




JMP Discovery Europe 2021

SVEM: A paradigm shift in Design and Analysis of Experiments



1

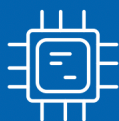
Agenda



- Introductions
- What, Why & How of SVEM
 - Quick overview of DOE and Machine learning
 - Describe blending them into SVEM
 - Analyze real-world SVEM experiments
 - Review Current Research
 - Demonstrate JMP's new Candidate Set Designer
- Next Steps & Q&A

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2



SVEM is a remarkable new method to extract more insights with fewer experimental cycles and build more accurate predictive models from small sets of data, including DOEs.

Less Cost / Faster to Market / Faster Problem-Solving

3

Speakers



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4

JMP's Contributions to Data Science

- DOE
 - Coordinate Exchange Algorithm
 - Definitive Screening Designs
 - SVEM



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5

SVEM

- Self-Validated, Ensemble Models
 - Combining two well-practiced methods
 - Machine Learning
 - Design of Experiments
- Overcoming limitations of limited amounts of data

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6

6

SVEM Field Trials: What We Have Learned

- SVEM works exceptionally well!
 - $n < p$
 - Higher order models
 - More accurate predictions
 - Recover from broken DOEs
 - DSDs that can't be run in Fit Definitive
 - Lacking power
 - Missing or questionable runs
- Does not help in Fractional Factorials

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7

7

SVEM Field Trials: What We Have Learned

- SVEM add-in full potential not tested
 - Need to design experiments with SVEM in mind
 - More factors
 - Bayesian I-optimal designs
 - Mixture designs

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8

8

jmp **SVEM: What, Why, and How** **sas**
THE POWER TO KNOW

Chris Gotwalt, JMP
Phil Ramsey, Univ. New Hampshire
Trent Lemkus, Univ. New Hampshire

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9

SVEM: What, Why, and How

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10

MACHINE LEARNING 101

	Compaction Pressure	Sintering Time	Sintering Temp C
1	75	30.80	1122
2	80	27.48	1111
3	100	27.48	1127
4	100	27.48	1118
5	85	20.85	1221
6	100	27.48	1118
7	95	27.48	1138
8	65	30.80	1122
9	75	30.80	1113
10	70	30.80	1127
11	90	20.85	1221
12	70	30.80	1221

$f(\text{pressure}, \text{time}, \text{temp})$

	Shrinkage
1	-0.37
2	-0.58
3	-0.16
4	0.05
5	-1.50
6	-0.09
7	0.55
8	-1.43
9	-0.93
10	-0.79
11	-1.22
12	-1.99

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11

MACHINE LEARNING 101

	Compaction Pressure	Sintering Time	Sintering Temp C
1	75	30.80	1122
2	80	27.48	1111
3	100	27.48	1127
4	100	27.48	1118
5	85	20.85	1221
6	100	27.48	1118
7	95	27.48	1138
8	65	30.80	1122

Fit models to the training set

	Shrinkage
1	-0.37
2	-0.58
3	-0.16
4	0.05
5	-1.50
6	-0.09
7	0.55
8	-1.43

Assess models using the validation set

9	75	30.80	1113
10	70	30.80	1127
11	90	20.85	1221
12	70	30.80	1221

9	-0.93
10	-0.79
11	-1.22
12	-1.99

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12

DOE MODELING 101

	Compaction Pressure	Sintering Time	Sintering Temp
1	60	15	1000
2	60	15	1000
3	60	15	1250
4	60	30	1000
5	60	30	1250
6	60	30	1250
7	115	15	1000
8	115	15	1250
9	115	15	1250
10	115	30	1000
11	115	30	1000
12	115	30	1250

$$f(\text{pressure, time, temp})$$



	Shrinkage
1	-1.85
2	-2.01
3	-3.75
4	-1.74
5	-3.8
6	-3.74
7	-1.95
8	-3.32
9	-2.79
10	-1.23
11	-1.31
12	-2.35

13

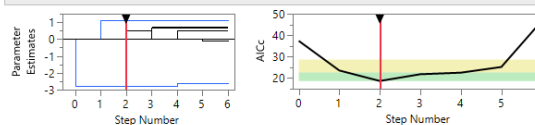
DOE MODELING 101

Effect Tests

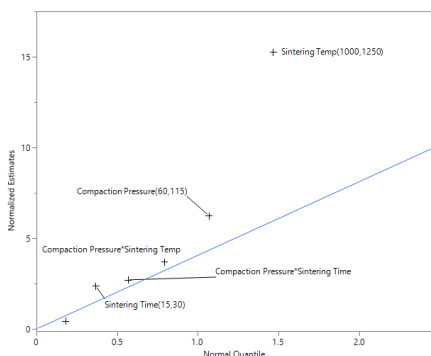
Source	Nparm	DF	Sum of Squares	F Ratio	Prob > F
Compaction Pressure(60,115)	1	1	1.1009611	33.0087	0.0022*
Sintering Time(15,30)	1	1	0.4005065	12.0079	0.0179*
Sintering Temp(1000,1250)	1	1	6.0741458	182.1132	<.0001*
Compaction Pressure*Sintering Time	1	1	0.2442070	7.3217	0.0425*
Compaction Pressure*Sintering Temp	1	1	0.4566092	13.6899	0.0140*
Sintering Time*Sintering Temp	1	1	0.0064027	0.1920	0.6796

Forward Selection via AICc

Solution Path



Half-Normal Plot



Effect Tests

Source	Nparm	DF	Sum of Squares	F Ratio	Prob > F
Sintering Temp(1000,1250)	1	1	7.777386	65.978133	<.0001*
Compaction Pressure(60,115)	1	1	1.3023508	11.048271	0.0089*
Sintering Time(15,30)	1	0	0	0	1.0000 Removed
Compaction Pressure*Sintering Time	1	0	0	0	1.0000 Removed
Compaction Pressure*Sintering Temp	1	0	0	0	1.0000 Removed
Sintering Time*Sintering Temp	1	0	0	0	1.0000 Removed

14

WHY NOT APPLY MACHINE LEARNING TO DOES?

	Compaction Pressure	Sintering Time	Sintering Temp
1	60	15	1000
2	60	15	1000
3	60	15	1250
4	60	30	1000
5	60	30	1250
6	60	30	1250
7	115	15	1000
8	115	15	1250


Training Set

	Shrinkage
1	-1.85
2	-2.01
3	-3.75
4	-1.74
5	-3.8
6	-3.74
7	-1.95
8	-3.32


9	115	15	1250
10	115	30	1000
11	115	30	1000
12	115	30	1250

Validation Set

9	-2.79
10	-1.23
11	-1.31
12	-2.35




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15

WHY NOT APPLY MACHINE LEARNING TO DOES?

Effect Tests						
Source	Nparm	DF	Sum of Squares	F Ratio	Prob > F	
Sintering Temp(1000,1250)	1	1	3.2802443	240.26823	0.0041*	
Compaction Pressure*Sintering Temp	1	1	0.0464416	3.4017103	0.2064	
Sintering Time*Sintering Temp	1	1	0.002799	0.205015	0.6951	
Compaction Pressure(60,115)	1	0	0	0	1.0000	LostDFs
Sintering Time(15,30)	1	0	0	0	1.0000	LostDFs
Compaction Pressure*Sintering Time	1	0	0	0	1.0000	LostDFs



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16

WHY NOT APPLY MACHINE LEARNING TO DOEs?

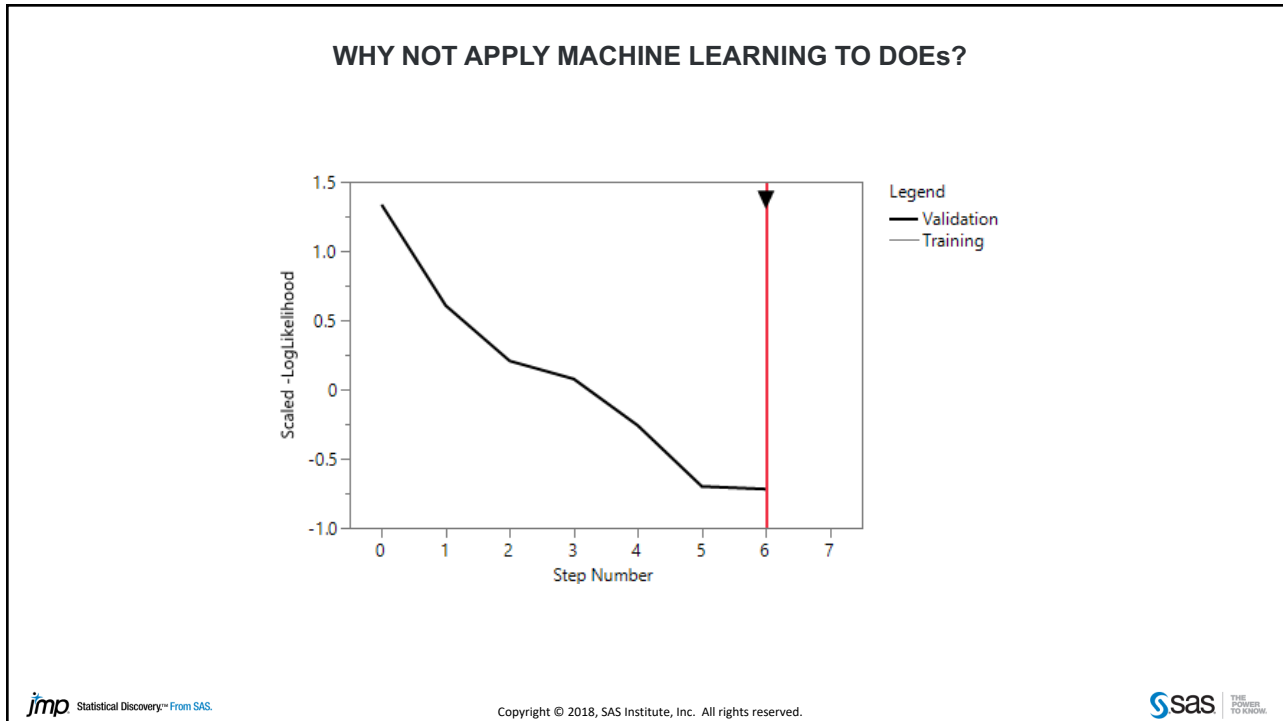
	Compaction Pressure	Sintering Time	Sintering Temp	Shrinkage
1	60	15	1000	-1.85
2	60	15	1000	-2.01
3	60	15	1250	-3.75
4	60	30	1000	-1.74
5	60	30	1250	-3.8
6	60	30	1250	-3.74
7	115	15	1000	-1.95
8	115	15	1250	-3.32
9	115	15	1250	-2.79
10	115	30	1000	-1.23
11	115	30	1000	-1.31
12	115	30	1250	-2.35

17

WHY NOT APPLY MACHINE LEARNING TO DOEs?

	Compaction Pressure	Sintering Time	Sintering Temp	Shrinkage	Validation
1	60	15	1000	-1.85	Training
2	60	15	1000	-1.85	Validation
3	60	15	1000	-2.01	Training
4	60	15	1000	-2.01	Validation
5	60	15	1250	-3.75	Training
6	60	15	1250	-3.75	Validation
7	60	30	1000	-1.74	Training
8	60	30	1000	-1.74	Validation
9	60	30	1250	-3.8	Training
10	60	30	1250	-3.8	Validation
11	60	30	1250	-3.74	Training
12	60	30	1250	-3.74	Validation
13	115	15	1000	-1.95	Training
14	115	15	1000	-1.95	Validation
15	115	15	1250	-3.32	Training
16	115	15	1250	-3.32	Validation
17	115	15	1250	-2.79	Training
18	115	15	1250	-2.79	Validation
19	115	30	1000	-1.23	Training
20	115	30	1000	-1.23	Validation
21	115	30	1000	-1.31	Training
22	115	30	1000	-1.31	Validation
23	115	30	1250	-2.35	Training
24	115	30	1250	-2.35	Validation

18

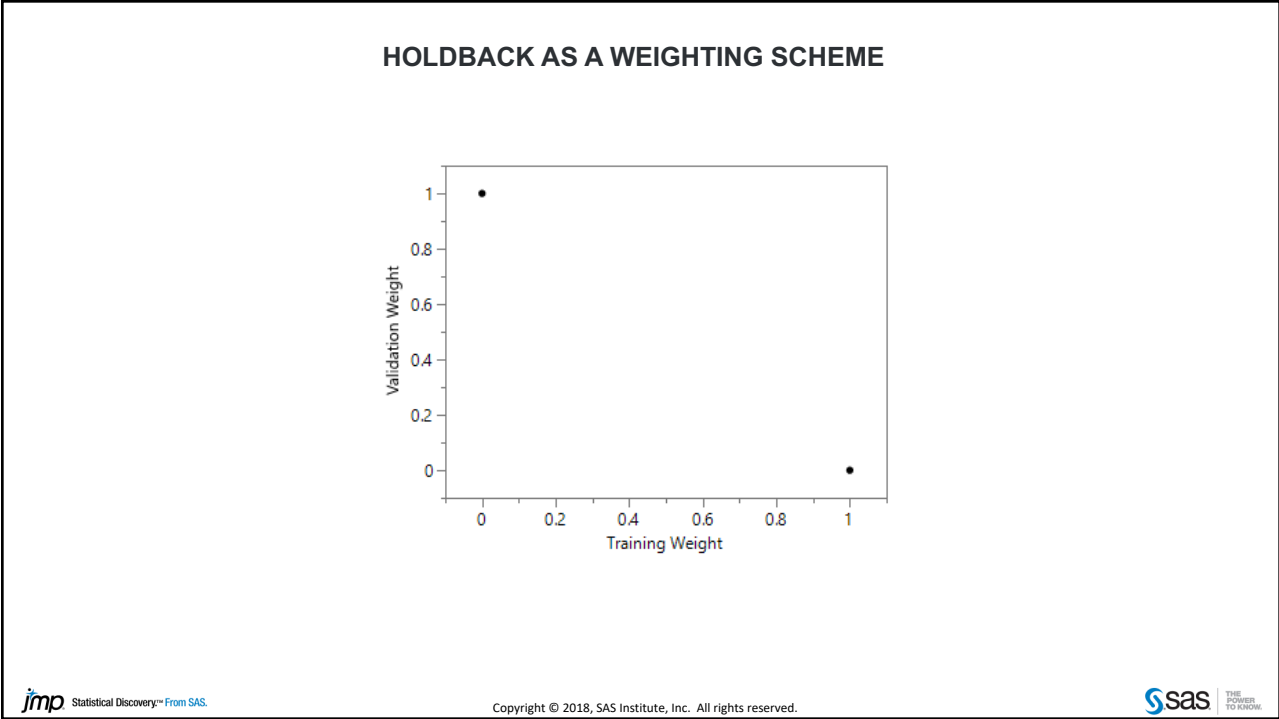


19

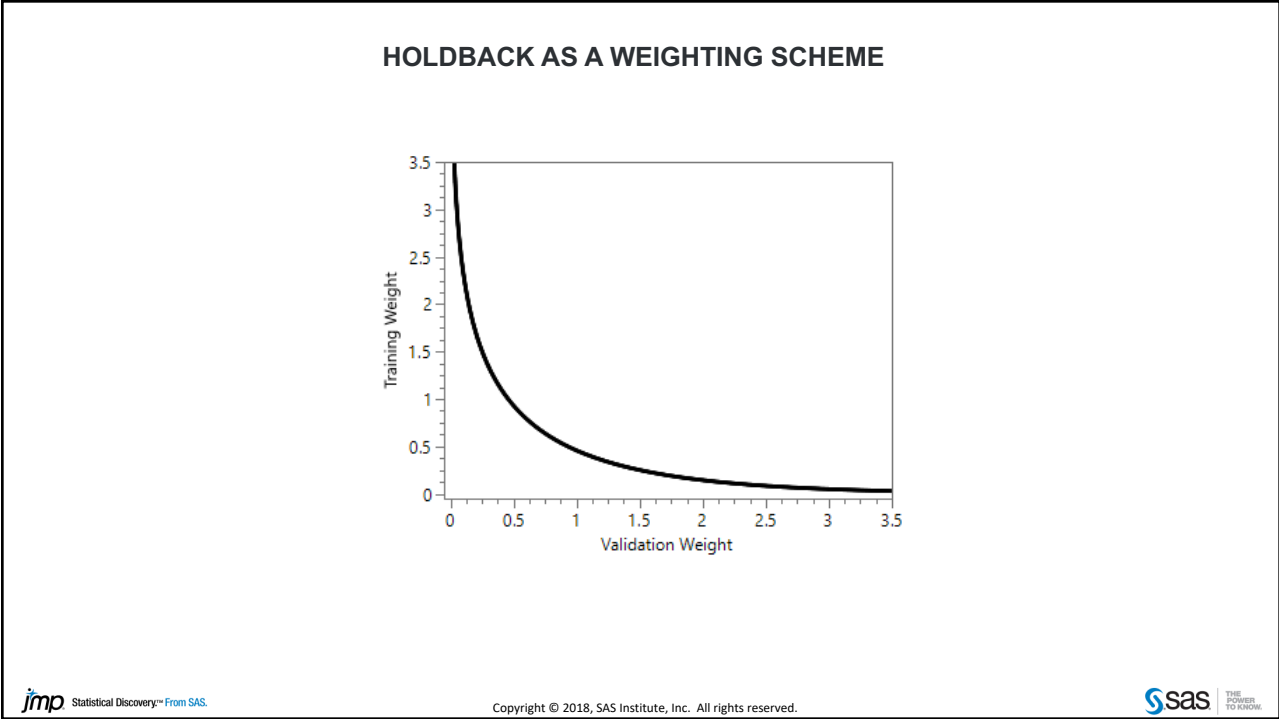
HOLDBACK AS A WEIGHTING SCHEME

	Compaction Pressure	Sintering Time	Sintering Temp	Shrinkage	Training Weight	Validation Weight
1	60	15	1000	-1.85	1	0
2	60	15	1000	-2.01	1	0
3	60	15	1250	-3.75	1	0
4	60	30	1000	-1.74	1	0
5	60	30	1250	-3.8	1	0
6	60	30	1250	-3.74	1	0
7	115	15	1000	-1.95	1	0
8	115	15	1250	-3.32	1	0
9	115	15	1250	-2.79	0	1
10	115	30	1000	-1.23	0	1
11	115	30	1000	-1.31	0	1
12	115	30	1250	-2.35	0	1

20



21



22

AUTOVALIDATION WEIGHTS

Create exponentially distributed weights by the *probability integral transform*:

$$u_i \sim \text{Uniform}(0,1)$$

$$w_i^T = \text{Exponential Quantile}(u_i, 1)$$

w_i^{Training} has an Exponential distribution with mean and variance 1.0.

$1 - u_i$ is also Uniform(0,1) and perfectly anticorrelated with u_i

$$w_i^V = \text{Exponential Quantile}(1 - u_i, 1)$$

w_i^V will be highly anticorrelated with w_i^T and yet has the exact same distribution!

23

AUTOVALIDATION WEIGHTS

	Compaction Pressure	Sintering Time	Sintering Temp	Shrinkage	U	Training Weight	Validation Weight
1	60	15	1000	-1.85	0.077	2.565	0.08
2	60	15	1000	-2.01	0.154	1.872	0.167
3	60	15	1250	-3.75	0.231	1.466	0.262
4	60	30	1000	-1.74	0.308	1.179	0.368
5	60	30	1250	-3.8	0.385	0.956	0.486
6	60	30	1250	-3.74	0.462	0.773	0.619
7	115	15	1000	-1.95	0.538	0.619	0.773
8	115	15	1250	-3.32	0.615	0.486	0.956
9	115	15	1250	-2.79	0.692	0.368	1.179
10	115	30	1000	-1.23	0.769	0.262	1.466
11	115	30	1000	-1.31	0.846	0.167	1.872
12	115	30	1250	-2.35	0.923	0.08	2.565

24

ROAD TO SVEM

	Compaction Pressure	Sintering Time	Sintering Temp	Shrinkage	U	Training Weight	Validation Weight											
1	60	15	1000	-1.85	0.804	0.218	1.63											
2	60	15	1000	-2.01	0.033	3.426	0.033											
3	60	15	1250	-3.75	0.624	0.472	0.978											
4	60	30	1000	-1.74	0.972	0.028	3.585											
5	60	30	1250	-3.8	0.992	0.008	4.85											
6	60	30	1250	-3.74	0.146	1.922	0.158											
7	115	15	1000	-1.95	0.046	3.082	0.047											
8	115	15	1250	-3.32	0.362	1.017	0.449											
9	115	15	1250	-2.79	0.154	1.871	0.167											
10	115	30	1000	-1.23	0.052	2.955	0.053											
11	115	30	1000	-1.31	0.019	3.962	0.019											
12	115	30	1250	-2.35	0.587	0.533	0.884											

25

ROAD TO SVEM

	Compaction Pressure	Sintering Time	Sintering Temp	Shrinkage	U	Training Weight	Validation Weight	Shrinkage Prediction Formula 1										
1	60	15	1000	-1.85	0.804	0.218	1.63	-2.01										
2	60	15	1000	-2.01	0.033	3.426	0.033	-2.01										
3	60	15	1250	-3.75	0.624	0.472	0.978	-3.62										
4	60	30	1000	-1.74	0.972	0.028	3.585	-2.01										
5	60	30	1250	-3.8	0.992	0.008	4.85	-3.62										
6	60	30	1250	-3.74	0.146	1.922	0.158	-3.62										
7	115	15	1000	-1.95	0.046	3.082	0.047	-1.351										
8	115	15	1250	-3.32	0.362	1.017	0.449	-2.962										
9	115	15	1250	-2.79	0.154	1.871	0.167	-2.962										
10	115	30	1000	-1.23	0.052	2.955	0.053	-1.351										
11	115	30	1000	-1.31	0.019	3.962	0.019	-1.351										
12	115	30	1250	-2.35	0.587	0.533	0.884	-2.962										

26

ROAD TO SVEM

$$-2.485909079 + 0.3294377473 \cdot \left(\frac{(\text{Compaction Pressure} - 87.5)}{27.5} \right) + -0.80505621 \cdot \left(\frac{(\text{Sintering Temp} - 1125)}{125} \right)$$

27

ROAD TO SVEM

	Compaction Pressure	Sintering Time	Sintering Temp	Shrinkage	U	Training Weight	Validation Weight	Shrinkage Prediction Formula 1										
1	60	15	1000	-1.85	0.804	0.218	1.63	-2.01										
2	60	15	1000	-2.01	0.033	3.426	0.033	-2.01										
3	60	15	1250	-3.75	0.624	0.472	0.978	-3.62										
4	60	30	1000	-1.74	0.972	0.028	3.585	-2.01										
5	60	30	1250	-3.8	0.992	0.008	4.85	-3.62										
6	60	30	1250	-3.74	0.146	1.922	0.158	-3.62										
7	115	15	1000	-1.95	0.046	3.082	0.047	-1.351										
8	115	15	1250	-3.32	0.362	1.017	0.449	-2.962										
9	115	15	1250	-2.79	0.154	1.871	0.167	-2.962										
10	115	30	1000	-1.23	0.052	2.955	0.053	-1.351										
11	115	30	1000	-1.31	0.019	3.962	0.019	-1.351										
12	115	30	1250	-2.35	0.587	0.533	0.884	-2.962										

28

ROAD TO SVEM

	Compaction Pressure	Sintering Time	Sintering Temp	Shrinkage	U	Training Weight	Validation Weight	Shrinkage Prediction Formula 1	Shrinkage Prediction Formula 2
1	60	15	1000	-1.85	0.529	0.637	0.753	-2.01	-1.902
2	60	15	1000	-2.01	0.055	2.9	0.057	-2.01	-1.902
3	60	15	1250	-3.75	0.637	0.451	1.014	-3.62	-3.82
4	60	30	1000	-1.74	0.714	0.337	1.251	-2.01	-1.826
5	60	30	1250	-3.8	0.93	0.073	2.654	-3.62	-3.745
6	60	30	1250	-3.74	0.721	0.326	1.278	-3.62	-3.745
7	115	15	1000	-1.95	0.466	0.763	0.628	-1.351	-1.951
8	115	15	1250	-3.32	0.254	1.372	0.293	-2.962	-3.082
9	115	15	1250	-2.79	0.644	0.44	1.033	-2.962	-3.082
10	115	30	1000	-1.23	0.694	0.366	1.183	-1.351	-1.229
11	115	30	1000	-1.31	0.627	0.467	0.985	-1.351	-1.229
12	115	30	1250	-2.35	0.231	1.464	0.263	-2.962	-2.36

ROAD TO SVEM

$$\begin{aligned}
 & -2.489456343 \\
 & + 0.3340338147 \cdot \left(\frac{\left(\left(\text{Compaction Pressure} - 87.5 \right) \right)}{27.5} \right) \\
 & + 0.1993742337 \cdot \left(\frac{\left(\left(\text{Sintering Time} - 22.5 \right) \right)}{7.5} \right) \\
 & + -0.762447346 \cdot \left(\frac{\left(\left(\text{Sintering Temp} - 1125 \right) \right)}{125} \right) \\
 & + \left(\frac{\left(\left(\text{Compaction Pressure} - 87.5 \right) \right)}{27.5} \right) \cdot \left(\frac{\left(\left(\text{Sintering Time} - 22.5 \right) \right)}{7.5} \right) \cdot 0.1615679978 \\
 & + \left(\frac{\left(\left(\text{Compaction Pressure} - 87.5 \right) \right)}{27.5} \right) \cdot \left(\frac{\left(\left(\text{Sintering Temp} - 1125 \right) \right)}{125} \right) \cdot 0.1967542111
 \end{aligned}$$

ROAD TO SVEM

	Compaction Pressure	Sintering Time	Sintering Temp	Shrinkage	U	Training Weight	Validation Weight	Shrinkage Prediction Formula 1	Shrinkage Prediction Formula 2									
1	60	15	1000	-1.85	0.529	0.637	0.753	-2.01	-1.902									
2	60	15	1000	-2.01	0.055	2.9	0.057	-2.01	-1.902									
3	60	15	1250	-3.75	0.637	0.451	1.014	-3.62	-3.82									
4	60	30	1000	-1.74	0.714	0.337	1.251	-2.01	-1.826									
5	60	30	1250	-3.8	0.93	0.073	2.654	-3.62	-3.745									
6	60	30	1250	-3.74	0.721	0.326	1.278	-3.62	-3.745									
7	115	15	1000	-1.95	0.466	0.763	0.628	-1.351	-1.951									
8	115	15	1250	-3.32	0.254	1.372	0.293	-2.962	-3.082									
9	115	15	1250	-2.79	0.644	0.44	1.033	-2.962	-3.082									
10	115	30	1000	-1.23	0.694	0.366	1.183	-1.351	-1.229									
11	115	30	1000	-1.31	0.627	0.467	0.985	-1.351	-1.229									
12	115	30	1250	-2.35	0.231	1.464	0.263	-2.962	-2.36									

31

ROAD TO SVEM

	Compaction Pressure	Sintering Time	Sintering Temp	Shrinkage	U	Training Weight	Validation Weight	Shrinkage Prediction Formula 1	Shrinkage Prediction Formula 2	Shrinkage Prediction Formula 3								
1	60	15	1000	-1.85	0.033	3.418	0.033	-2.01	-1.902	-1.885								
2	60	15	1000	-2.01	0.146	1.925	0.158	-2.01	-1.902	-1.885								
3	60	15	1250	-3.75	0.367	1.001	0.458	-3.62	-3.82	-3.885								
4	60	30	1000	-1.74	0.093	2.379	0.097	-2.01	-1.826	-1.746								
5	60	30	1250	-3.8	0.027	3.606	0.028	-3.62	-3.745	-3.747								
6	60	30	1250	-3.74	0.425	0.857	0.553	-3.62	-3.745	-3.747								
7	115	15	1000	-1.95	0.761	0.273	1.431	-1.351	-1.951	-1.935								
8	115	15	1250	-3.32	0.189	1.667	0.209	-2.962	-3.082	-3.031								
9	115	15	1250	-2.79	0.416	0.876	0.539	-2.962	-3.082	-3.031								
10	115	30	1000	-1.23	0.687	0.375	1.162	-1.351	-1.229	-1.255								
11	115	30	1000	-1.31	0.56	0.581	0.82	-1.351	-1.229	-1.255								
12	115	30	1250	-2.35	0.982	0.018	4.026	-2.962	-2.36	-2.35								

32

ROAD TO SVEM

	Compaction Pressure	Sintering Time	Sintering Temp	Shrinkage	U	Training Weight	Validation Weight	Shrinkage Prediction Formula 1	Shrinkage Prediction Formula 2	Shrinkage Prediction Formula 3	Shrinkage Prediction Formula 4
1	60	15	1000	-1.85	0.746	0.293	1.372	-2.01	-1.902	-1.885	-1.921
2	60	15	1000	-2.01	0.315	1.157	0.378	-2.01	-1.902	-1.885	-1.921
3	60	15	1250	-3.75	0.163	1.814	0.178	-3.62	-3.82	-3.885	-3.797
4	60	30	1000	-1.74	0.474	0.747	0.642	-2.01	-1.826	-1.746	-1.856
5	60	30	1250	-3.8	0.286	1.253	0.336	-3.62	-3.745	-3.747	-3.731
6	60	30	1250	-3.74	0.896	0.11	2.265	-3.62	-3.745	-3.747	-3.731
7	115	15	1000	-1.95	0.184	1.69	0.204	-1.351	-1.951	-1.935	-2.025
8	115	15	1250	-3.32	0.075	2.596	0.078	-2.962	-3.082	-3.031	-3.236
9	115	15	1250	-2.79	0.838	0.177	1.819	-2.962	-3.082	-3.031	-3.236
10	115	30	1000	-1.23	0.452	0.793	0.602	-1.351	-1.229	-1.255	-1.237
11	115	30	1000	-1.31	0.133	2.014	0.143	-1.351	-1.229	-1.255	-1.237
12	115	30	1250	-2.35	0.252	1.38	0.29	-2.962	-2.36	-2.35	-2.448

33

ROAD TO SVEM

	Compaction Pressure	Sintering Time	Sintering Temp	Shrinkage	U	Training Weight	Validation Weight	Shrinkage Prediction Formula 1	Shrinkage Prediction Formula 2	Shrinkage Prediction Formula 3	Shrinkage Prediction Formula 4	Shrinkage Prediction Formula 5
1	60	15	1000	-1.85	0.096	2.345	0.101	-2.01	-1.902	-1.885	-1.921	-1.873
2	60	15	1000	-2.01	0.671	0.399	1.112	-2.01	-1.902	-1.885	-1.921	-1.873
3	60	15	1250	-3.75	0.161	1.827	0.175	-3.62	-3.82	-3.885	-3.797	-3.749
4	60	30	1000	-1.74	0.987	0.013	4.322	-2.01	-1.826	-1.746	-1.856	-1.707
5	60	30	1250	-3.8	0.592	0.524	0.898	-3.62	-3.745	-3.747	-3.731	-3.778
6	60	30	1250	-3.74	0.736	0.307	1.331	-3.62	-3.745	-3.747	-3.731	-3.778
7	115	15	1000	-1.95	0.797	0.227	1.594	-1.351	-1.951	-1.935	-2.025	-1.947
8	115	15	1250	-3.32	0.932	0.07	2.693	-2.962	-3.082	-3.031	-3.236	-2.82
9	115	15	1250	-2.79	0.342	1.073	0.418	-2.962	-3.082	-3.031	-3.236	-2.82
10	115	30	1000	-1.23	0.694	0.366	1.183	-1.351	-1.229	-1.255	-1.237	-1.287
11	115	30	1000	-1.31	0.3	1.203	0.357	-1.351	-1.229	-1.255	-1.237	-1.287
12	115	30	1250	-2.35	0.395	0.928	0.503	-2.962	-2.36	-2.35	-2.448	-2.354

34

ROAD TO SVEM

$$\text{Mean} \begin{pmatrix} \text{Shrinkage Prediction Formula 1} \\ \text{Shrinkage Prediction Formula 2} \\ \text{Shrinkage Prediction Formula 3} \\ \text{Shrinkage Prediction Formula 4} \\ \text{Shrinkage Prediction Formula 5} \end{pmatrix}$$

35

ROAD TO SVEM

	Compaction Pressure	Sintering Time	Sintering Temp	Shrinkage	U	Training Weight	Validation Weight	Shrinkage Prediction Formula 1	Shrinkage Prediction Formula 2	Shrinkage Prediction Formula 3	Shrinkage Prediction Formula 4	Shrinkage Prediction Formula 5						
1	60	15	1000	-1.85	0.096	2.345	0.101	-2.01	-1.902	-1.885	-1.921	-1.873						
2	60	15	1000	-2.01	0.671	0.399	1.112	-2.01	-1.902	-1.885	-1.921	-1.873						
3	60	15	1250	-3.75	0.161	1.827	0.175	-3.62	-3.82	-3.885	-3.797	-3.749						
4	60	30	1000	-1.74	0.987	0.013	4.322	-2.01	-1.826	-1.746	-1.856	-1.707						
5	60	30	1250	-3.8	0.592	0.524	0.898	-3.62	-3.745	-3.747	-3.731	-3.778						
6	60	30	1250	-3.74	0.736	0.307	1.331	-3.62	-3.745	-3.747	-3.731	-3.778						
7	115	15	1000	-1.95	0.797	0.227	1.594	-1.351	-1.951	-1.935	-2.025	-1.947						
8	115	15	1250	-3.32	0.932	0.07	2.693	-2.962	-3.082	-3.031	-3.236	-2.82						
9	115	15	1250	-2.79	0.342	1.073	0.418	-2.962	-3.082	-3.031	-3.236	-2.82						
10	115	30	1000	-1.23	0.694	0.366	1.183	-1.351	-1.229	-1.255	-1.237	-1.287						
11	115	30	1000	-1.31	0.3	1.203	0.357	-1.351	-1.229	-1.255	-1.237	-1.287						
12	115	30	1250	-2.35	0.395	0.928	0.503	-2.962	-2.36	-2.35	-2.448	-2.354						

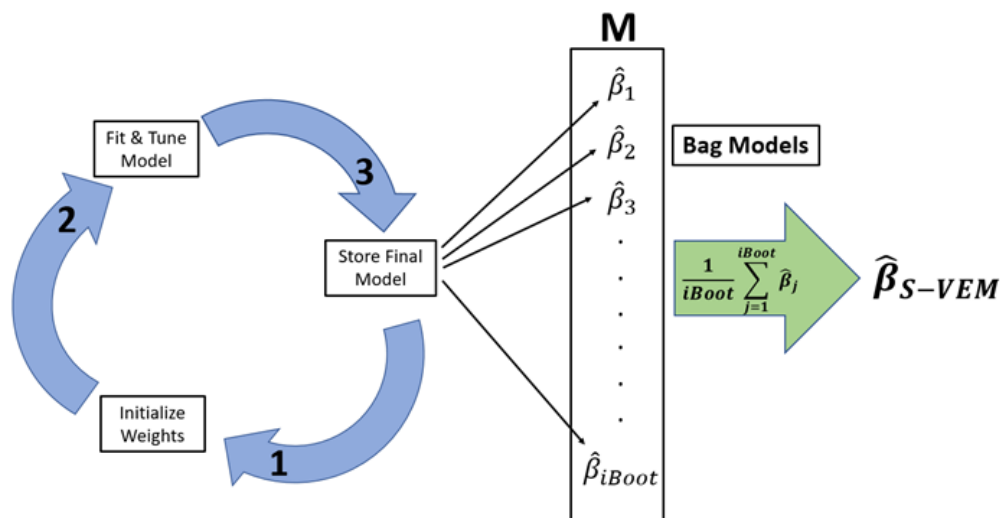
36

ROAD TO SVEM

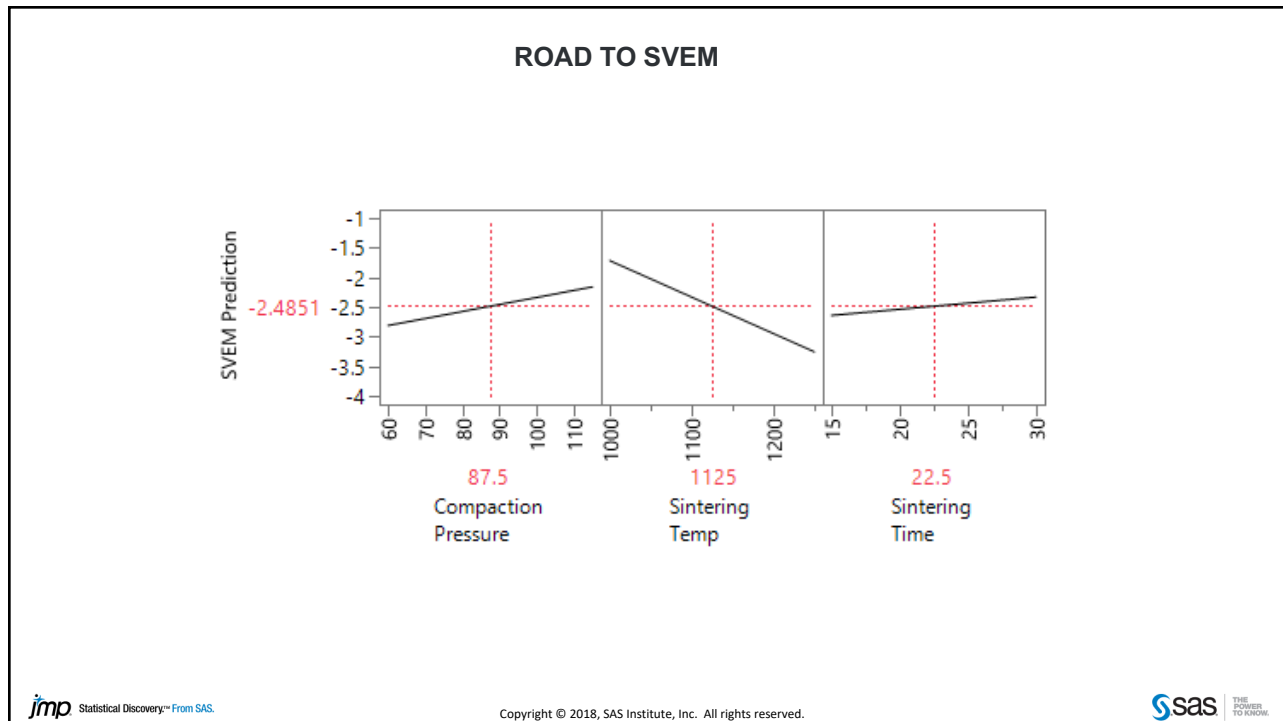
	Compaction Pressure	Sintering Time	Sintering Temp	Shrinkage	U	Training Weight	Validation Weight	Shrinkage Prediction Formula 1	Shrinkage Prediction Formula 2	Shrinkage Prediction Formula 3	Shrinkage Prediction Formula 4	Shrinkage Prediction Formula 5	SVEM Prediction
1	60	15	1000	-1.85	0.096	2.345	0.101	-2.01	-1.902	-1.885	-1.921	-1.873	-1.9183
2	60	15	1000	-2.01	0.671	0.399	1.112	-2.01	-1.902	-1.885	-1.921	-1.873	-1.9183
3	60	15	1250	-3.75	0.161	1.827	0.175	-3.62	-3.82	-3.885	-3.797	-3.749	-3.7744
4	60	30	1000	-1.74	0.987	0.013	4.322	-2.01	-1.826	-1.746	-1.856	-1.707	-1.8293
5	60	30	1250	-3.8	0.592	0.524	0.898	-3.62	-3.745	-3.747	-3.731	-3.778	-3.7242
6	60	30	1250	-3.74	0.736	0.307	1.331	-3.62	-3.745	-3.747	-3.731	-3.778	-3.7242
7	115	15	1000	-1.95	0.797	0.227	1.594	-1.351	-1.951	-1.935	-2.025	-1.947	-1.842
8	115	15	1250	-3.32	0.932	0.07	2.693	-2.962	-3.082	-3.031	-3.236	-2.82	-3.026
9	115	15	1250	-2.79	0.342	1.073	0.418	-2.962	-3.082	-3.031	-3.236	-2.82	-3.026
10	115	30	1000	-1.23	0.694	0.366	1.183	-1.351	-1.229	-1.255	-1.237	-1.287	-1.272
11	115	30	1000	-1.31	0.3	1.203	0.357	-1.351	-1.229	-1.255	-1.237	-1.287	-1.272
12	115	30	1250	-2.35	0.395	0.928	0.503	-2.962	-2.36	-2.35	-2.448	-2.354	-2.4948

37

SVEM



38



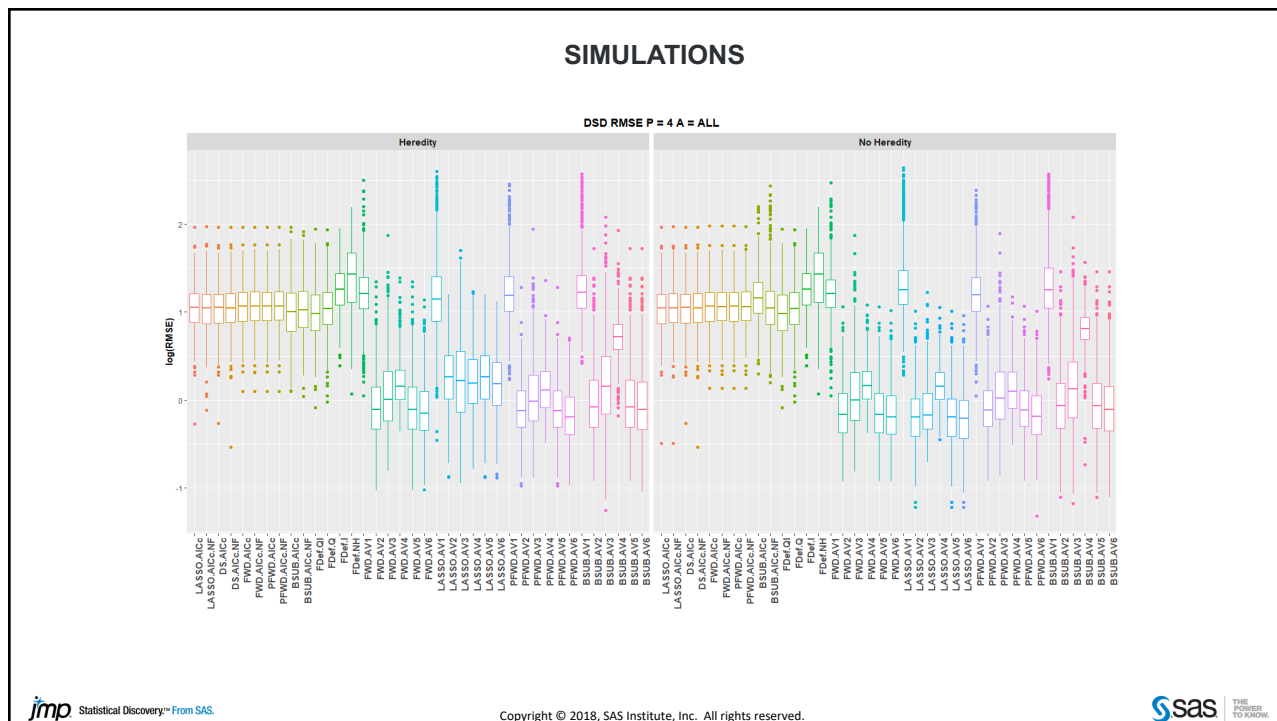
39

SIMULATIONS

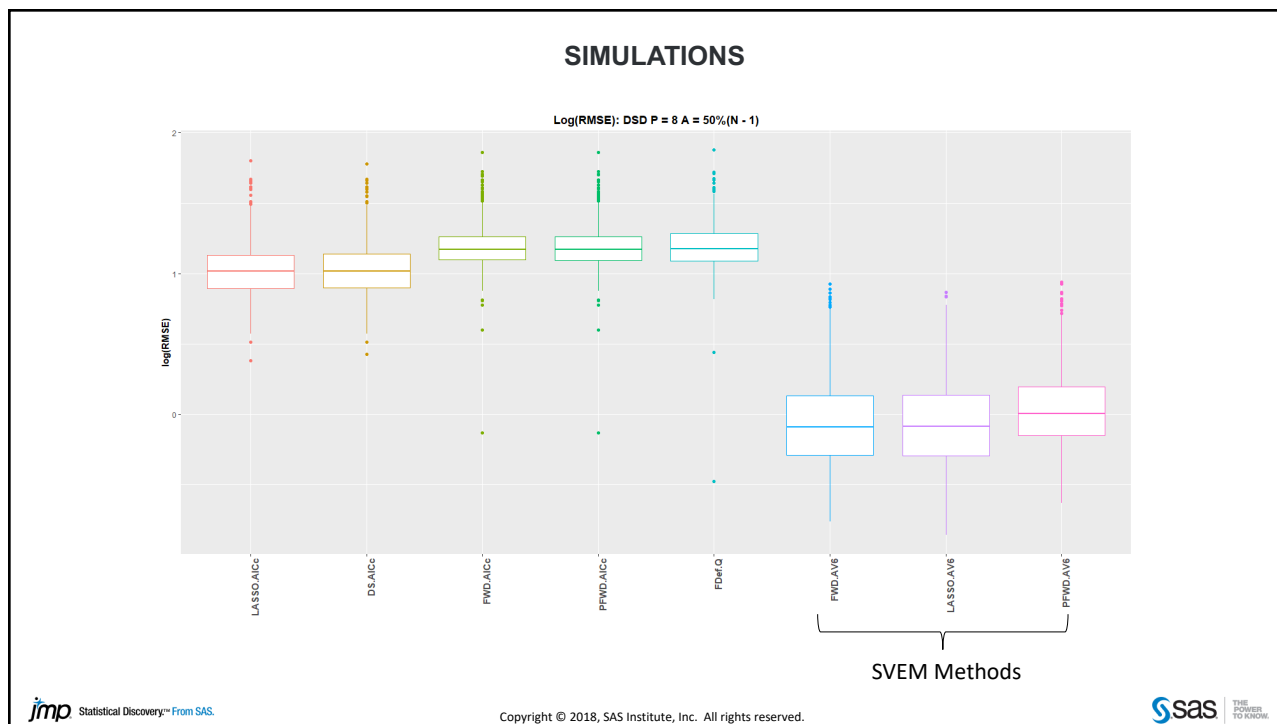
- Definitive Screening and Box Behnken Designs in 4 and 8 DoE factors
- Tried many “classical” and autovalidation based modeling approaches, all based on quadratic RSM.
- 1000 simulation reps per situation
- Each sim rep had its own set of “true” model coefficients
 - Nonzero model coefficients were double exponentially distributed
 - “True” nonzero coefficients represented 50%-100% of all possible coefficients
- Models evaluated on an independent set RMSE (n=10k, spacefilling design) vs. true model

jmp Statistical Discovery.™ From SAS. Copyright © 2018, SAS Institute, Inc. All rights reserved. sas THE POWER TO KNOW.

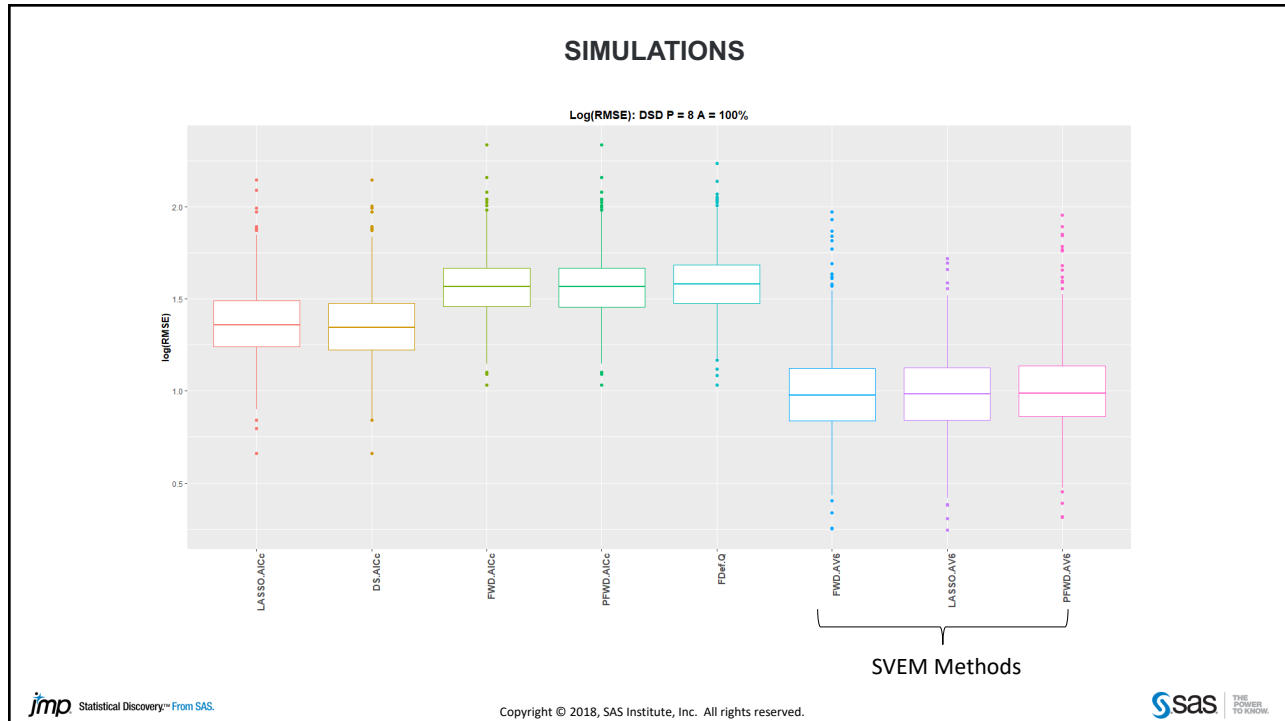
40



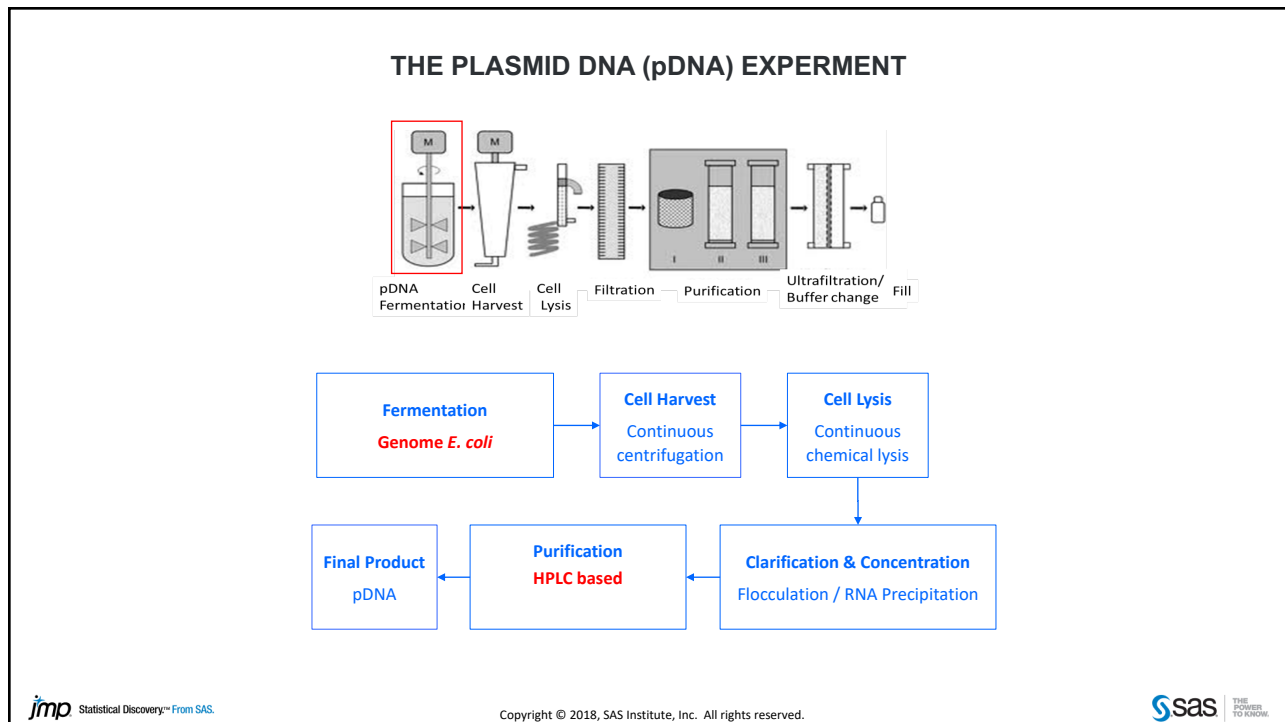
41



42



43



44

THE PLASMID DNA (pDNA) EXPERIMENT

	pH	%DO	Induction TempC	Induction OD600	Feed Rate	pDNA Titer mg/L
1	6.8	30	42.5	40	1.9	285.60
2	7.0	30	41.0	30	2.7	364.00
3	7.0	30	41.0	30	2.7	348.08
4	7.0	30	41.0	30	2.7	434.74
5	7.0	30	41.0	30	2.7	339.74
6	7.2	40	42.5	20	1.9	154.46
7	7.0	30	41.0	30	2.7	430.35
8	6.8	20	42.5	40	3.5	341.00
9	7.2	20	42.5	20	3.5	303.82
10	7.2	30	39.5	20	3.5	398.00
11	7.0	30	41.0	30	2.7	411.74
12	6.8	20	39.5	20	3.5	517.23
13	6.7	30	41.0	30	2.7	338.68
14	7.2	20	41.0	40	1.9	229.00
15	7.0	30	41.0	17	2.7	282.29
16	6.8	40	41.0	20	3.5	377.00

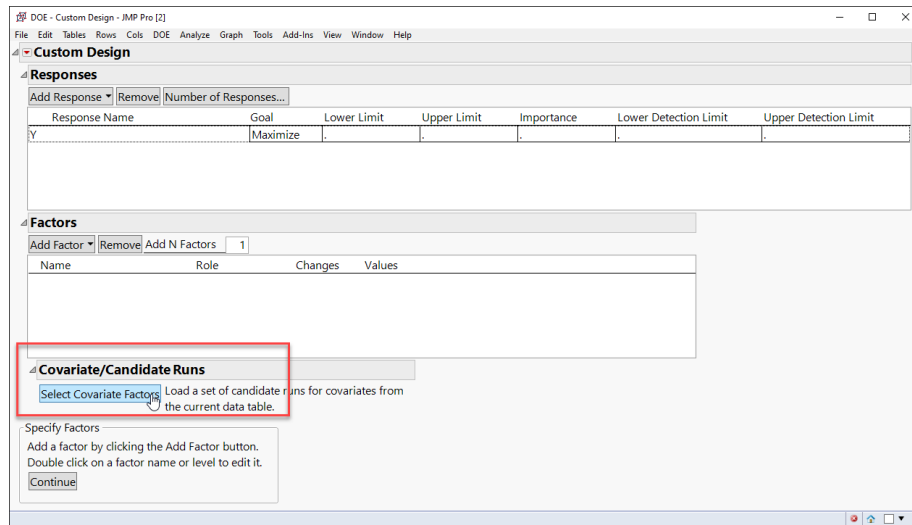
45

THE PLASMID DNA (pDNA) EXPERIMENT

The screenshot shows the JMP Pro software interface for a Design of Experiments (DOE) project. The main window displays a table of experimental runs with columns for pH, %DO, Induction TempC, Induction OD600, Feed Rate, and pDNA Titer mg/L. A 'Custom Design' menu is open, showing options like Augment Design, Definitive Screening, Classical, Design Diagnostics, Consumer Studies, Special Purpose, and Sample Size Explorers. The 'Columns (7/0)' list on the left includes pH *, %DO *, Induction TempC *, Induction OD600 *, Feed Rate *, and pDNA Titer mg/L *. The 'Rows' section shows 46 total rows, with 0 selected, 0 excluded, and 0 hidden. The status bar at the bottom indicates 'evaluations done'.

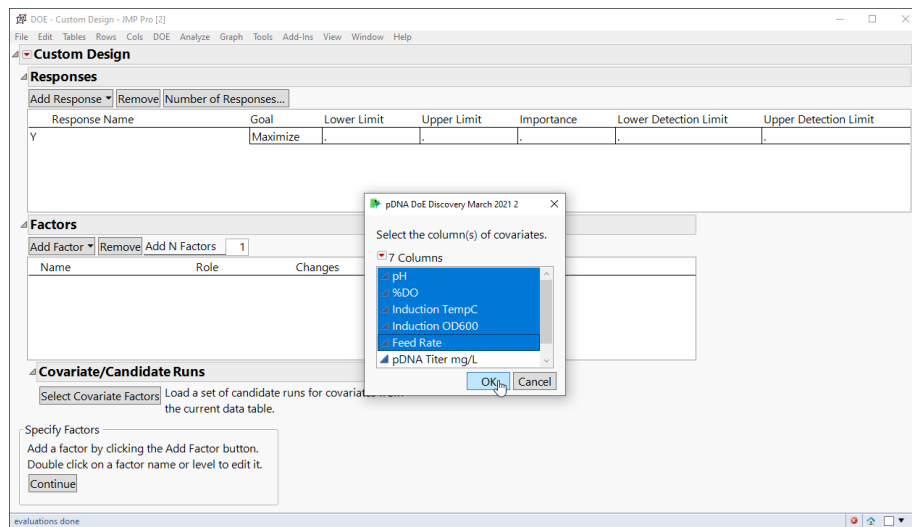
46

THE PLASMID DNA (pDNA) EXPERIMENT



47

THE PLASMID DNA (pDNA) EXPERIMENT



48

THE PLASMID DNA (pDNA) EXPERIMENT

Custom Design

Responses

Response Name	Goal	Lower Limit	Upper Limit	Importance	Lower Detection Limit	Upper Detection Limit
Y	Maximize

Factors

Name	Role	Changes	Values
pH	Covariate	Easy	6.74 7.26
%DO	Covariate	Easy	17 43
Induction TempC	Covariate	Easy	39.05 42.95
Induction OD600	Covariate	Easy	17 43
Feed Rate	Covariate	Easy	1.66 3.74

Covariate/Candidate Runs

Specify Factors
 Add a factor by clicking the Add Factor button.
 Double click on a factor name or level to edit it.

[Continue](#)

49

THE PLASMID DNA (pDNA) EXPERIMENT

Custom Design

Responses

Factors

Name	Role	Changes	Values
pH	Covariate	Easy	6.74 7.26
%DO	Covariate	Easy	17 43
Induction TempC	Covariate	Easy	39.05 42.95
Induction OD600	Covariate	Easy	17 43
Feed Rate	Covariate	Easy	1.66 3.74

Covariate/Candidate Runs

Define Factor Constraints

None
 Specify Linear Constraints
 Use Disallowed Combinations Filter
 Use Disallowed Combinations Script

Model

Main Effects Interactions **RSM** Cross Powers Remove Term

Name	Estimability
Intercept	Necessary
pH	Necessary
%DO	Necessary
Induction TempC	Necessary
Induction OD600	Necessary
Feed Rate	Necessary

Alias Terms

Design Generation

Include all selected covariate rows in the design
 Allow covariate rows to be repeated
 Number of Runs: 46

[Make Design](#)

50

THE PLASMID DNA (pDNA) EXPERIMENT

Custom Design

Responses

Factors

Name	Role	Changes	Values
pH	Covariate	Easy	6.74, 7.26
%DO	Covariate	Easy	17, 43
Induction TempC	Covariate	Easy	39.05, 42.95
Induction OD600	Covariate	Easy	17, 43
Feed Rate	Covariate	Easy	1.66, 3.74

Covariate/Candidate Runs

Define Factor Constraints

Model

Name	Estimability
Intercept	Necessary
pH	Necessary
%DO	Necessary
Induction TempC	Necessary
Induction OD600	Necessary
Feed Rate	Necessary
pH*pH	Necessary
pH*%DO	If Possible

Design Generation

Number of Runs: 46

51

THE PLASMID DNA (pDNA) EXPERIMENT

Custom Design

Responses

Factors

Name	Role	Changes	Values
pH	Covariate	Easy	6.74, 7.26
%DO	Covariate	Easy	17, 43
Induction TempC	Covariate	Easy	39.05, 42.95
Induction OD600	Covariate	Easy	17, 43
Feed Rate	Covariate	Easy	1.66, 3.74

Covariate/Candidate Runs

Define Factor Constraints

Model

Name	Estimability
Intercept	Necessary
pH	If Possible
%DO	If Possible
Induction TempC	If Possible
Induction OD600	If Possible
Feed Rate	If Possible
pH*pH	If Possible
pH*%DO	If Possible

Design Generation

Number of Runs: 16

52

THE PLASMID DNA (pDNA) EXPERIMENT

The screenshot shows the 'Custom Design' window in SAS JMP. The 'Responses' section is empty. Under 'Factors', a table lists five covariate factors with their roles, change types, and values. The 'Define Factor Constraints' section has 'None' selected. The 'Model' section shows 'Main Effects' selected, with a list of terms including pH, %DO, Induction TempC, Induction OD600, Feed Rate, pH*pH, and pH*%DO. The 'Design Generation' section has 'Number of Runs' set to 16, and the 'Make Design' button is highlighted with a red box.

Name	Role	Changes	Values
pH	Covariate	Easy	6.74 7.26
%DO	Covariate	Easy	17 43
Induction TempC	Covariate	Easy	39.05 42.95
Induction OD600	Covariate	Easy	17 43
Feed Rate	Covariate	Easy	1.66 3.74

53

THE PLASMID DNA (pDNA) EXPERIMENT

This screenshot shows the same 'Custom Design' window as above, but with the 'Make Design' button clicked. A design table is now visible on the right side of the window, showing 16 rows of experimental runs with columns for various factors and the resulting pDNA titer.

Run	pH	%DO	Induction TempC	Induction OD600	Feed Rate	pDNA Titer mg/L
1	6.74	17	39.05	17	1.66	285.60
2	7.26	17	39.05	17	1.66	364.00
3	6.74	43	39.05	17	1.66	348.08
4	7.26	43	39.05	17	1.66	434.74
5	6.74	17	42.95	17	1.66	339.74
6	7.26	17	42.95	17	1.66	154.46
7	6.74	43	42.95	17	1.66	430.35
8	7.26	43	42.95	17	1.66	341.00
9	6.74	17	39.05	43	1.66	303.82
10	7.26	17	39.05	43	1.66	398.00
11	6.74	43	39.05	43	1.66	411.74
12	7.26	43	39.05	43	1.66	517.23
13	6.74	17	42.95	43	1.66	338.68
14	7.26	17	42.95	43	1.66	229.00

54

THE PLASMID DNA (pDNA) EXPERIMENT

	pH *	%DO *	Induction TempC *	Induction OD600 *	Feed Rate *	pDNA Titer mg/L *
1	6.8	30	42.5	40	1.9	285.60
2	7.0	30	41.0	30	2.7	364.00
3	7.0	30	41.0	30	2.7	348.08
4	7.0	30	41.0	30	2.7	434.74
5	7.0	30	41.0	30	2.7	339.74
6	7.2	40	42.5	20	1.9	154.46
7	7.0	30	41.0	30	2.7	430.35
8	6.8	20	42.5	40	3.5	341.00
9	7.2	20	42.5	20	3.5	303.82
10	7.2	30	39.5	20	3.5	398.00
11	7.0	30	41.0	30	2.7	411.74
12	6.8	20	39.5	20	3.5	517.23
13	6.7	30	41.0	30	2.7	338.68
14	7.2	20	41.0	40	1.9	229.00
15	7.0	30	41.0	17	2.7	282.29
16	6.8	40	41.0	20	3.5	377.00

55

THE PLASMID DNA (pDNA) EXPERIMENT

	pH *	%DO *	Induction TempC *	Induction OD600 *	Feed Rate *	pDNA Titer mg/L *	Bayesian I Optimal Subset n=16
1	6.8	30	42.5	40	1.9	285.60	0
2	7.0	30	41.0	30	2.7	364.00	0
3	7.0	30	41.0	30	2.7	348.08	0
4	7.0	30	41.0	30	2.7	434.74	0
5	7.0	30	41.0	30	2.7	339.74	0
6	7.2	40	42.5	20	1.9	154.46	1
7	7.0	30	41.0	30	2.7	430.35	0
8	6.8	20	42.5	40	3.5	341.00	0
9	7.2	20	42.5	20	3.5	303.82	0
10	7.2	30	39.5	20	3.5	398.00	1
11	7.0	30	41.0	30	2.7	411.74	0
12	6.8	20	39.5	20	3.5	517.23	0
13	6.7	30	41.0	30	2.7	338.68	1
14	7.2	20	41.0	40	1.9	229.00	1
15	7.0	30	41.0	17	2.7	282.29	0
16	6.8	40	41.0	20	3.5	377.00	0

56

THE PLASMID DNA (pDNA) EXPERIMENT

	pH	%DO	Induction TempC	Induction OD600	Feed Rate	pDNA Titer mg/L	Bayesian I Optimal Subset n=16
1	6.8	30	42.5	40	1.9	285.60	0
2	7.0	30	41.0	30	2.7	364.00	0
3	7.0	30	41.0	30	2.7	348.08	0
4	7.0	30	41.0	30	2.7	434.74	0
5	7.0	30	41.0	30	2.7	339.74	0
6	7.2	40	42.5	20	1.9	154.46	1
7	7.0	30	41.0	30	2.7	430.35	0
8	6.8	20	42.5	40	3.5	341.00	0
9	7.2	20	42.5	20	3.5	303.82	0
10	7.2	30	39.5	20	3.5	398.00	1
11	7.0	30	41.0	30	2.7	411.74	0
12	6.8	20	39.5	20	3.5	517.23	0
13	6.7	30	41.0	30	2.7	338.68	1
14	7.2	20	41.0	40	1.9	229.00	1
15	7.0	30	41.0	17	2.7	282.29	0
16	6.8	40	41.0	20	3.5	377.00	1

57

THE PLASMID DNA (pDNA) EXPERIMENT

	pH	%DO	Induction TempC	Induction OD600	Feed Rate	pDNA Titer mg/L	Bayesian I Optimal Subset n=16
1	6.8	30	42.5	40	1.9	285.60	0
2	7.0	30	41.0	30	2.7	364.00	0
3	7.0	30	41.0	30	2.7	348.08	0
4	7.0	30	41.0	30	2.7	434.74	0
5	7.0	30	41.0	30	2.7	339.74	0
6	7.2	40	42.5	20	1.9	154.46	1
7	7.0	30	41.0	30	2.7	430.35	0
8	6.8	20	42.5	40	3.5	341.00	0
9	7.2	20	42.5	20	3.5	303.82	0
10	7.2	30	39.5	20	3.5	398.00	1
11	7.0	30	41.0	30	2.7	411.74	0
12	6.8	20	39.5	20	3.5	517.23	0
13	6.7	30	41.0	30	2.7	338.68	1
14	7.2	20	41.0	40	1.9	229.00	1
15	7.0	30	41.0	17	2.7	282.29	0
16	6.8	40	41.0	20	3.5	377.00	1

58

THE PLASMID DNA (pDNA) EXPERIMENT

	pH	%DO	Induction TempC	Induction OD600	Feed Rate	pDNA Titer mg/L	Bayesian I Optimal Subset n=16
1	6.8	30	42.5	40	1.9	285.60	0
				30	2.7	364.00	0
				30	2.7	348.08	0
				30	2.7	434.74	0
				30	2.7	339.74	0
				20	1.9	154.46	1
				30	2.7	430.35	0
				40	3.5	341.00	0
				20	3.5	303.82	0
				20	3.5	398.00	1
				30	2.7	411.74	0
				20	3.5	517.23	0
				30	2.7	338.68	1
14	7.2	20	41.0	40	1.9	229.00	1
15	7.0	30	41.0	17	2.7	282.29	0
16	6.8	40	41.0	20	3.5	377.00	1

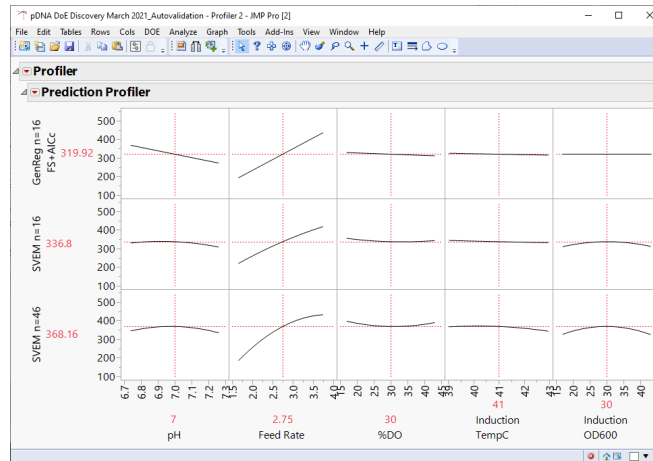
59

THE PLASMID DNA (pDNA) EXPERIMENT

Bayesian I Optimal Subset n=16	Predictor	Creator	.2.4.6.8	RSquare	RASE	AAE	Freq
0	GenReg n=16 FS+AICc	Fit Generalized Forward Selection		0.2192	91.452	72.827	60
0	SVEM n=16			0.5023	73.011	58.265	60
1	GenReg n=16 FS+AICc	Fit Generalized Forward Selection		0.7885	48.687	40.628	32
1	SVEM n=16			0.7740	50.327	38.406	32

60

THE PLASMID DNA (pDNA) EXPERIMENT

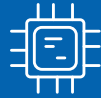


61

OBSERVATIONS AND FURTHER QUESTIONS

- Diminishes the role of p-value based thinking.
- Establish best practice for uncertainty analysis of predictions
- What are the best families of base models?
- What kinds of designs do we use?
- What is the role of screening experiments?
- What are power and sample size considerations?
- What about blocking and split-plot experiments?

62



Philip Ramsey

Case-Study & SVEM Product Demonstration

63



Philip Ramsey

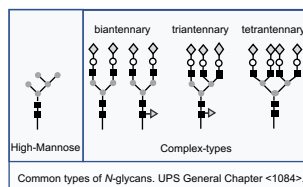
Characterizing a Glycoprofiling Analytical Method for Therapeutic Proteins using SVEM and Bayesian I- optimal Designs

64

Glycoproteins are the largest group of biologically-derived drugs.

ICH Q6B guideline requires extensive physicochemical characterization of biopharmaceuticals including inherent structural heterogeneity due to glycosylation (post-translationally modified) and lot-to-lot consistency is required.

- Carbohydrate content.
- Carbohydrate chain structure.
- Oligosaccharide pattern (antennary profile).
- Glycosylation site.



Currently, there is a lack of a universally accepted analysis technique for glycosylation characterization.

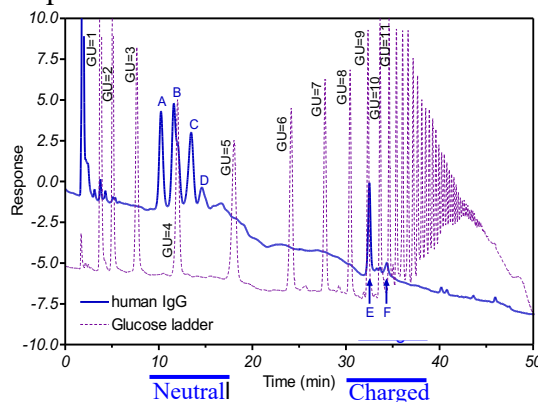
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65

65

The goal is to develop a robust, cost effective characterization method.

Two independent experiments was used to characterize & optimize an HPAE-PAD¹ method. The approach uses a glucose ladder (GU) as a reference to identify glycoform peaks from an actual human antibody sample.



Glycoform glycan	G-Unit
A	3.59
B	3.89
C	4.23
D	4.42
E	9.17
F	10.8

¹High Performance Anion Exchange Chromatography with Pulsed Amperometric Detection

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66

66

The first experiment consisted of a 16 run, 3-level design.

The second experiment consisted of 28 run, 3-level design.

Since the two experiments are independent, one set can be used as a training set and the second set as a validation set.

For purposes of today's discussion, both designs were combined into a single 44 run design.

The candidate selector in Custom Design (JMP 16) was used to create Bayesian I-optimal designs from the 44 combined runs:

- **Design #1 n = 10 runs**, Full Quadratic Model specified
- **Design #2 n = 13 runs**, Full Quadratic Model specified
- **Design #3 n = 16 runs**, Partial Cubic Model specified

The runs not selected are used as a validation set for predictive models.

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67

67

Traditionally in statistics the full quadratic model (FQM) is used to build models to optimize physical processes. The FQM has the following mathematical form.

$$Y = \beta_0 + \sum_{i=1}^K \beta_i X_i + \sum_{i=1}^{K-1} \sum_{j=i+1}^K \beta_{ij} X_i X_j + \sum_{i=1}^K \beta_{ii} X_i^2$$

Although popular for optimization the FQM is often not sufficient to model the complex kinetics over the experimental region.

Cornell and Montgomery (1998) suggested augmenting the FQM with partial cubic (PC) terms that allow more flexibility to fit the complex kinetics in physical systems. The form of the PC model is:

$$Y = \beta_0 + \sum_{i=1}^K \beta_i X_i + \sum_{i=1}^{K-1} \sum_{j=i+1}^K \beta_{ij} X_i X_j + \sum_{i=1}^K \beta_{ii} X_i^2 + \sum_{i=1}^K \sum_{j \neq i}^K \beta_{ij} X_i^2 X_j + \sum_{i=1}^K \sum_{j \neq i}^K \beta_{ij} X_i X_j^2 + \sum_{i=1}^K \sum_{j \neq i}^K \beta_{ij} X_i^2 X_j^2$$

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68

68

Five factors were selected to manipulate in the experiments.

Factor (level)	-1	0	1
Initial %NaOAc (% A)	0	10	20
Initial %NaOH (% B)	30	40	50
Gradient_01-12 (mM NaOAc /min)	0.415	1.25	2.085
Gradient_12-24 (mM NaOAc /min)	1.25	2.085	2.915
Gradient_24-42 (mM NaOAc /min)	4.72	5.555	6.39

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69

69

Two responses were chosen to optimize in the experiment; a total of 28 responses exist.

Retention Time for glycan 3 (RT_G03) was most important as it anchors the position of the glucose ladder used to identify specific glycan peaks.

The second response is peak resolution for charged glycan G10 (Resol_G10).

Glycan G10 elutes late with a number of other charged glycans.

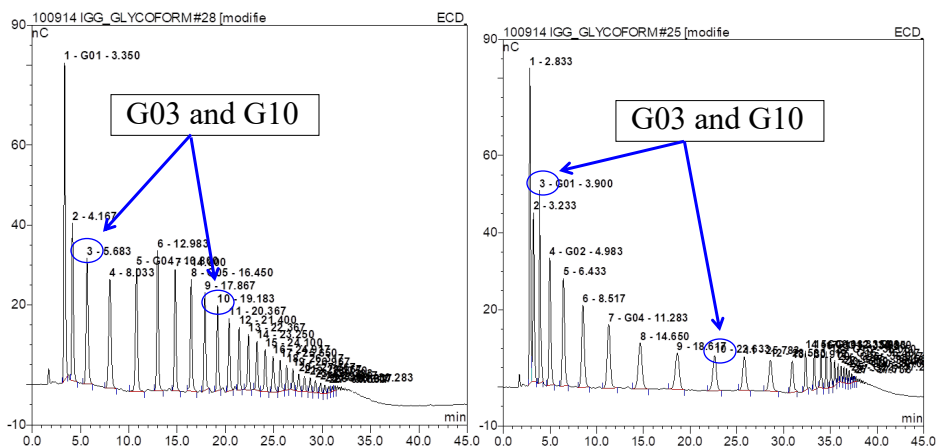
Response	Description	Optimization
RT_G03	Retention Time	Target ~ 8.5 min
RT_G10	Retention Time	None

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70

70

Below are pictures of typical chromatograms from two different runs.
Notice the shapes change markedly with experimental conditions.



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71

71

The original experimental designs and the experimental work was performed by Dr. Eliza Yeung of Cytovance Biologics, OKC.

Dr. Yeung is the Director of Process Characterization for Cytovance.

We thank Dr. Yeung and Cytovance for their willingness to share the experimental results.

Dr. Yeung is an outstanding scientist and a JMP user.



Eliza Yeung, Ph.D.

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72

72

A limitation of the PC model for traditional statistical modeling methods is the number of potential PC model terms.

For K factors the full PC model has $2K^2+1$ terms including the intercept; e.g., for K = 5 there exist 51 potential model terms.

Using **SVEM** it is actually possible to fit full PC models with relatively small number of design points.

We illustrate **SVEM** with the case study.

All of the SVEM modeling was performed using the **SVEM addin available from Predictum Inc.**

If interested in the addin send email to Wayne Levin at Predictum: levin@predictum.com.

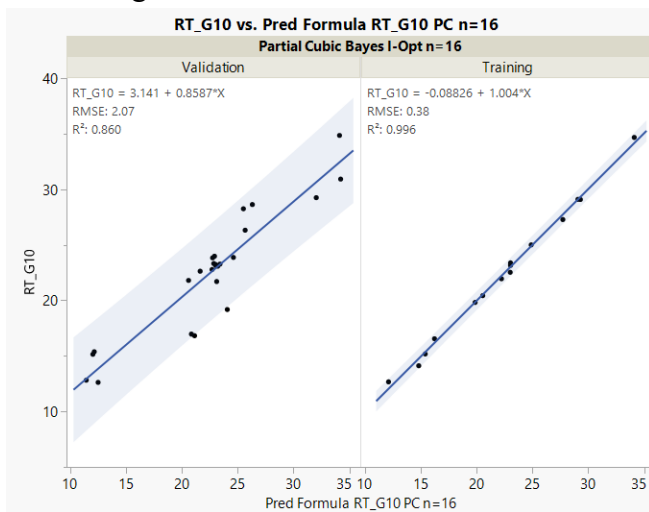
73

Below is summary table of the SVEM modeling results. All SVEM runs used Best Subsets with Max Model Size = 5 and Nboot = 1,000 FWB runs for each model.

Bayesian I-optimal Design	Response	Full Model (No. Predictors)	RASE Training	RASE Validation	R ² Validation
N = 16	RT_G03	PC (40)	0.190	0.387	0.98
N = 16	RT_G10	PC (40)	0.353	2.148	0.84
N = 13	RT_G03	PC (40)	0.065	0.480	0.97
N = 13	RT_G10	PC (40)	0.172	2.671	0.75
N = 10	RT_G03	FQ (20)	0.225	0.730	0.94
N = 10	RT_G10	FQ (20)	0.240	2.354	0.75

74

Below are typical actual by predicted plots for RT_G10 and N = 16, the hardest modeling scenario.

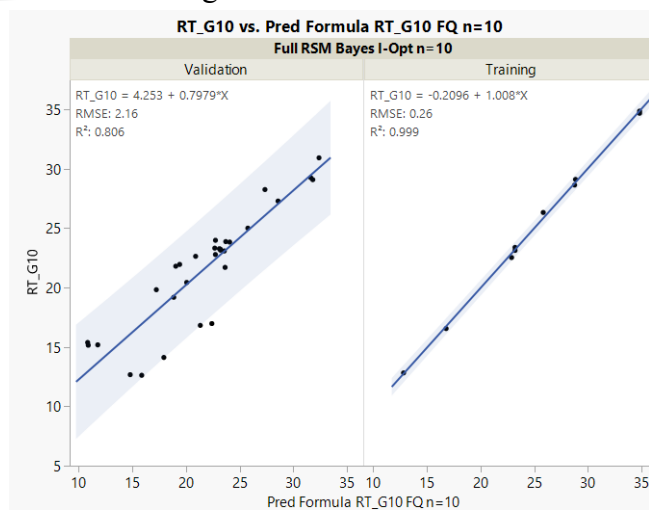


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75

75

Below are typical actual by predicted plots for RT_G10 and N = 10, the more difficult modeling scenario.

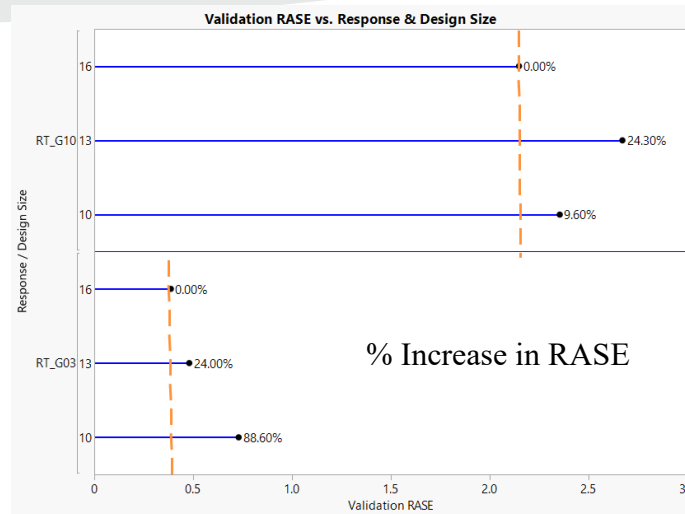


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76

76

Validation RASE values vs Design Size for RT_G03 and RT_G10.



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77

77

SVEM provides the ability to fit supersaturated models ($p > n$) that are often needed for prediction in complex physical systems.

With SVEM we combine methods from machine learning with Design of Experiments.

Modeling with SVEM allows one to use highly efficient experimental designs speeding up the pace of innovation and reducing overall cost of experimentation.

Bayesian I-optimal designs, available in Custom Design, have the potential to substantially decrease the sizes of experimental designs when use in combination with SVEM.

With a 10 run Bayesian I-optimal design we were able to use SVEM to fit a 20 predictor model that successfully predicted responses on a true validation dataset.

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78

78

SVEM Add-in



Predictum

- Capability Analysis
- Auto Word Output
- Developer Panel
- Self-Validating Ensemble Modeling**
- Demo Central

	pH	%DO	Induction Temperature C	Induction OD600	Feed rate	pDNA Titer mg/L
1	7.0	40	42.5	20	1.9	156.20
2	7.0	20	39.5	40	3.5	318.45
3	7.2	30	39.5	20	3.5	398.00
4	6.8	30	42.5	40	1.9	285.60
5	7.2	20	41.0	40	1.9	229.00
6	6.8	40	41.0	20	3.5	377.00
7	7.2	20	42.5	30	3.5	290.00
8	6.8	40	39.5	30	1.9	123.00
9	7.2	40	42.5	40	2.7	299.00
10	6.8	20	39.5	20	2.7	428.00
11	7.0	30	41.0	30	2.7	327.80
12	7.0	30	41.0	30	2.7	339.74
13	7.0	30	41.0	30	2.7	387.35
14	7.0	30	41.0	30	2.7	393.97
15	7.0	30	41.0	30	2.7	348.08

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79

Fit Full Quadratic Model +



Report: Fit Model

Model Specification

Select Columns

- 9 Columns
- ▲ pH
- ▲ %DO
- ▲ Induction Temperature C
- ▲ Induction OD600
- ▲ Feed rate
- ▲ pDNA Titer mg/L
- ▲ Null Factor
- ▲ Paired Fractionally ...hted Bootstrap Weight
- ▲ Validation

Pick Role Variables

Y: **pDNA Titer mg/L** (optional)

Freq: Paired Fra...rap Weight

Validation: Validation

Personality: Generalized Regression

Distribution: Normal

Censor Code: []

Buttons: Help, Run, Recall, Keep dialog open, Remove

Construct Model Effects

Add: Null Factor, pH, pH*pH, %DO, %DO*%DO, Induction Temperature C, Induction Temperature C*Induction Temperature C, Induction OD600, Induction OD600*Induction OD600, Feed rate

Attributes: [v] Transform: [v] No Intercept: []

Fit the full quadratic model with the addition of interactions between main effects and quadratic terms.

80

Saving the Prediction Formula

S-VEM Settings at bottom

Generalized Regression for pDNA Titer mg/L

Model Launch

Singularity Details

Response Distribution: Normal

Estimation Method: Lasso

Adaptive

Advanced Controls

Validation Method: Validation Column

Early Stopping

S-VEM settings

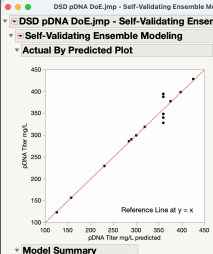
Number of simulations:

Random sampling of effects

RASE-weighted averaging

Go

S-VEM Output



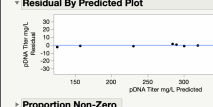
Model Summary

Response: pDNA Titer mg/L
 Distribution: Normal
 Estimation Method: Lasso
 Validation Method: Auto Validation
 Mean Model Link: Identity
 Scale Model Link: Identity

Measure

	Training
Number of rows	15
RISquare	0.0675011
RMSE	15.29963
Number of Simulations	1000

Residual By Predicted Plot



Proportion Non-Zero

Parameter Estimates

predictum® [Feedback to Predictum](#)

Saved Prediction Formula

```

3630.236924
+ -118.6664268 • %DO
+ 194.18576035 • Feed rate
+ -22.37217396 • Induction OD600
+ 23.922831578 • Induction Temperature C
+ -467.4523557 • pH
+ -0.390109988 • %DO2
+ -44.55917313 • Feed rate2
+ -0.06801447 • Induction Temperature C2
+ -0.051062686 • pH2
+ 5.8496304578 • %DO • Feed rate
+ 0.0091653996 • %DO • Induction OD600
+ 0.0079227233 • %DO • Induction Temperature C
+ 17.508688819 • %DO • pH
+ -7.520564e-7 • Feed rate • Induction OD600
+ -1.071051792 • Feed rate • Induction Temperature C
+ 0.5041794505 • Induction OD600 • Induction Temperature C
+ -4.006032248 • Induction Temperature C • pH
                    
```

81



ANNOUNCEMENT


ON-DEMAND COURSE MIXTURE DESIGN EXPERIMENTS

82



83

Next Steps







- 2 ways to try this yourself
 1. Try the Product
 - JMP Pro v14.3 or 15.x
 - JMP Pro trials are available via this [link](#)
 2. Contact Predictum
 - Working with your historical DOEs, we can run a proof-of-concept, comparative analyses for you, on a limited basis
 - Review the analysis with you

Contact Wayne:
levin@predictum.com

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84

84




Self-Validated, Ensemble Models (S-VEM)

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85



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86