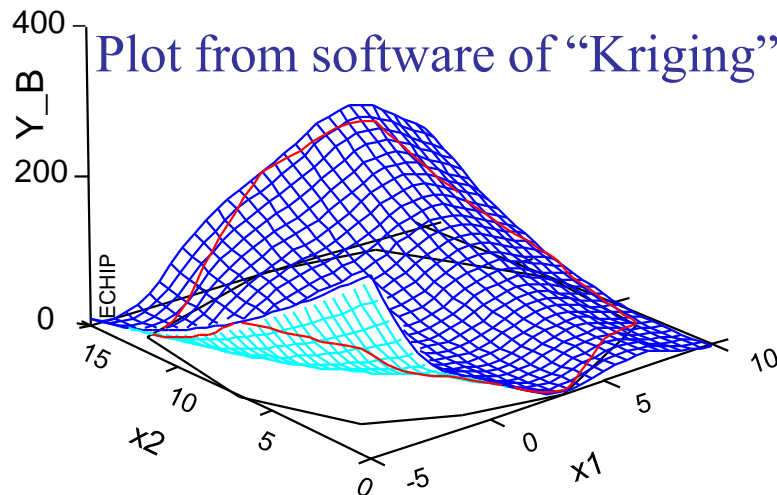


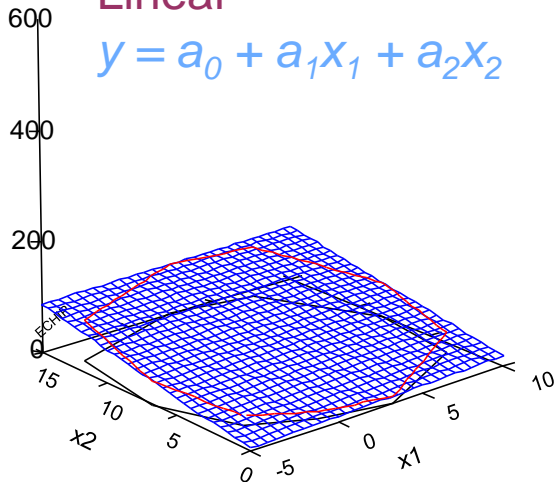
FIGURE C.1. The Branin function on  $[-5, 10] \times [0, 15]$

Plot from software of “Kriging” fit



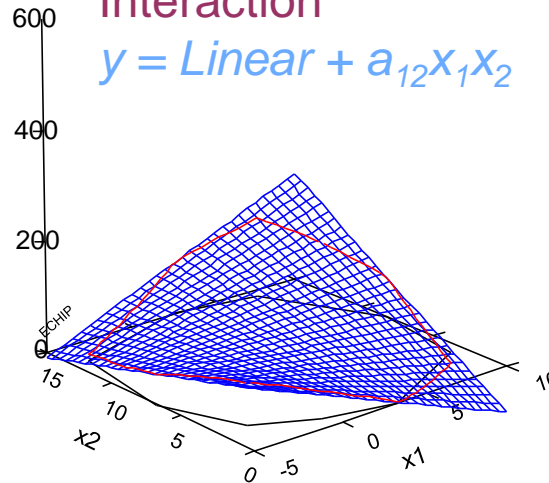
## Linear

$$y = a_0 + a_1x_1 + a_2x_2$$



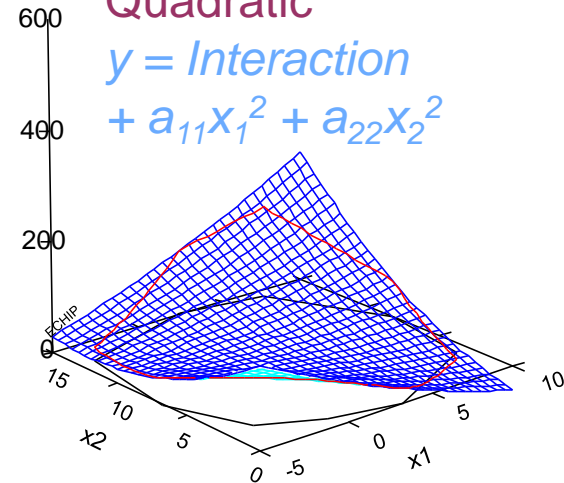
## Interaction

$$y = \text{Linear} + a_{12}x_1x_2$$



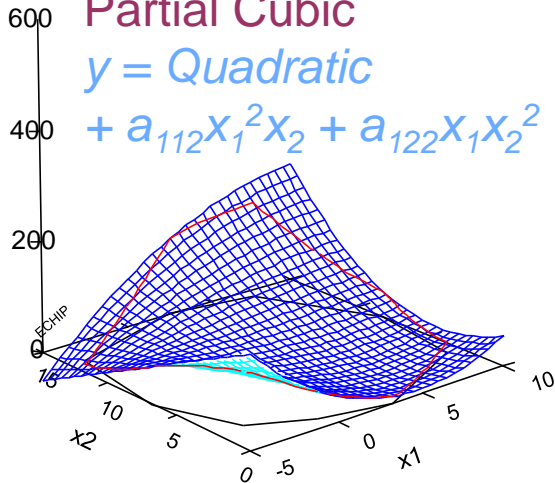
## Quadratic

$$y = \text{Interaction} + a_{11}x_1^2 + a_{22}x_2^2$$



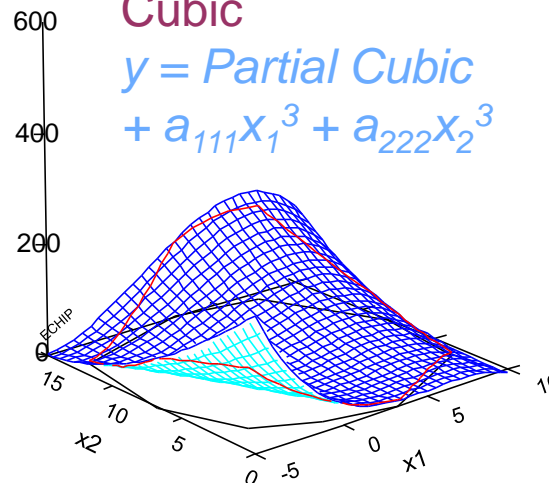
## Partial Cubic

$$y = \text{Quadratic} + a_{112}x_1^2x_2 + a_{122}x_1x_2^2$$

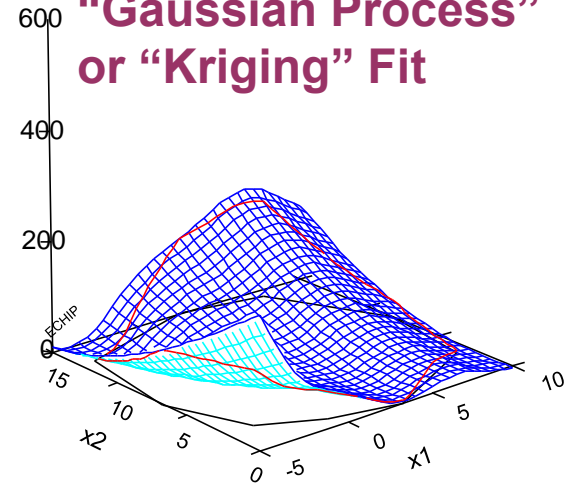


## Cubic

$$y = \text{Partial Cubic} + a_{111}x_1^3 + a_{222}x_2^3$$



## "Gaussian Process" or "Kriging" Fit



The full *cubic* model closely approximates the Branin function, but still cannot capture the ripples seen in the fit using the Kriging method.

- Branin function example is trivial. With 2 control variables the full cubic model has 10 terms.
- What if your simulation has 10 control variables?
  - Full cubic model has 166 terms!
  - And still may not be complex enough to accurately approximate the simulation

We wanted to not just do sensitivity analysis of the factors, but **provide an interactive surrogate model of the long-running simulation so that analysts could evaluate “what if?” scenarios.**

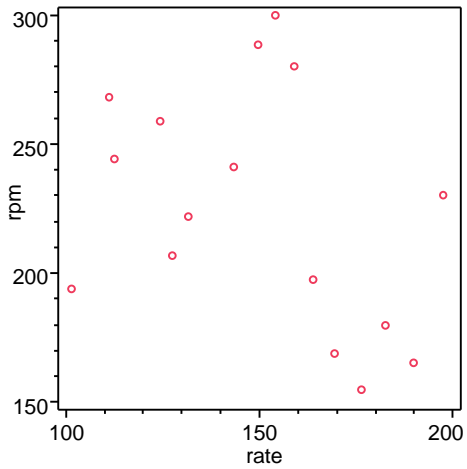
The problem was that the Computational Fluid Dynamics models we were looking to run could take a week on a single CPU or **12 hours on 50 CPU cluster.** With on the order of 10 factors we expected to need to run on the order of **100 simulations.** **This meant it could be weeks or months before we could start our analysis.**

**Nested Latin Hypercube Designs gave us a way to start analyzing data after about the first 20% of the simulations were run.** We also wanted to be able to run just enough simulations to achieve a surrogate model accuracy of 90%.

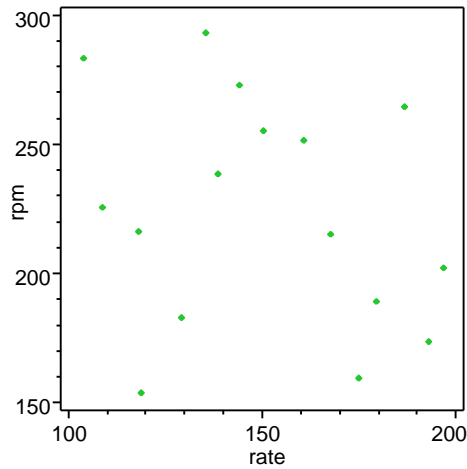


# Projections of Trial Locations in 2 factors for a 10-factor, 128-trial, Nested Latin Hypercube Design\* (NLHD) with 4 Blocks

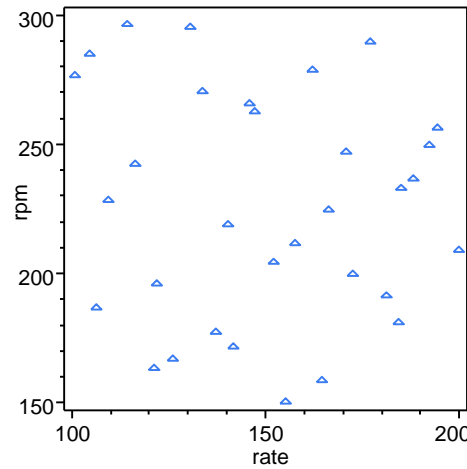
Block 1, 16 trials



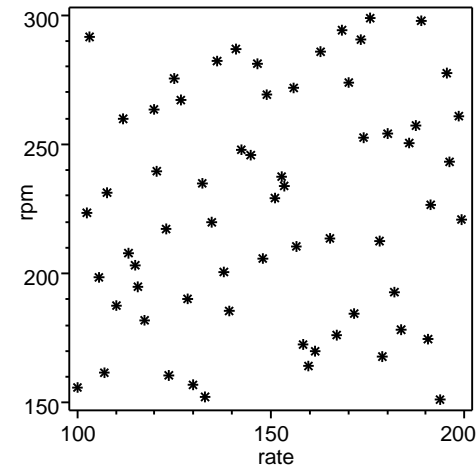
Block 2, 16 trials



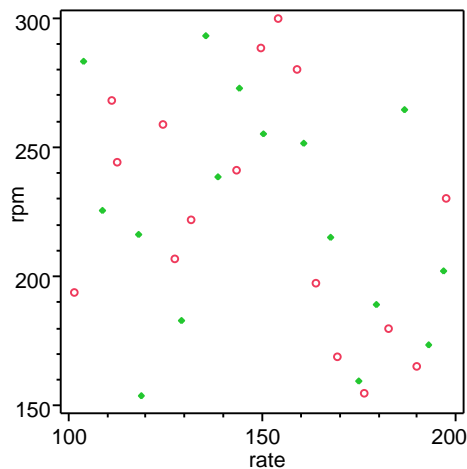
Block 3, 32 trials



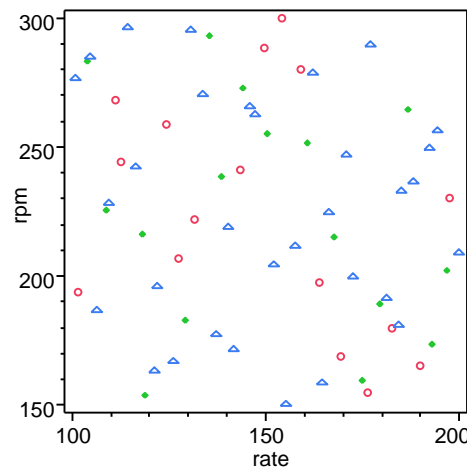
Block 4, 64 trials



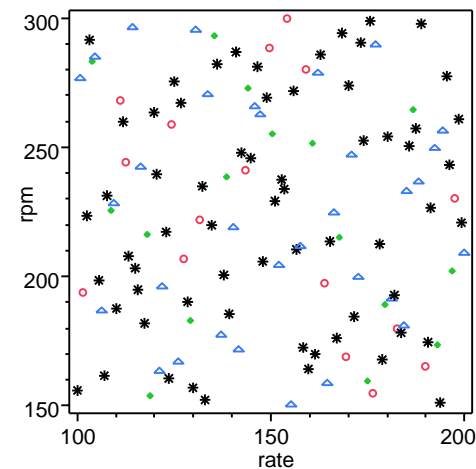
Blocks 1 & 2, 32 trials



Blocks 1, 2 & 3, 64 trials



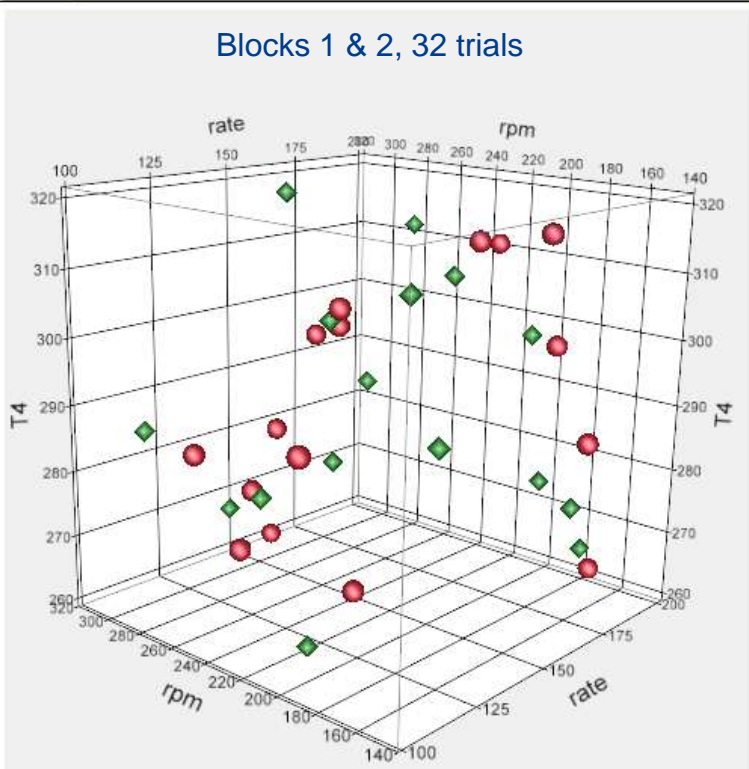
Blocks 1, 2, 3 & 4, 128 trials



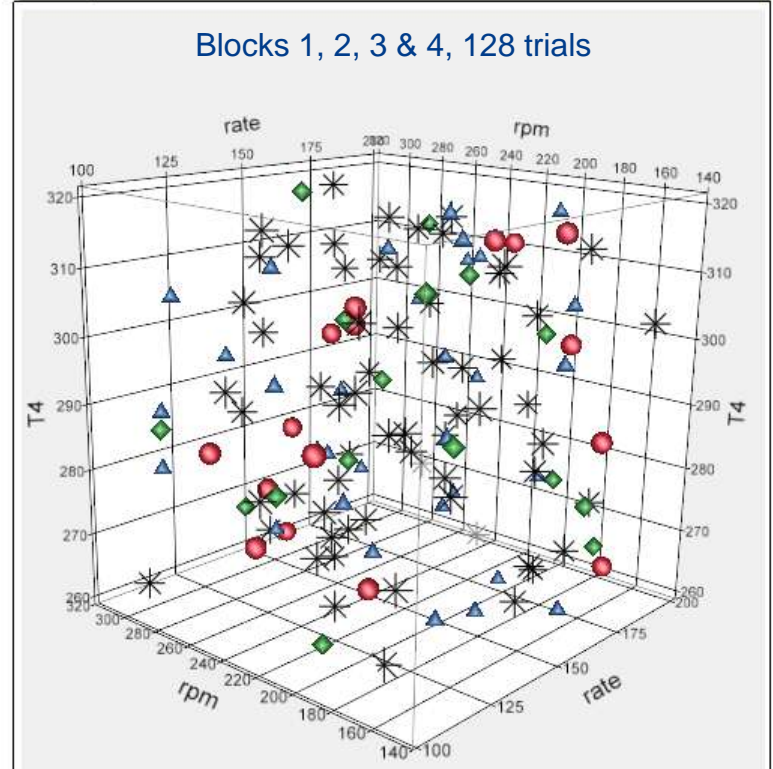
Running totals of blocks are also Latin Hypercube Designs

\*Generated with Matlab Code Received from Prof. Peter Qian of U of Wi.

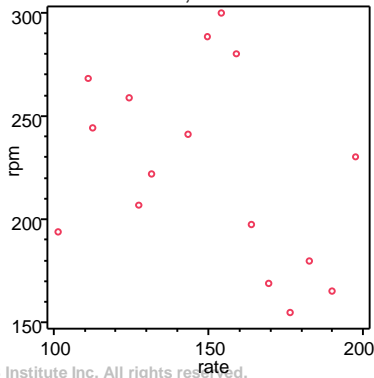
Scatterplot 3D



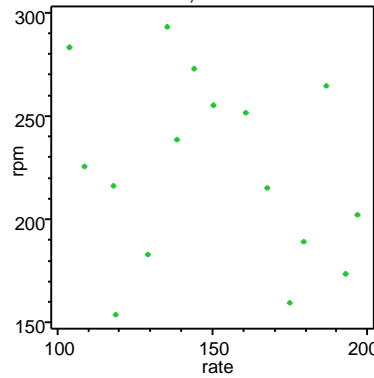
Scatterplot 3D



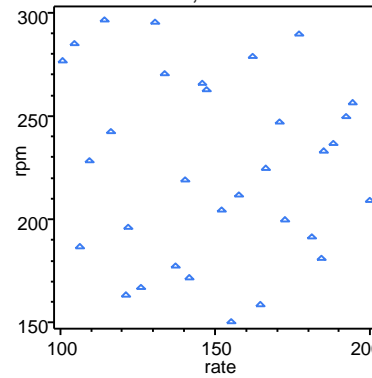
Block 1, 16 trials



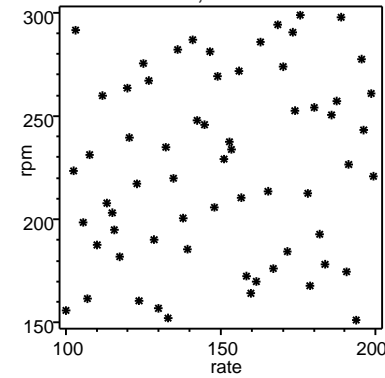
Block 2, 16 trials



Block 3, 32 trials

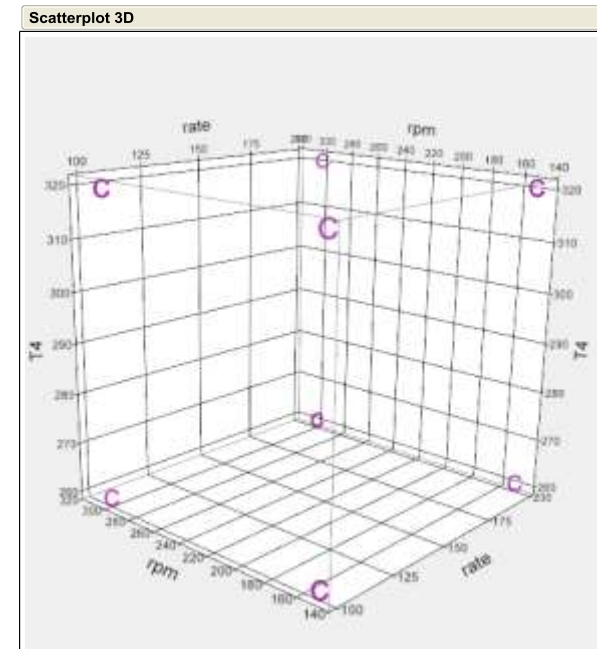
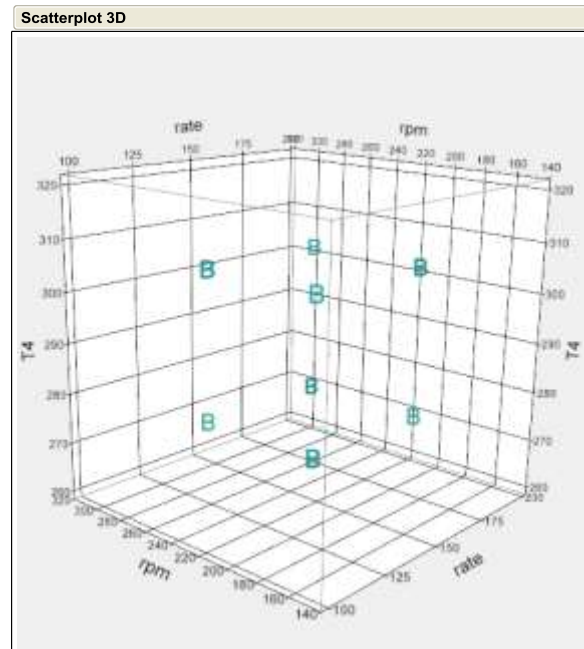
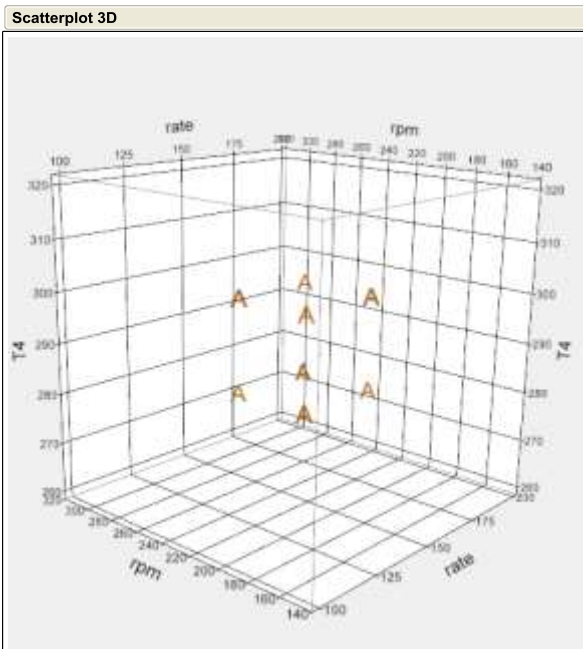


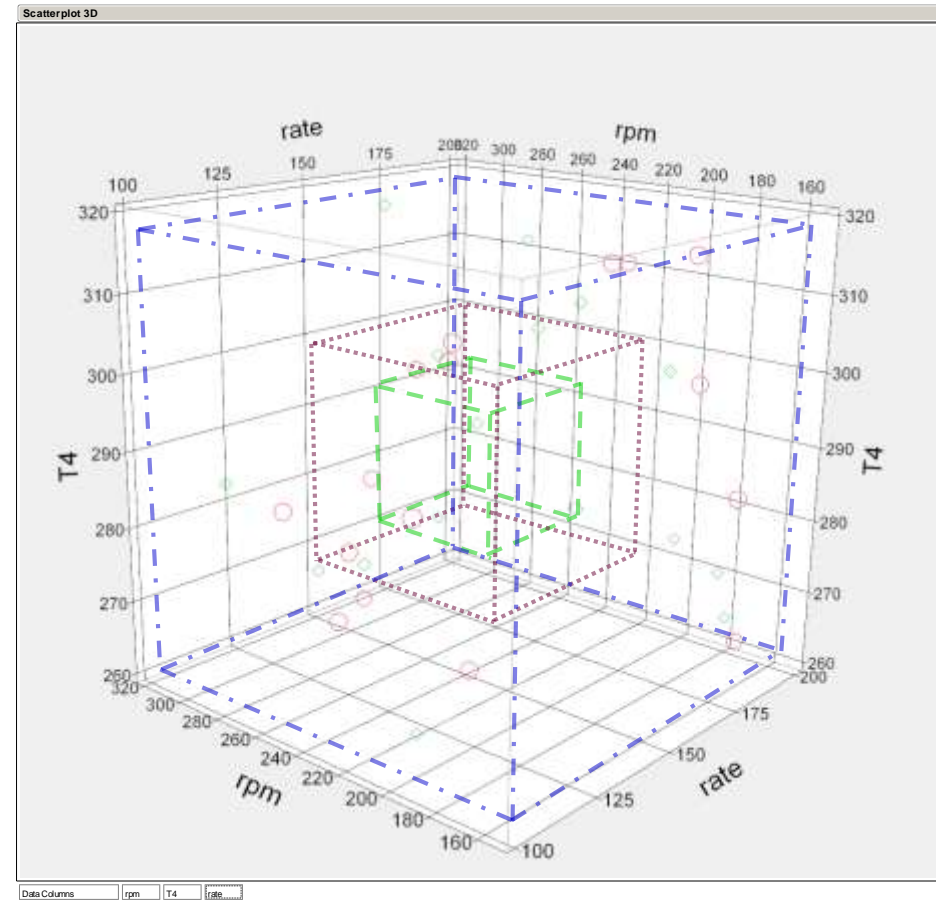
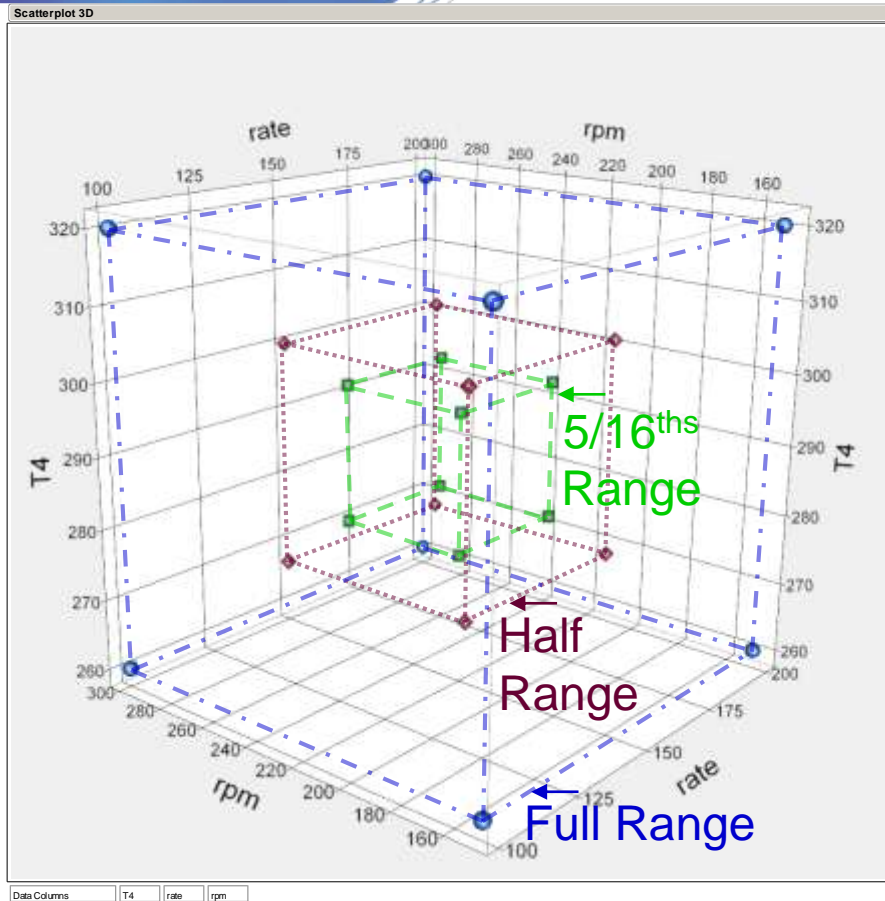
Block 4, 64 trials



The point of running this sequence of blocks is to be able to evaluate the surrogate model after each stage to see how accurately it is predicting observed values of 3 sets of checkpoint trials. If it proves to be sufficiently accurate, then subsequent blocks of simulation trials need not be run.

Without the NLHD approach one has to choose the “right” size space-filling design in order to get useful results. If you choose too small a design, one has to start over with a larger design.



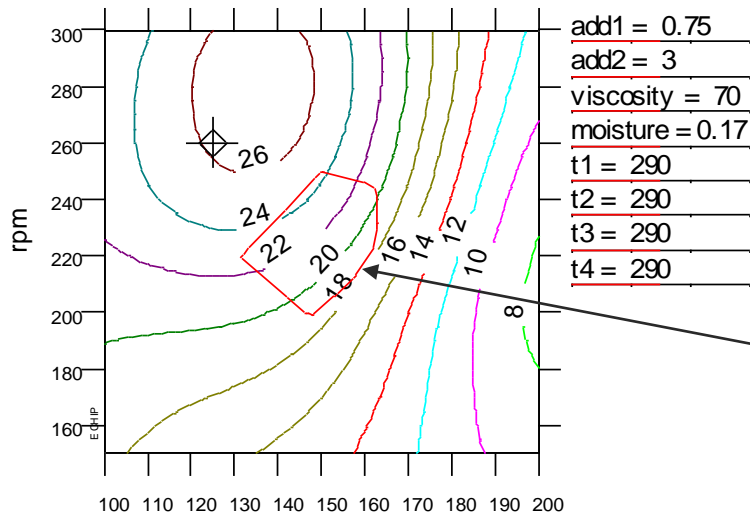


In the full design space over 10 factors there are 10 dimensions and 1024 corners. The 12 trials in a Plackett-Burman design populate only about 0.1% of these combinations of settings.

Today, I would use Definitive Screening Design with 21 trials for 10 factors but also get information at midpoints of each factor.



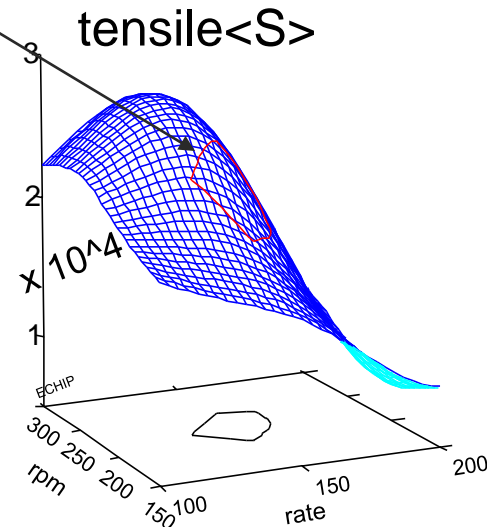
tensile<S> x 10<sup>3</sup>



rate		
rate=125.00		rpm=260.00
Value	LowLimit	HighLimit
26259.40	-1.#J	1.#J

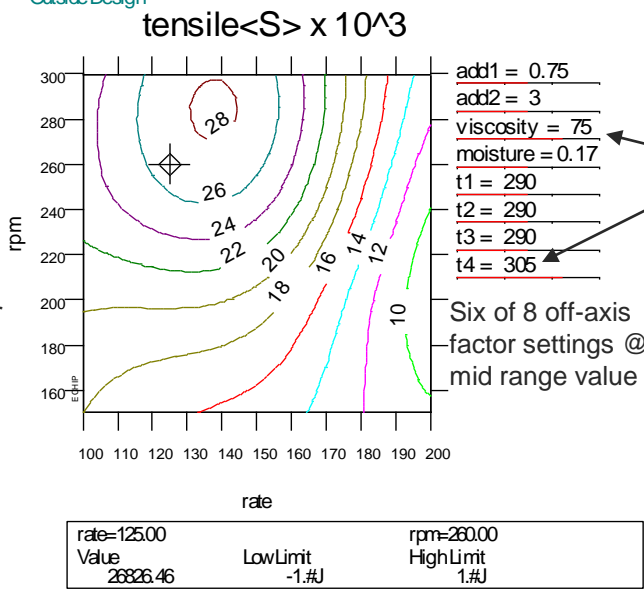
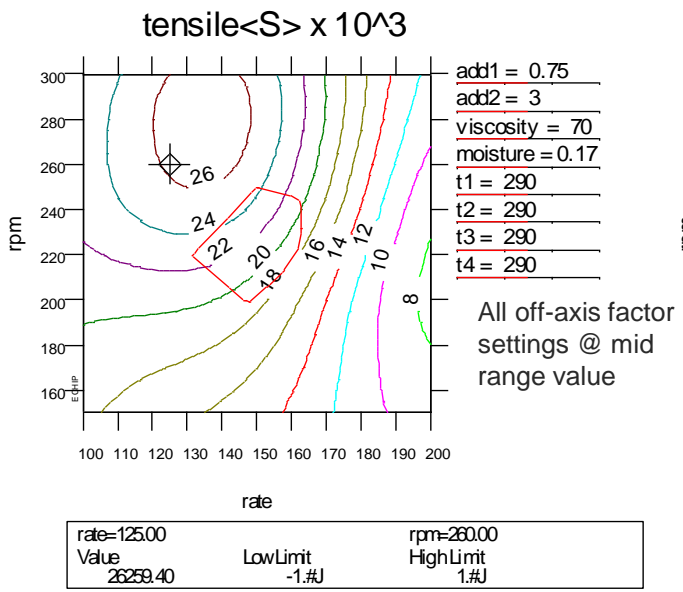
The 10-dimensional design space is only sparsely covered by the initial 16-trial NLHD Block. As a result only a small fraction of the full design region is valid for interpolation with the Kriging analysis.

Red polygon marks boundary between regions of interpolation (inside) and extrapolation (outside). Statistical name for the design boundary is the "Convex Hull."



# NLHD, 16 trials Block 1 only

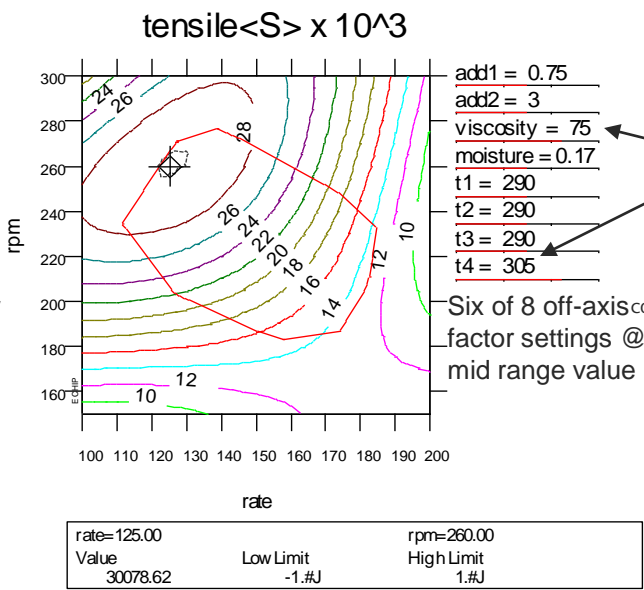
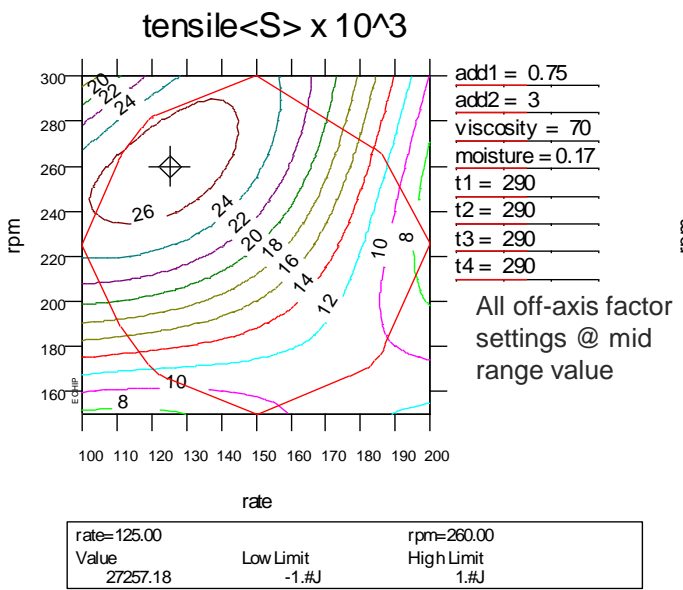
← Note that this entire plot is extrapolation.



viscosity and t4 factor settings @ 75% of range

COEFFICIENTS	SD	P	CONDITION	TERM
18903.8				0 CONSTANT
-6319.34	8425.11	0.4870-	0.559	1 add1
-882.924	1658.5	0.6173-	0.703	2 add2
139.372	143.796	0.3769-	0.820	3 viscosity
-25974.4	19170.6	0.2334	0.813	4 moisture
32.8283	74.8572	0.6793-	0.515	5 t1
21.6944	79.5117	0.7959-	0.488	6 t2
61.0479	88.6668	0.5218-	0.432	7 t3
44.7973	81.6762	0.6070-	0.477	8 t4
-128.299	42.7544	0.0301	0.550	9 rate
96.4185	27.5245	0.0172	0.546	10 rpm

N trials = 16  
Closer to 1.000 the CONDITION is, the closer the term is to being orthogonal



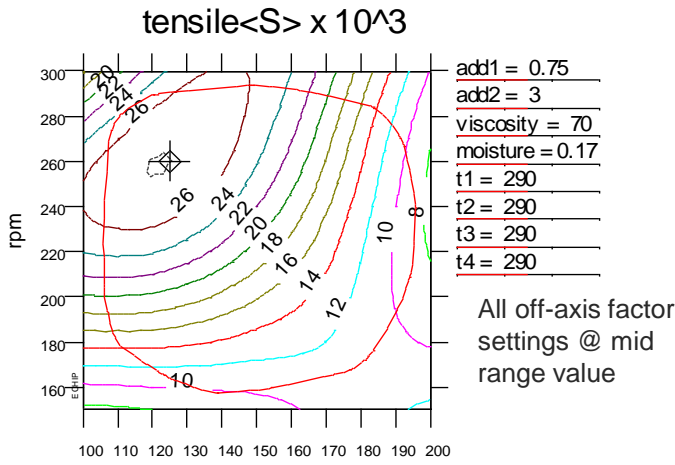
viscosity and t4 factor settings @ 75% of range

COEFFICIENTS	SD	P	CONDITION	TERM
15760.2				0 CONSTANT
-721.328	6720.15	0.9158-	0.958	1 add1
966.423	1647.02	0.5651-	0.974	2 add2
65.0763	163.369	0.6953-	0.986	3 viscosity
4164.81	21651.0	0.8497-	0.989	4 moisture
-7.23148	54.8311	0.8966-	0.973	5 t1
1.39981	54.9105	0.9800-	0.973	6 t2
-38.7242	56.0048	0.4986-	0.951	7 t3
47.8879	54.5957	0.3926-	0.981	8 t4
-49.521	32.9741	0.1515	0.976	9 rate
43.0792	21.8001	0.0646	0.971	10 rpm

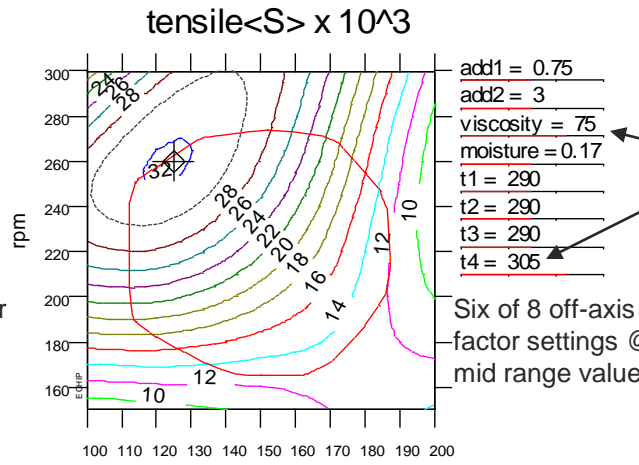
N trials = 28  
Closer to 1.000 the CONDITION is, the closer the term is to being orthogonal

Inclusion of checkpoints – here the 12 over the full range of the factors – increases the size of the design boundary and the volume of interpolation region.

# NLHD, 16+16+32+64=128 Blocks 1, 2, 3 & 4



rate=125.00	rpm=260.00
Value	LowLimit HighLimit
28043.68	1.#R 1.#R

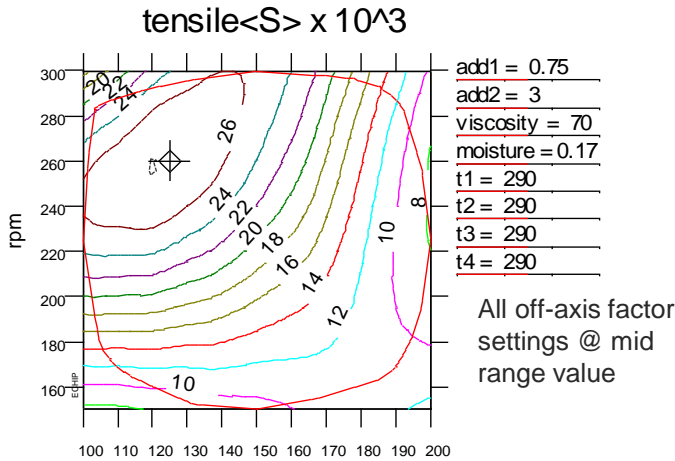


rate=125.00	rpm=260.00
Value	LowLimit HighLimit
32266.97	1.#R 1.#R

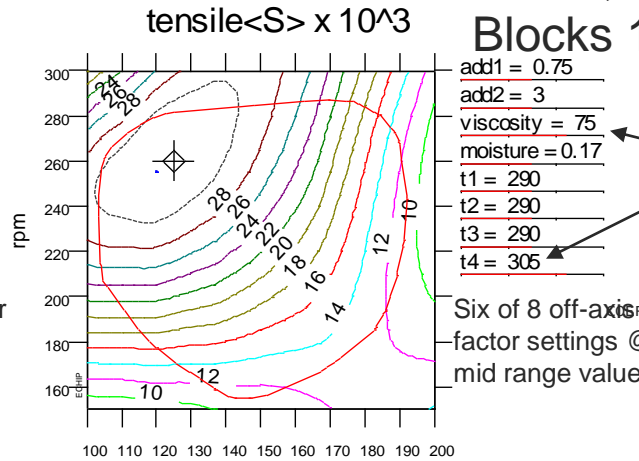
viscosity and t4 factor settings @ 75% of range

COEFFICIENTS	SD	P	CONDITION	TERM
17631.9				0 CONSTANT
-682.306	2233.14	0.7605-	0.945	1 add1
884.808	540.923	0.1046	0.975	2 add2
300.046	54.081	0.0000	0.976	3 viscosity
-3680.05	7202.41	0.6104-	0.977	4 moisture
5.82534	18.0142	0.7470-	0.976	5 t1
-11.1487	18.0473	0.5379-	0.975	6 t2
31.8328	18.7587	0.0924	0.938	7 t3
86.4875	17.8697	0.0000	0.984	8 t4
-130.079	10.864	0.0000	0.971	9 rate
90.3335	7.22455	0.0000	0.974	10 rpm

N trials = 128



rate=125.00	rpm=260.00
Value	LowLimit HighLimit
27892.11	1.#R 1.#R



rate=125.00	rpm=260.00
Value	LowLimit HighLimit
31901.21	1.#R 1.#R

# NLHD, 16+16+32+64+36ckps=164 Blocks 1- 4, & 3 sets of checkpoints

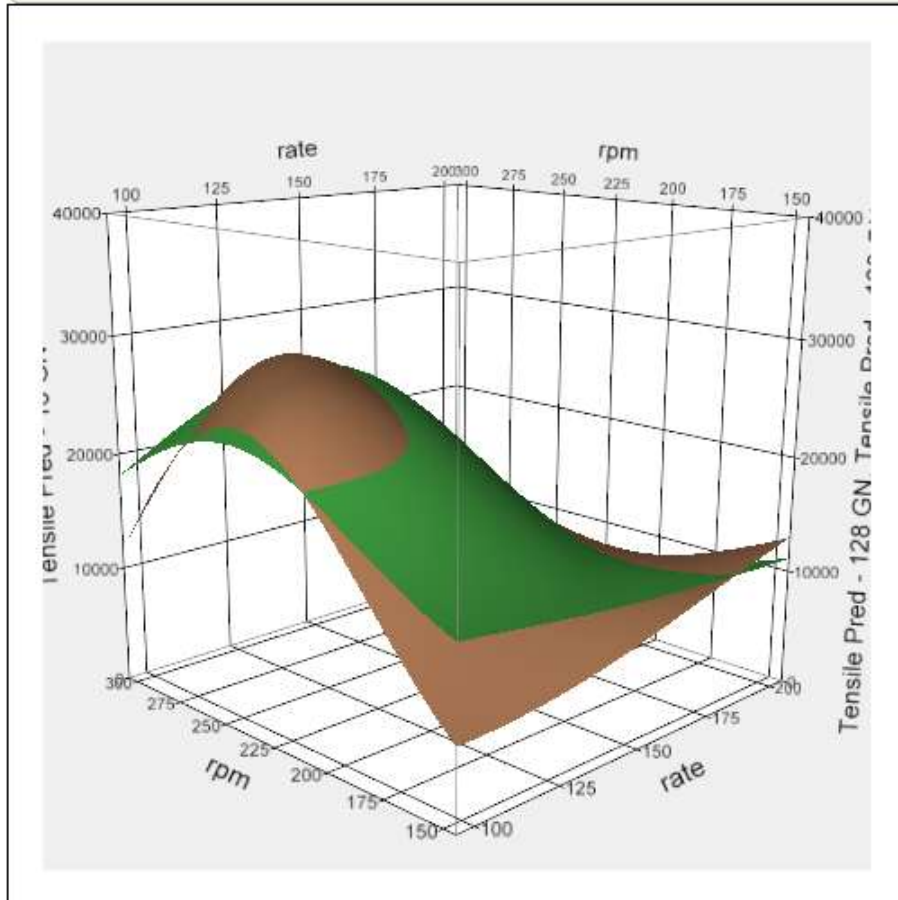
viscosity and t4 factor settings @ 75% of range

COEFFICIENTS	SD	P	CONDITION	TERM
17357				0 CONSTANT
-843.041	2246.05	0.7079-	0.973	1 add1
1016.04	553.662	0.0684	0.987	2 add2
226.532	55.3433	0.0001	0.987	3 viscosity
2098.09	7370.9	0.7763-	0.988	4 moisture
8.98137	18.4363	0.6268-	0.988	5 t1
-16.0393	18.4583	0.3862-	0.987	6 t2
5.38914	18.7981	0.7747-	0.969	7 t3
88.5134	18.3523	0.0000	0.992	8 t4
-106.215	11.086	0.0000	0.986	9 rate
76.3942	7.37976	0.0000	0.987	10 rpm

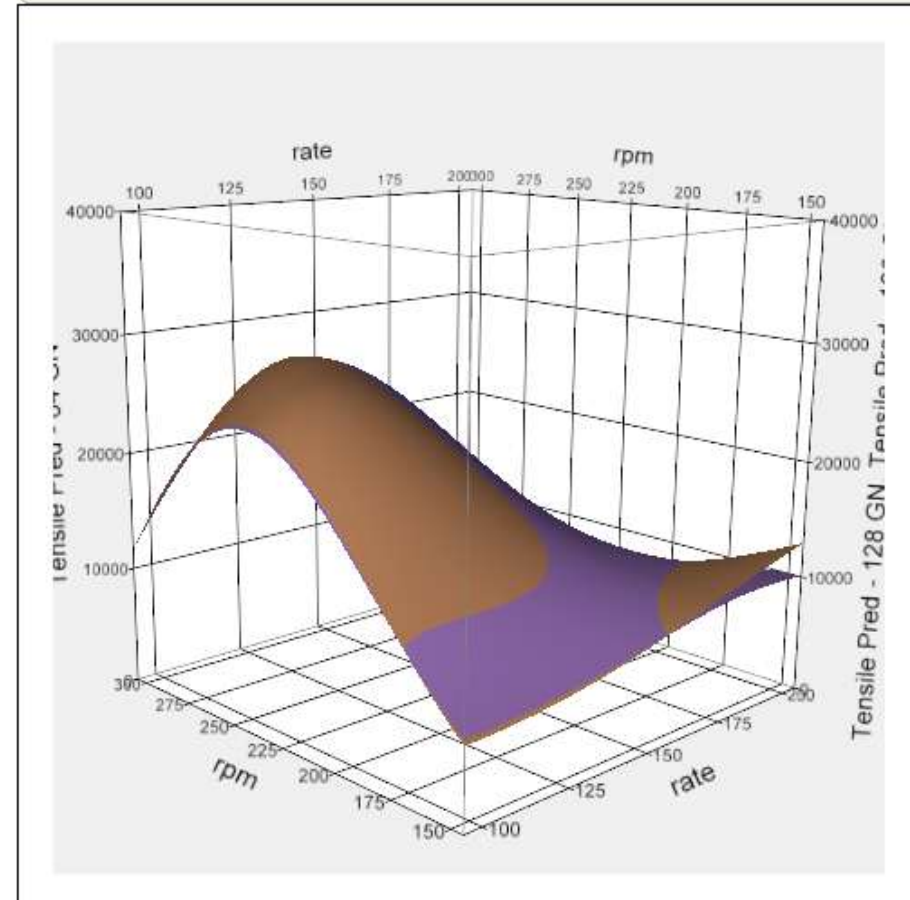
N trials = 164

Stage 1 fit of 16 trials colored green  
 Stage 4 fit 128 trials colored brown  
 Stage 3 fit 64 trials colored purple

Surface Plot

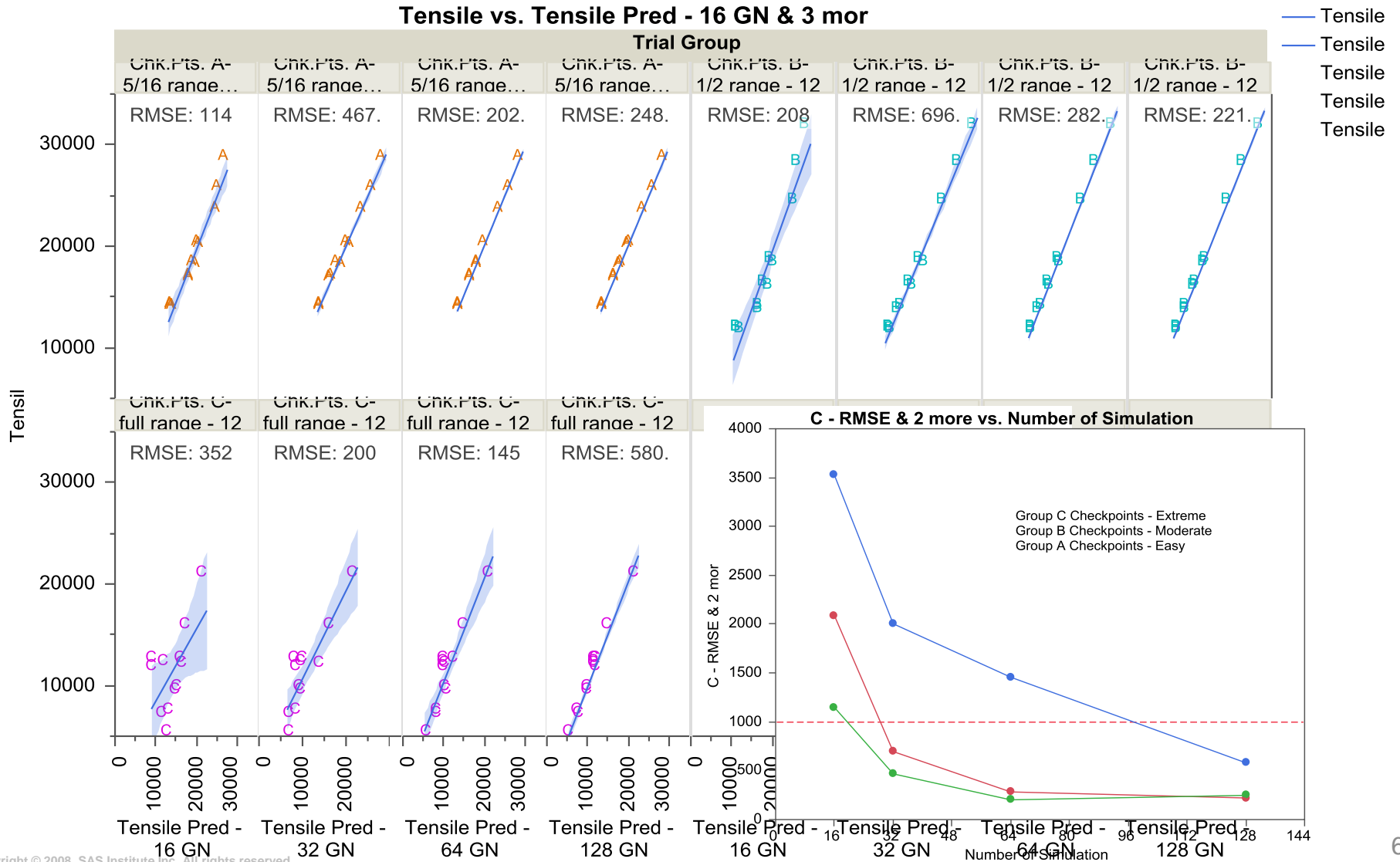


Surface Plot

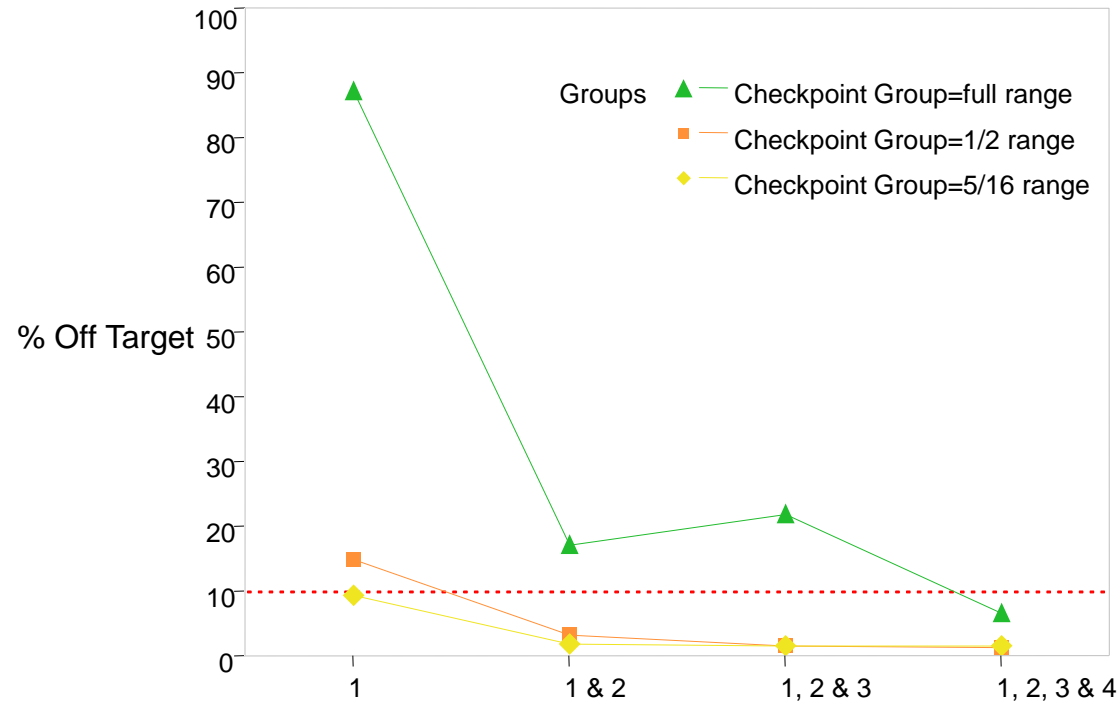


Checkpoint Groups A & B show diminishing return in prediction improvement for running past stage 3

## Tensile vs. Tensile Pred - 16 GN & 3 mor



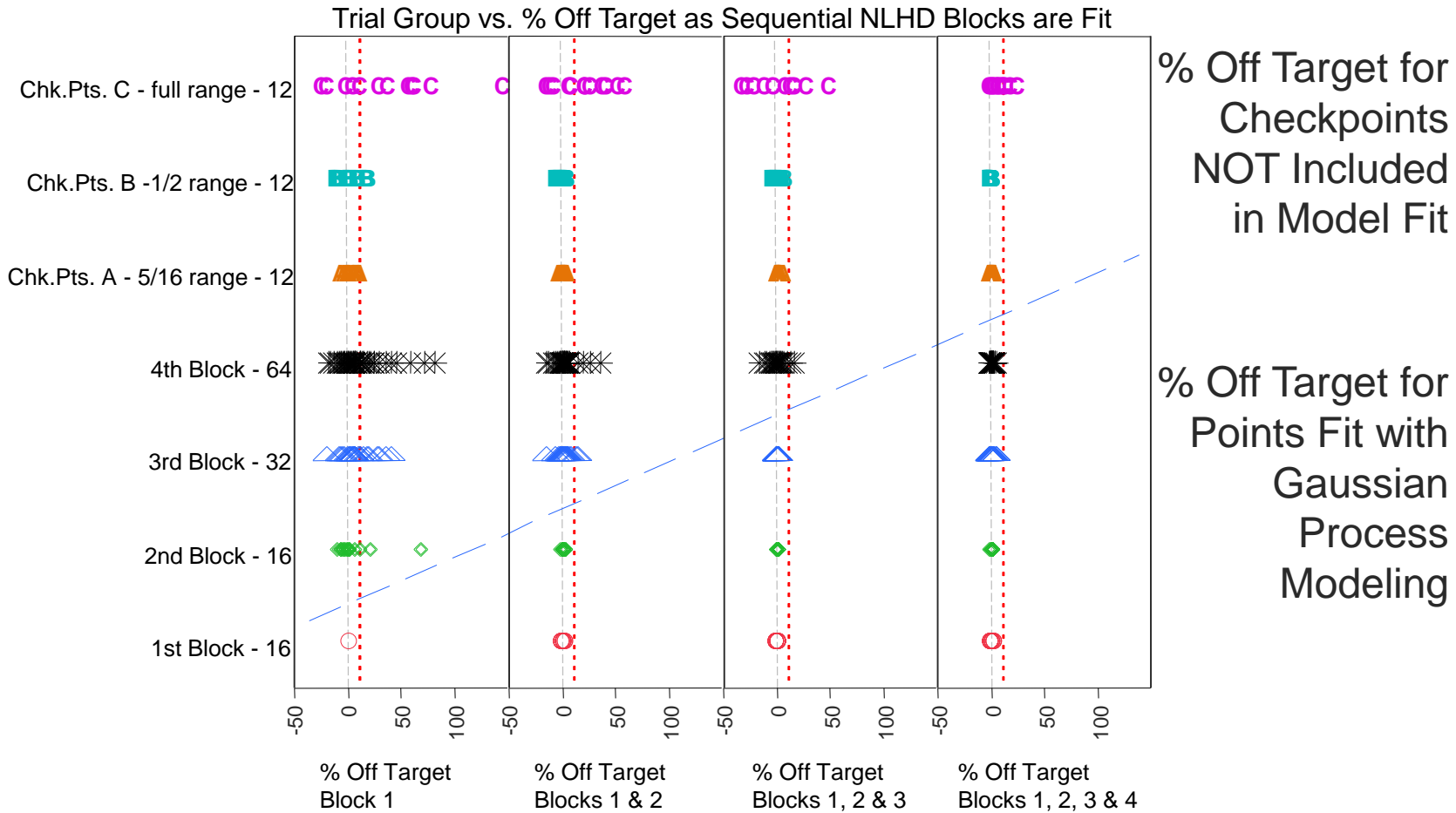
**Overlay Plot**



Percent Off Target - Root Mean Square of 12 Checkpoints				
Blocks	1	1 & 2	1, 2 & 3	1, 2, 3 & 4
5/16 range	9.39	2.08	1.72	1.53
1/2 range	14.94	3.33	1.79	1.27
full range	87.16	17.17	21.96	6.72

Percent Off Target - worst Case of 12 Checkpoints				
Blocks	1	1 & 2	1, 2 & 3	1, 2, 3 & 4
5/16 range	17.13	4.52	3.48	2.74
1/2 range	33.74	7.11	-3.38	2.31
full range	225.70	34.69	46.98	16.66

Each checkpoint group consisted of a 12-trial Plackett-Burman DOE. The ranges of the factors relative to the ranges used for the NLHD were 5/16ths (marginal extrapolation), half (moderate extrapolation) and full (extreme extrapolation).



- NLHD designs can be run sequentially so that surrogate model accuracy can be evaluated after each block and decision made as to whether or not to move forward with the next block
- Generally as more NLHD blocks are run, the surrogate model accuracy increases
- Inclusion of extreme (full range) extrapolation checkpoints will expand interpolation volume of Kriging analysis – assuming Kriging analysis remains stable
- Caveat: These conclusions were reached using a moderately complex transcendental function in lieu of a CFD simulation model that is believed to do a good job of stressing extrapolation with the surrogate model..



*Quicker answers, lower costs, solve bigger problems*

- Obtain a fast surrogate model of the simulation
  - Individual simulations can run for hours, days, weeks
    - Computational Fluid Dynamics (CFD)
    - Simulation runs in real-time
  - Numbers of factors can be very large (40+)
  - Numbers of simulations needed can be large (thousands in many cases)
  - Simulations can be stochastic requiring many replications
- Surrogate model yields a fast approximation of the simulation
  - more rapidly answer “what if?” questions
  - do sensitivity analysis of the control factors
  - optimize multiple responses and make trade-offs
- By running efficient subsets of all possible combinations, one can – for the same resources and constraints – *solve bigger problems*
- By running sequences of designs one can be as *cost effective as possible & run no more trials than are needed to get a useful answer*