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Representative Models for History Matching and Robust Optimization



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Table of Contents

Acknowledgement	i
List of Figures	iv
List of Tables	V
Nomenclature	vi
Abbreviations	vi
Symbols	vi
Alphabetical	vi
Greek	vii
Abstract	1
1. Introduction	2
1.1 Problem statement	2
1.2 Literature Review	2
1.2.1 Parallel Computing	2
1.2.2 Reduced-Order Model (ROM)	3
1.2.3 Representative Models	3
1.3 Hypothesis	3
1.4 Thesis Outline	4
2. Background	5
2.1 Feature and Distance	5
2.2 Clustering	5
2.3 Dimensionality Reduction and Projection	7
2.3.1 Tensor Decomposition	7
2.3.2 Multidimensional Scaling (MDS)	7
2.4 Self-organizing Map (SOM)	8
2.5 Robust Optimization	
2.6 History Matching	
3. Methodology	14
3.1 Representative Model Selection	14
3.1.1 Feature Selection	14
3.1.2 Projection M ethod	
3.1.3 Clustering	
3.1.4 Weighting	15
3.1.5 SOM	
3.2 Validation	
3.3 Workflow of Representative Robust Optimization	

3.4 Workflow of Representative History Matching19
4. Examples and Results
4.1 Cas e Studies
4.1.1 2D Model
4.1.2 3D Model
4.2 Robust Optimization Results
4.2.1 2D Model
4.2.3 3D Model
4.3 History Matching Results
4.3.1 2D Model
4.3.2 3D Model
5. Discussion
6. Conclusion
References
Appendix AI
Appendix BV

List of Figures

Figure 1 Workflow of K-means clustering algorithm	6
Figure 2 Workflow of multidimensional scaling	8
Figure 3 SOM Kohonen network structure. Light pink denotes the winning node, pink denotes the immediat	æ
neighbors and purple denotes further neighbors. (from Kohonen Network - Background Information, 2012)	9
Figure 4 Workflow of SOM	0
Figure 5 Workflow of history matching	3
Figure 6 Workflow for selection of representative models (Repr. Select)	4
Figure 7 Representative model selection using SOM	6
Figure 8 Stages of validation	7
Figure 9 Workflow of robust optimization on full ensemble and representative ensemble. (Left) Unoptimize	d
reference (Middle) Representative robust optimization (Right) Full robust optimization reference	8
Figure 10 Workflow of history matching on full ensemble and representative ensemble. (Left) Full ensemble histor	v
matching (Right) Representative history matching	9
Figure 11: 2D inverted five-spot configuration	0
Figure 12 Permeability image of sixteen randomly chosen realizations from simsim model ensemble 50	1
Figure 13 Permeability image of sixteen randomly chosen realizations from channel model ensemble 50	1
Figure 14 (Left) Reservoir model displaying the position of the injectors (blue) and producers (red). (Right) Si	x
randomly chosen realizations. (from Jansen et al., 2014)	2
Figure 15 Simsim model NPV CDF of 4 ensembles using permeability as feature. (Left) MDS (Right) Tenso)r
decomposition	3
Figure 16 Simsim model NPV CDF of 4 ensembles using NPV time series as feature. (Left) MDS (Right) Tenso)r
decomposition	4
Figure 17 Simsim model NPV CDF of 4 ensembles using oil saturation snapshots as feature. (Left) MDS (Right) Tenso	r
decomposition	4
Figure 18 Simsim model NPV CDF of 4 ensembles using (Left) random selection (Right) oil saturation snapshots wit	h
SOM	5
Figure 19 Mean NPV comparison of MDS and tensor decomposition with four ensembles in Simsim model	5
Figure 20 Channel model NPV CDF of 4 ensembles using permeability as feature. (Left) MDS (Right) Tenso	r
decomposition	6
Figure 21 Channel model NPV CDF of 4 ensembles using NPV time series as feature. (Left) MDS (Right) Tenso	٥r
decomposition2	6
Figure 22 Channel model NPV CDF of 4 ensembles using oil saturation snapshots as feature. (Left) MDS (Right) Tenso	r
decomposition2	7
Figure 23 Channel model NPV CDF of 4 ensembles using (Left) random selection (Right) oil saturation snapshots wit	h
SOM	7
Figure 24 NPV comparison of MDS and Tensor with four ensembles in Channel model 2	8
Figure 25 Egg Model robust optimization results	9
Figure 26 Permeability results of history matching using MDS. (Top) Simsim Model (Bottom) Channel Model	9
Figure 27 Simsim Model field production data of representative ensemble using all oil saturation snapshots at tim	e
1,500 days. Red is oil and blue is water. (Left) The priori (Right) The posterior	0
Figure 28 Simsim Model field production data of posterior of 10 representative realizations using MDS and all o	il
saturation snapshots at various history matching time. Red is oil and blue is water	1
Figure 29 Simsim Model field production data of posterior of 10 representative realizations using MDS and o	il
saturation snapshots up to specified history matching time. Red is oil and blue is water	1
Figure 30 Simsim Model NPV comparison of 10 representative realizations using MDS. Repr prior all and repr post a	11
are selected using all oil saturation snapshots whereas repr prior and repr post are selected using oil saturatio	n
snapshots up to the history matching time. (Left) The prior NPV CDF plot. (Right) The posterior NPV CDF plot 3	2
Figure 31 Channel Model field production data of representative ensemble using all oil saturation snapshots at tim	e
1,500 days. Red is oil and blue is water. (Left) The prior (Right) The posterior	3

Figure 32 Channel Model field production data of posterior of 10 representative realizations using MDS and all oil Figure 33 Channel Model field production data of posterior of 10 representative realizations using MDS and oil Figure 34 Channel Model NPV comparison of 10 representative realizations using MDS. Repr prior all and repr post all are selected using all oil saturation snapshots whereas repr prior and repr post are selected using oil saturation Figure 35 Examples of history matched layer 4 of Egg Model permeability field using (Top) MDS (Bottom) Tensor Figure 36 Egg Model water (blue) and oil (red) production rates. Representative ensembles selected using MDS.....36 Figure 37 Egg Model injector rates of representative ensembles using MDS as projection method. (Left) Prior (Right) Figure 38 Egg Model water (blue) and oil (red) production rates. Representative ensembles selected using tensor Figure 39 Egg Model injector rates of representative ensembles using tensor decomposition as projection method. Figure 41 Egg Model history matching prior and posterior final NPV CDF using full ensemble and representative Figure 42 Simsim model NPV CDF of 6 additional ensembles using permeability as feature. (Left) MDS (Right) Tensor decomposition.....I Figure 43 Simsim model NPV CDF of 6 additional ensembles using NPV time series as feature. (Left) MDS (Right) Tensor decompositionI Figure 44 Simsim model NPV CDF of 6 additional ensembles using oil saturation snapshots as feature. (Left) MDS (Right) Tensor decompositionII Figure 45 Simsim model NPV CDF of 6 additional ensembles using (Left) Random selection (Right) oil saturation snapshots with SOM......II Figure 46 Channel model NPV CDF of 6 additional ensembles using permeability as feature. (Left) MDS (Right) Tensor decomposition.....III Figure 47 Channel model NPV CDF of 6 additional ensembles using NPV time series as feature. (Left) MDS (Right) Tensor decompositionIII Figure 48 Channel model NPV CDF of 6 additional ensembles using oil saturation snapshots as feature. (Left) MDS (Right) Tensor decompositionIV Figure 49 Channel model NPV CDF of 6 additional ensembles using (Left) Random selection (Right) oil saturation snapshots with SOMIV

List of Tables

Table 1: Reservoir Properties for 2D Model	20
Table 2 Reservoir properties for Egg Model	

Nomenclature

Abbreviations

AI	Artificial intelligence
ANN	Artificial neural network
CDF	Cumulative distribution function
CLRM	Closed-loop reservoir management
KS test	Kolmogorov-Smirnov Test
MDS	Multidimensionalscaling
NPV	Net present value
PCA	Principal Component Analysis
POD	Proper orthogonal decomposition
Post.	Posterior
Repr.	Representative
ROM	Reduced-order model
SOM	Self-organizing Map
SSE	Sum of squared error
SVD	Singular value decomposition
TPWL	Trajectory piecewise linearization
UQ	Uncertainty quantification
VOI	Value of information

Symbols

Alphabetical

Α	Amplitude of neighborhood adaptation
С	Covariance matrix
С	Cluster set
С	K-means centroid
D	Dissimilarity matrix
D	Projected dissimilarity matrix
dI	Projected distance in low dimensional space
â	Monotonic transformation of dissimilarity
d _{obs}	Observation data vector
<i>e</i> ²	Squared error criterion
i _{it}	SOM iteration number
J	Objective function
Κ	Number of clusters
М	Number of feature data dimension
Μ	Ensemble parameter matrix
M _{full}	Full ensemble parameter matrix
M _{repr.}	Representative ensemble parameter matrix
M ^{prior}	Prior ensemble parameter matrix
M ^{post} .	Posterior ensemble parameter matrix
m	Model parameters vector
\mathbf{m}_{full}	Full model parameters vector
m _{repr.}	Representative model parameters vector
m ^{prior}	Prior model parameter
$\widetilde{\mathbf{m}}^{prior}$	Projected prior model parameter

m ^{post.}	Posterior model parameter vector			
$\widetilde{\mathbf{m}}^{post.}$	Projected posterior model parameter vector			
Ν	Number of Kohonen nodes			
N _{repr}	Number of representative realizations			
R	Number of realizations			
S	3D tensor			
S	Stress function			
u	Control input vector			
W	Kohonen weight vector			
w _{repr}	Representative Realizations weight vector			
X	2D grid property matrix			
x	State vector			
Greek				
α	Monotonic decreasing learning coefficient			
δ	Pairwise distance (dissimilarity)			
Θ	Feature data matrix			
Õ	Projected feature data matrix			
Θ_{sub}	Sub ensemble feature data matrix			
θ	Feature data vector			
μ	Mean			
μ_{NPV}	Ensemble mean			
σ	Standard deviation			
φ	1 st orthonormal basis functions (vectors)			
Ψ	2 nd orthonormal basis functions (vectors)			
X	3 rd orthonormal basis functions (vectors)			

Abstract

Reservoir management has been widely implemented in the petroleum industry to attain the best performance out of the asset. Highly efficient computer-assisted reservoir management is getting more common and therefore enabling the incorporation of ensembles to provide uncertainty quantification (UQ). Closed-loop reservoir management (CLRM) further enhances reservoir management by combining robust optimization and history matching while accounting for UQ. However, CLRM workflow is very computationally intensive. In addition, value of information (VOI) workflows that make use of CLRM framework are currently unfeasible mainly due to multiplication of the already immense computational cost required. Therefore, this thesis proposes a method to select representative models forming a reduced ensemble that can replace the full ensemble in robust optimization and history matching.

We use clustering algorithm to select the representative models. Features were extracted based on various model parameters and projected into lower dimensional space using ordination techniques. Different number of representative models were investigated to explore the performance and discover the minimal number of models required in a representative ensemble.

The method is tested in two simple 2D models and in a larger 3D Model. The results showed a very promising future for representative ensembles to be applied in robust optimization where an order of magnitude speedup is estimated. Whereas the implementation of representative ensembles in history matching may require higher number of representative models, although achieving a commendable result. Depending on the size of original ensemble, using reduced ensemble can greatly decrease the computational cost associated with optimization and simulation while providing very comparable results to using full ensemble.

1. Introduction

1.1 Problem statement

In the exploration and production sector, companies have to make difficult decisions regarding the development strategies for their assets. The fact that every reservoir is the only one of its kind means that there is no room for experiments to be carried out to determine the best development strategy. Because of that, numerical simulations are extensively used in reservoir engineering to characterize the reservoirs and predict their production. In essence, simulation models are populated with parameters derived from all the available data from the reservoir (e.g., lithological and pore fluid data). Due to the limited knowledge of the true reservoir, it is common to generate more than one interpretation from collected field data, which, in most cases, results in several models of the subsurface.

To extensively account for uncertainty in reservoir model parameters, an ensemble of reservoir realizations is employed in most modern reservoir management workflows. Uncertainty quantification (UQ) is of major interest in the petroleum industry where quantitative characterization and assessment of uncertainties are paramount to reduce the risk for all field development. While considering an ensemble of models allows to better define, ideally, all possible reservoir characteristics, it leads to additional demand in computational cost to achieve optimal reservoir management.

Closed-loop reservoir management (CLRM) framework combines model-based life-cycle optimization and computerassisted history matching (Jansen et al., 2009) to maximize the reservoir performance (i.e., recovery or financial measures) and obtain the optimal strategy for reservoir management. In a nutshell, CLRM makes use of data collected throughout the reservoir life-cycle to update the reservoir models which, in turn, improve the optimization of the field production strategy.

In combination with UQ by using an ensemble of model realizations, the computational cost of CLRM workflows increases significantly. Workflows to assess value of information (VOI) in CLRM with UQ as proposed by Barros et al. (2016a) further multiply the cost of computation by order of tens or hundreds, making real-field implementation unfeasible with the current advancement in computational power. Therefore, some alternatives are needed to reduce the computational cost to an acceptable range.

1.2 Literature Review

Since the main barrier for a wide implementation of VOI assessment is the computational cost; techniques for accelerating these workflows need to be found. Three main categories of solutions for speeding-up simulations are identified. The first one corresponds to the 'brute-force' approach by increasing computing power to solve larger problems. The second method seeks to speedup simulations by using surrogate models (i.e., approximate or proxy models). And the third one aims at reducing the number of required simulations directly by approximating UQ (i.e., considering few representative models).

1.2.1 Parallel Computing

This may be the simplest approach where the great amount of simulations is simply solved by increasing the computing power. The advancement of ever faster processors and the advent of parallel computing have made this approach rather attractive for companies that are well-funded.

Despite showing many advantages, the development of software for parallel computing can be very complex. Ouenes et al. (1995) and Salazar et al. (1996) have shown that it is possible to have parallel computers and a network of workstations to run simulations without modification by using Parallel Virtual Machine. Schiozer (1999) introduced Module for Parallel Simulations which uses Parallel Virtual Machine to distribute the simulations efficiently by taking into account the speed and dynamic characteristics of each machine. He also concluded that, by using Module for Parallel Simulations in parallel computing, it is possible to reduce the cost of hardware by automating the simulation process and taking advantage of idle workstations.

1.2.2 Reduced-Order Model (ROM)

The aim of ROM is to create a simpler model that can to an extent accurately reproduce the output of a simulator. A good ROM should be accurate while requiring significantly less computational cost than the full-order model.

One approach is implemented by reconstructing a grid based numerical model into a coarser grid model. This approach is considered as grid-based reduced-order model and can be constructed using either upscaling or multiscale method. The former have the disadvantage of losing finer grid resolution while the latter retain information on finer scale commonly with dual-grid methods that are coupled by the prolongation (coarse to fine) and restriction (fine to coarse) operators. However, multiscale methods require extra computation of the operators before simulation, thus limiting the potential for speedup of our workflows. Krogstad et al. (2011) have shown that using multiscale methods can achieve a speedup of an order magnitude in water flooding optimization.

Another approach is to use snapshots of time-variant problems to create basis functions in order to have a reduced model. This method utilizes proper orthogonal decomposition (POD) to calculate the basis functions. However, the large number of variable changes in history matching heavily diminishes the speedup of POD when applied in reservoir simulation. He (2013) utilized trajectory piecewise linearization (TPWL) in conjunction with POD for model-order reduction in history matching whereas Hewson (2015) tested on ensemble-based robust optimization. This method achieved an order of magnitude acceleration in terms of simulation time although it sacrifices the accuracy of the results. However, when accounting for the preprocessing required to build the proxy models, the application of POD-TPWL in CLRM workflow achieved a speedup that is much lower than in simulation alone.

Insuasty et al. (2015a) introduced tensor-based ROM using tensor decomposition and representations of flow characteristics to quantify the features of flow simulations. They compared the tensor approach to POD for adjoint-based optimization where the tensor approach achieved better financial performance. They also showed that tensor models provide higher approximation accuracy over classical POD models, although the computational gain is low.

1.2.3 Representative Models

Sarma et al. (2013) proposed a method for selecting representative realizations for UQ. They claim that their minimax method is able to efficiently select a few reservoir models from a large ensemble by matching target percentiles of multiple output responses while obtaining maximally different models in the parameter uncertainty space. The idea behind the minimax method is to select representative models that are statistical representative while maximizing the spread in the parameter uncertainty space. The authors also claimed that the solution from minimax is generally better than clustering and that the computation is orders of magnitude faster. Although their method ensures good spread of selected models in parameter and output spaces, Sarma et al. (2013) did not address the effectiveness of using selected representative models in optimization or history matching workflows.

Insuasty et al. (2015b) proposed a measure of dissimilarity that is based on reservoir flow patterns in numerical simulation using flow variables such as oil saturation. They suggest that, by applying tensor decomposition on spatial-temporal representation of the reservoir flow patterns (e.g., snapshots of the temporal evolution of oil saturation distribution), the structure of the flow data can be preserved, which allows to better determine dominant flow patterns. Tensor decomposition provides a dissimilarity measure that offers low dimension data and thus easier to be used for model classification. Clustering of realizations from an example ensemble were compared in the paper, the results showed this method is able to provide better defined clusters compared to singular value decomposition (SVD. Insuasty et al. (2015b) showed that, by reducing the number of realizations from 1,000 to 50 realizations in flow-relevant ensemble, it is possible to obtain similar optimal production strategy and yield similar final net present value (NPV) distribution.

1.3 Hypothesis

While an ensemble can more effectively quantify uncertainties, it is arguably unnecessary for robust optimization (*Section 2.5 Robust Optimization*) and history matching (*Section 2.6 History Matching*) procedures to require hundreds of model realizations. By grouping similar realizations and selecting representative realizations for each group we can greatly reduce the ensemble size as well as directly decrease the computational cost of our workflows. Many studies have used representative ensemble for UQ, but very few have focused on using representative ensembles for

optimization. Of course, the implementation of representative ensembles need to have similar effect on robust optimization and history matching which are the main principles in CLRM. This thesis investigates whether, with clever selection of representative realizations to represent the original full ensemble, robust optimization and history matching can be carried out more efficiently while performing close to the full ensemble. This thesis also makes an attempt to understand the fundamentals of selecting good representative models and, more importantly, to determine the bare minimum of representative models needed for an accurate representation of the full ensemble.

1.4 Thesis Outline

In *Chapter 2. Background* we discuss about the fundamentals of robust optimization, history matching and techniques required for representative model selection. *Chapter 3. Methodology* introduces the procedure used in order to select and construct the representative ensembles followed by techniques on validating the performance. *Chapter 4. Examples* describes the case study examples and presents the results on the performance of representative ensemble. *Chapter 5. Discussion* presents the summary and challenges faced using the proposed method followed by various reasonings and future works that may to improve the method. Finally, *Chapter Conclusion* wrap up by presenting the essential findings of the thesis.

2. Background

2.1 Feature and Distance

Feature describes the property selected in order to distinguish between realizations. The feature data matrix $\boldsymbol{\Theta} = [\boldsymbol{\theta}_1 \ \boldsymbol{\theta}_2 \dots \boldsymbol{\theta}_R]$ contains the feature data vectors $\boldsymbol{\theta}$ of the individual realizations, which have dimension *M*. $\boldsymbol{\Theta}$ can be used for selecting representative realizations.

Here 'distances' are measures of dissimilarity between realizations. They are always defined as pairwise distances in terms of any form of feature of a reservoir. In petroleum engineering, the distances can be generally categorized into static and dynamic. Static distances are calculated from initial grid based properties or parameters (e.g., permeability and initial oil saturation). Dynamic distances, on the other hand, require simulated properties (e.g., NPV, streamlines and oil production rates). Suzuki et al. (2008) and Caers et al. (2010) have used permeability as a distance measure to differentiate model realizations based on geological features. Van Essen et al. (2009) and Jansen et al. (2009) have shown that, although reservoir models might have different geological properties, they may generate the same NPV under individually optimized strategies. Scheidt and Caers (2009) and Scheidt et al. (2011) use the cumulative oil and water production rates as the dissimilarity measures to assess the flow uncertainty. Park and Caers (2007), Scheidt et al. (2009) and Scheidt and Caers (2009) used streamline simulators to produce fast characterization of cumulative oil and water production to distinguish models.

The distance measures can be computed in multiple ways (e.g., Euclidean, Manhattan, and Minkowski). The Euclidean distance, defined as the 'straight-line' distance between two points in Euclidean space, is the one used in this thesis. It is formulated as

$$\delta_{ij} = \sqrt{\left\| \mathbf{\theta}_i - \mathbf{\theta}_j \right\|^2}$$
 ,

where δ_{ij} is the pairwise distance between realization *i* and *j* in terms of the selected feature. Given a set of *R* realizations, **D** is be the dissimilarity matrix of *R* × *R* containing the distance between two realizations δ_{ij} . δ_{ij} is be 0 when *i*=*j* as there is no dissimilarity among itself and δ_{ij} is equal to δ_{ii} .

Equation 2

Equation 1

$$\mathbf{D} = \begin{bmatrix} \delta_{11} & \cdots & \delta_{1N} \\ \vdots & \ddots & \vdots \\ \delta_{N1} & \cdots & \delta_{NN} \end{bmatrix} \; .$$

2.2 Clustering

One of the most obvious ways of finding representative models is to group them according to some criteria and select a model out of each group. Clustering analysis is a family of techniques used to partition a set of similar points (i.e. objects or observations) into clusters. Cluster analysis aims to unveil the internal organization of a dataset by detecting the structure within the data in the form of clusters. The goal of clustering is to categorize similar data together. Therefore it is useful for reducing the amount of data. Such uses of grouping are pervasive in how humans process information. Cluster analysis using numerical methods were introduced in biological classifications (Jardine and Sibson, 1971; Sneath and Sokal, 1973) and have been used in pattern recognition (Anderberg, 2014), image processing (Jain and Flynn, 1996), machine learning (Arabie and Hubert, 1996) and many other domains such as psychology, geology, marketing and archaeology (Jain et al. 2009).

At the top level of cluster analysis classification, there is an important distinction between hierarchical and partitional approaches. Hierarchical clustering produces nested groupings based on criteria for merging or splitting clusters. The nested groupings in hierarchical clustering can thus be presented in a dendrogram. On the other hand, partitional clustering separates the points into exclusive clusters by optimizing a defined criterion function. The most common criterion function used is the squared error criterion which performs well with compact and isolated clusters (Jain et al., 2009). The squared error criterion is defined by

Equation 3

$$e^{2} = \sum_{j=1}^{K} \sum_{i=1}^{n_{j}} \left\| x_{i}^{j} - c_{j} \right\|^{2}$$

where x_i^j is the *i*th point in cluster *j*, c_j is the centroid of cluster *j* and *K* is the predefined number of clusters and n_j is the nth point in cluster *j*.



Figure 1 Workflow of K-means clustering algorithm

K-means clustering is the simplest and most common clustering algorithm (McQueen, 1967; Caers, 2011) that employ the squared error criterion. The name K-means comes from the technique itself whereby it tries to partition data into *k* clusters of which each data point belongs to the cluster with the nearest mean (i.e., the centroid). *Figure 1* illustrates the workflow of K-means clustering algorithm where it starts with predefined *k* number of randomize centroid placement. Next, all points are assigned to the nearest centroid. The mean of each cluster is then calculated and the centroid is reassigned to the new mean. The criterion function is then calculated and reassignment of all points to new centroids is carried out if convergence is not met. Typically we minimize some measure of dissimilarity in the samples within each cluster (i.e., intra-cluster distance), while maximizing the dissimilarity between clusters (i.e., inter-cluster distance). With user predefined number *K* sets of cluster C_k , where k = 1, 2, ..., K, of which individual C_k contains n_k unique indices. When using K-means clustering on a fixed indices dataset and applying squared error criterion (*Equation 3*), K-means clustering can then be stated as an optimization problem defined as

Equation 4

$$C_{opt} = \arg\min_{C} \sum_{k=1}^{K} \sum_{i \in C_{k}} ||x_{i} - c_{k}||^{2}$$

where $c_k = \frac{1}{n_k} \sum_{i \in C_k} x_i$. In a sentence, K-means clustering is an iterative process which partitions the data by minimizing the within cluster sum of point-to-cluster centroid distances over all clusters. Many clustering algorithms suffer from inefficiency when performed on high dimensional data due to the inherent sparsity of data: as the number of dimensions increases, the distance measures becomes equidistant (Berchtold et al., 1997; Parsons et al., 2004). Therefore, dimensionality reduction is recommended to treat the high dimensional data before clustering.

Another major issue with K-means algorithm is the sensitivity to the initial randomized partition that may lead to local minimum convergence. However, this problem can be mitigated by repeating the clustering process with different random seeds. Many variant of K-means clustering have been introduced to improve and add new attributes. Arthur and Vassilvitskii (2007) proposed a useful variant named K-means++ which chooses the initial values that try to spread out clusters' centroids. Shindler (2008) pointed out that K-means++ is able to overcome some of the problems associated with defining initial clusters' centroids compared to K-means algorithm. In general, K-means clustering is a robust way for grouping seemingly unrelated data spread and is widely used in engineering to group scattered data points.

2.3 Dimensionality Reduction and Projection

High dimensional data suffer from a few drawbacks. The most inconvenient one concerns the data containing excessive and often unneeded information. When operations are carried out on high dimensional data, unwanted effects such as data over-fitting and suboptimal search are likely to occur, besides increasing the computational cost. The goal of projections is to represent the parameters in lower dimensional space that preserve certain properties of the data structure as faithfully as possible. Therefore, projections can be very helpful in providing a better dataset for further operations. Aggarwal et al. (1999) have shown that the projection of high dimensional data spaces into low dimension subspaces leads to improved clustering results.

2.3.1 Tensor Decomposition

One projection method is proposed by Insuasty et al. (2015) where dimension reduction is achieved through tensor decomposition. Tensor decomposition is strongly related to principal component analysis (PCA) or singular value decomposition (SVD). While PCA and SVD are also applicable in this context, tensor decomposition is able to reduce the dimension while honoring the original data structure and correlations (e.g., spatial and temporal). These structures are often lost with data vectorization needed in PCA and SVD techniques. For example on an ensemble of 2D models, rather than vectorization, tensor decomposition operates by constructing a 3D tensor. The first two dimension correspond to the original spatial structure of the feature data and the third dimension to the realization number (i.e., the uncertainty dimension). More dimensions, such as time, may be included by taking snapshots of the feature property. According to Insuasty et al. (2015), tensor decomposition is able to compress large datasets while having minimal approximation and reconstruction error.

Consider a 2D reservoir model with 2D grid properties matrix **X** (size $I \times J$) at different time snapshot *K* stacked into 3D tensor **S** = [**X**₁, **X**₂, ..., **X**_K]. Tensor **S** can be decomposed as

Equation 5

$$\mathbf{S} = \sum_{i=1}^{l} \sum_{j=1}^{J} \sum_{k=1}^{K} \sigma_{ijk} (\boldsymbol{\varphi}_i \otimes \boldsymbol{\Psi}_j \otimes \boldsymbol{\chi}_k) ,$$

where $\xi_{ijk} = \varphi_i \otimes \Psi_j \otimes \chi_k$ is now rank-one tensors, φ_i , Ψ_j and χ_k are orthonormal basis functions (vectors) and σ_{ijk} is the elements of $I \times J \times K$ core tensor (i.e. the 3D analogy of a diagonal matrix). Equation 12 can be formulated as an optimization problem as

$$\begin{split} \min_{\boldsymbol{\varphi}_{1:l}, \boldsymbol{\Psi}_{1:J}, \boldsymbol{\chi}_{1:K}} \left\| \mathbf{S} - \hat{\mathbf{S}} \right\|_{F} , \\ s.t. \quad \boldsymbol{\varphi}_{i'}^{T} \boldsymbol{\varphi}_{i''} = \delta_{i'i''} , \boldsymbol{\Psi}_{j'}^{T} \boldsymbol{\Psi}_{j''} = \delta_{j'j''} , \boldsymbol{\chi}_{k''}^{T} \boldsymbol{\chi}_{k''} = \delta_{k'k''} , \end{split}$$

where $\delta_{w'w''}$ is the Dirac delta function. Insuasty et al. (2015) showed that this approach allows comparison of model realizations based on very rich datasets, such as the temporal evolution of the spatial distribution of pressures and saturations inside the reservoir. They are able to select a subset of realizations representative in terms of dynamic flow patterns and form reduced ensembles to perform robust production optimization more efficiently.

2.3.2 Multidimensional Scaling (MDS)

Multidimensional scaling (MDS) is a method that represents measurements of dissimilarity among pairs of objects as distances among points in a low-dimensional space. MDS has been widely used in engineering to map dissimilarity

matrix or distance matrix **D** into points in metric space. MDS is not a factorization technique like SVD but rather a method to rearrange objects in an efficient manner, with the goal to find a configuration that best approximates the observed distances. The points in this spatial representation are also arranged in such a way that their Euclidean distances (i.e., dissimilarity) corresponds to the projected distance of each points (Borg and Groenen, 1997). Scheidt and Caers (2009) introduced MDS in the reservoir simulation community and many successful applications are documented in Caers (2011).

MDS can arguably achieve the same results as dimensional reduction by projecting the feature data Θ into a lower dimensional dataset $\tilde{\Theta}$. Different from most other ordination methods (e.g., PCA and SVD), MDS is a numerical technique that iteratively computes a solution until a pre-defined tolerance has been reached. As a result, the solution of MDS depends on the initial randomized projection. The number of axes or dimensions are explicitly chosen prior to computation and data are fitted to chosen dimensions rather than truncating it. As a numerical optimization technique, MDS suffers from the possibility that the solution may be the local optima, but this can be reduced by repeating the process with random initialization seed.



Figure 2 Workflow of multidimensional scaling

With the distance matrix **D**, the realizations can be mapped using MDS into specified *p*-dimensional Euclidean space. *Figure 2* illustrates the workflow of how MDS functions. The stress function, *s* is a measure of fit on how well the data are mapped and is defined by

Equation 7

$$s = \sqrt{\frac{\sum (d_{ij} - \hat{d}_{ij})^2}{\sum \hat{d}_{ij}^2}},$$

where d_{ij} is the projected distance between *i* and *j* and \hat{d}_{ij} is the monotone transformation of δ_{ij} . Zero stress indicates a perfect fit. Kruskal (1964) suggested that low value of stress (i.e., < 5%) indicate an excellent fit between projected space with distance matrix. Understandably, increasing the dimension (i.e., degrees of freedom) in projected space would eventually reduce the stress value to 0%. However, that would defeat the purpose of dimensionality reduction that we want to take advantage of. Kruskal (1976) state that MDS can be complementary to clustering techniques.

2.4 Self-organizing Map (SOM)

Self-organizing map (SOM) is a type of artificial neural network (ANN) (Kohonen, 1990). ANN started when research in machine learning and artificial intelligence (AI) developed a technique inspired by biological neural networks (i.e., the brain). One of the main differences to regular ANN is that SOM relies on an unsupervised learning algorithm, which means that it does not require any a priori information to function and that it excels at establishing unknown relationships in dataset (Deboeck, 1998; Penn, 2005).

SOM networks typically have two layers of nodes, as shown in *Figure 3*: the input and the Kohonen layers. The input layer is fully connected to the two-dimensional Kohonen layer. During the training process, input data pass through the input layer's nodes. Assuming *M*-dimensional input vector $\mathbf{x} = [x_1, x_2, ..., x_M]^T$ and *N* Kohonen layer nodes ($N = n_X \times n_Y$), *M* input nodes are connected to each of the *N* nodes in Kohonen layer. A weight vector $\mathbf{w}_i = [w_{i1}, w_{i2}, ..., w_{iM}]$ is associated with N_i nodes ($i = 1, 2, ..., n_X \times n_Y$), where w_{ij} is the weight associated with i^{th} Kohonen layer node and j^{th} input layer node. SOM utilizes competitive learning where each node gradually becomes sensitive to different input data. The node *N* that best represents an arriving input **x** wins the competition and is allowed to learn better (i.e., increasing the weight). 'Specialization' occurs in the network when nodes specialize to represent different types of inputs. In most SOM, neighbors of the winning node are allowed to learn albeit at a lower rate, making representation of nodes become ordered.



input vector

Figure 3 SOM Kohonen network structure. Light pink denotes the winning node, pink denotes the immediate neighbors and purple denotes further neighbors. (from Kohonen Network - Background Information, 2012)

The competition function of Euclidean distance is defined by

$$f_{comp}(\mathbf{x}) = \arg\min\{\|\mathbf{x} - \mathbf{w}_i\|^2\}$$
.

The winning node, together with its neighbors, can better represent the input by modifying its weight. The amount of learning is dictated by the amplitude of neighborhood adaptation $A_i(i_{it})$ and defined by

Equation 9

Equation 8

$$\mathbf{w}_{i}(i_{it}+1) = \mathbf{w}_{i}(i_{it}) + \alpha(i_{it})A_{i}(i_{it})[\mathbf{w}_{i}(i_{it}) - \mathbf{x}(i_{it})] ,$$

where i_{it} is the iteration step index and $\alpha(i_{it})$ is the monotonically decreasing learning coefficient. The workflow of SOM is illustrated in *Figure 4*. At every iteration, the node with the minimum distance from the competition function is the winner and adjusts its weight to be closer to the value of input data. Each input data point is then assigned to the winning node. This process is repeated until specified iteration limit. In the end, the nodes are able to show the topological relations of the data and input data points that are in the same node are similar.

Kohonen (1996) claimed that SOM is a new and powerful tool used to visualize high dimensional data by converting complex and nonlinear relationships present in the data into simple geometrical relationships on a low-dimensional display. SOM is especially suitable for data surveys due to its prominent visualization properties and ability to obtain qualitative information. For this reason, SOM as a projection method has been extensively used in data exploratory research especially in pattern recognition (Kohonen et al., 1996). SOM has also been very successfully applied in seismic data interpretation, where similar seismic reflectors are grouped as indicators for lithology and pore content to assist interpretation (Klose, 2006). Kiang (2001) showed that using SOM in conjunction with clustering techniques (Vesanto and Alhoniemi, 2000; Cabanes and Bennani, 2010) has multiple advantages over other approaches.

While SOM has proven to be a good projection method, it can also be used for clustering. It has been shown that while for large number of nodes, SOM rearranges data in a way that is fundamentally topological in character, for small number of nodes, SOM behaves in a way that is similar to k-means clustering (Kaski, 1997). Kaski (1997) showed that SOM's cost function closely resembles the one minimized in K-means clustering. Kaski (1997) also stated that SOM can function as a conventional clustering algorithm if the amplitude of neighborhood adaptation is zero.



2.5 Robust Optimization

Van Essen et al. (2009) have presented robust optimization as a way to obtain the optimal control strategy that accounts for geological uncertainty by performing optimization over an ensemble; see also Chen et al. (2012) and Yasari et al. (2013). Although, in theory, with an ensemble we could derive a multitude of strategies that may improve the reservoir performance, in practice we can only apply one strategy for reservoir management and that is what motivates robust optimization. While an ensemble enables better inclusion of possible reservoir characteristics, optimizing production strategies for the ensemble becomes a more computationally demanding problem to solve as all realizations need to be considered. Besides controls for all the wells, the unique flow patterns of each realization also have to be examined. The main problem is that each realization has different reservoir characteristics which require different control strategies to maximize a given objective function, typically net present value (NPV) or cumulative volume of oil produced. Therefore, robust optimization requires the optimization of all realizations simultaneously in order to maximize the objective function. Given an ensemble, $\mathbf{M} = \{\mathbf{m}_1, \mathbf{m}_2, ..., \mathbf{m}_N\}$, where \mathbf{m}_i is the model realizations, the objective function of NPV is computed as

Equation 10

$$\mu_{NPV} = \frac{1}{N} \sum_{i}^{N} J_{i} \quad ,$$

where μ_{NPV} is the ensemble mean of objective function of each realization, J_i which is defined by

Equation 11

$$J_{i} = \int_{t=0}^{1} \frac{q_{o}(t, \mathbf{m}_{i})r_{o} - q_{wp}(t, \mathbf{m}_{i})r_{wp} - q_{wi}(t, \mathbf{m}_{i})r_{wi}}{(1+b)^{t/\tau}} dt$$

where *t* is time, *T* is the lifetime of reservoir, q_o is the oil production rate, q_{wp} is the water production rate, q_{wi} is the water injection rate, r_o is the oil price, r_{wp} is the water production cost, r_{wi} is the water injection cost, *b* is the discount rate and τ is discount time reference. The mathematical formulation of the optimization problem is

Equation 12

$$\max \mu_{NPV}(\mathbf{u}) ,$$

s.t. $\mathbf{g}(\mathbf{u}, \dot{\mathbf{x}}, \mathbf{x}, \mathbf{m}) = 0$
s.t. $\mathbf{c}(\mathbf{u}, \mathbf{x}) \le 0$,

where **g** is the generalized nonlinear vector-valued function of the reservoir simulator (system equation), **u** is the control vector to be optimized, **x** is the state vector, **m** is the model parameters vector and **c** are the constraints (e.g. on the inputs, outputs and state). The optimized strategy is a vector **u** containing the control settings usually for each well in a field over the lifetime of the reservoir. Typically strategy **u** is the model) or quarterly well head pressure, water injection rates and valve opening settings. Although there are *N* realizations in the ensemble, only one single optimal strategy **u** exists which we refer to as the robust optimal strategy which maximize the given objective function for the ensemble.

The optimization is formulated as finding **u** which maximizes J_{NPV} subject to **g** and **c**. Most of the time the relationship between inputs and outputs is nonlinear and the optimization is often nonconvex. Many numerical techniques are available for solving this type of optimization problem; this thesis employs adjoint-based method. For more information on the application of adjoint-based method see Brower and Jansen (2002), Van Essen et al. (2006), Zandvliet et al.(2007) and Jansen et al.(2008).

2.6 History Matching

Reservoir simulation models integrate knowledge of many domains in petroleum engineering such as geology, petrophysics, etc. Most if not all, reservoir models that are made for simulations have parameters that are uncertain. In order to reduce the uncertainty and obtain a set of reservoir models that reflect observed measurements, history matching (or data assimilation) can be utilized. History matching seeks to incorporate the presently observed information into existing numerical models. The intuition behind it is that, if what is simulated matches what is observed, then the model used is correct and more importantly reliable. In other words, history matching can also be defined as the act of adjusting parameters of the numerical models until it closely fits the observed data.

Since the reservoir parameters are changed to minimize the mismatch between historical production data and the simulated model response, it effectively makes history matching an inverse problem. This minimization is regarded as an optimization problem. The problem is compounded by the fact that the relationship between observed data and model parameters are highly complex and nonlinear. The model parameters may be porosity and permeability, fault transmissibility, initial saturation of phases and many more properties. The observed data on the other hand can be the production rates, bottom hole pressures, phase saturations or even 4D seismic data. By matching the simulated production data with real production data, we presume that we improve our reservoir models to better predict the response of the real reservoir and account for uncertainty.

One of the first history matching applications in petroleum engineering was done by Kruger (1961), where he manually calculated the areal permeability distribution of the reservoir. Jacquard and Jain (1965) then developed an initial framework for automated history matching and many further works have been done based on this framework. This technique is also known as computer-assisted history matching. An overview of the methods more recently used in petroleum engineering can be found in Oliver and Chen (2011). To simplify the inverse nature of this problem, assumptions are made that (1) All distributions are Gaussian, (2) Initial reservoir models are correct to some extent, (3) Measurements always contain Gaussian noise, and (4) Simulator numerical model is correct.

The Gaussian probability is defined by

$$f_x(x) = \frac{1}{\sigma\sqrt{2\pi}} \exp\left(-\frac{(x-\mu)^2}{2\sigma^2}\right)$$
,

where σ is the standard deviation and μ is the mean. As for multivariate Gaussian with *M*-dimensional vector **x**

Equation 13

Equation 14

$$f_{\mathbf{x}}(\mathbf{x}) = \frac{1}{\sqrt{(2\pi)^n |\mathbf{C}_{\mathbf{x}}|}} \exp\left(-\frac{1}{2} (\mathbf{x} - \mu)^T \mathbf{C}_{\mathbf{x}}^{-1} (\mathbf{x} - \mu)\right)$$

where $\mathbf{C}_{\mathbf{x}}$ is the covariance matrix and $|\mathbf{C}_{\mathbf{x}}|$ is its determinant.

To account for uncertainty in reservoir simulation, Bayesian statistics is used by using probability as a measure for uncertainty. In general, observations, \mathbf{d}_{obs} which are more certain are used to improve a less certain reservoir model parameters, \mathbf{m} . In Bayesian framework, the unknown model parameters are treated as random variables with multivariate probability distributions. The conditional probability density function for model parameters given observation data is

Equation 15

$$p(\mathbf{m}|\mathbf{d}_{obs}) = \frac{p(\mathbf{d}_{obs}|\mathbf{m})p(\mathbf{m})}{p(\mathbf{d}_{obs})} .$$

Likewise for the multivariate model parameters and observation data, the probability distribution of the model parameters conditional to the data can be defined as

Equation 16

$$p(\mathbf{m}|\mathbf{d}_{obs}) \propto \exp^{\left\{-\frac{1}{2}(\mathbf{m} - \mathbf{m}^{\text{prior}})^{T} \mathbf{C}_{M}^{-1}(\mathbf{m} - \mathbf{m}^{\text{prior}}) - \frac{1}{2}(\mathbf{g}(\mathbf{m}) - \mathbf{d}_{obs})^{T} \mathbf{C}_{D}^{-1}(\mathbf{g}(\mathbf{m}) - \mathbf{d}_{obs})\right\}} ,$$

where **m** corresponds to the vector of model parameters, $\mathbf{g}(\mathbf{m})$ to the simulated data and \mathbf{d}_{obs} to the observed data. \mathbf{C}_{M} is the covariance matrix of model parameters and \mathbf{C}_{D} the covariance matrix of observed data. To maximize $p(\mathbf{m}|\mathbf{d}_{obs})$ we must minimize the term

Equation 17

$$I(\mathbf{m}) = \frac{1}{2} (\mathbf{m} - \mathbf{m}^{\text{prior}})^T \mathbf{C}_{\mathsf{M}}^{-1} (\mathbf{m} - \mathbf{m}^{\text{prior}}) + \frac{1}{2} (\mathbf{g}(\mathbf{m}) - \mathbf{d}_{obs})^T \mathbf{C}_{D}^{-1} (\mathbf{g}(\mathbf{m}) - \mathbf{d}_{obs}) \quad .$$

The objective function is usually defined as a sum of weighted squared differences between observed and modeled data. The first term of the exponent is the regularization term which constrains to geological sensibility and reduces the ill-posedness of the problem in terms of model parameters. The regularization term acts as an anchor on the prior knowledge (e.g., the input from geologists) which in part limits the changes to the parameters and constrains the problem.

There are many available techniques for history matching and one of them are the gradient-based methods. They are generally very efficient but suffers from two limitations. Firstly, they tend to result in local optima rather than global optima. Secondly, the geological constraints are not preserved in standard gradient-based techniques due to geostatistical correlations between model parameters are not maintained during optimization. Sarma et al. (2006) proposed a method utilizing PCA to circumvent these two difficulties by efficiently reparameterizing the permeability field.

Figure 5 illustrates how history matching is carried out by adapting PCA reparameterization of realizations' parameters. According to them, this technique is more efficient than stochastic search procedures and is able to utilize adjoint computed for production optimization. The workflow of this method starts with generating the prior ensemble parameter, \mathbf{m}^{prior} and computing reduced dimension prior parameter, $\mathbf{\tilde{m}}^{prior}$ using PCA. For more information regarding reducing the dimension of parameter using PCA please refer to Sarma et al. (2006). History matching is performed on reduced dimension prior parameter with observed data, \mathbf{d}_{obs} . Various methods may be implemented to generate sensitivity coefficients at this stage. In this thesis we use the adjoint method. When we have the sensitivity coefficients, the parameters updates are performed using gradient-based line-search algorithms. After convergence, $\mathbf{\tilde{m}}^{post.}$ is then converted back into original dimension parameter, $\mathbf{m}^{post.}$ to obtain posterior ensemble, $\mathbf{M}^{post.}$.



Figure 5 Workflow of history matching

3. Methodology

3.1 Representative Model Selection

Representative model selection has been used in many domains of science; yet it has been somewhat neglected in the petroleum industry. A few studies that use representative models were highlighted in *Section 1.2.3 Representative Models*, but most of them are employed in UQ and only a few used in optimization processes. While each realization seemingly has unique parameters, some realizations, in fact, behave similarly in terms of flow characteristics under a given field development configuration. Thus, grouping similar realizations together and using only one to represent each group seem to be tenable.

Insuasty et al. (2015) presented the use of representative models in robust optimization with reduction from 1,000 to 50 realizations (i.e., 5% of the full ensemble). With highly complex numerical models, 50 realizations may still have prohibitively high computational cost. Therefore, this thesis seeks to further reduce the number of representative realizations to the bare minimum without sacrificing too much accuracy in the results. The deliberation behind this is that order of magnitude and percentage of reduction are not a good measure of bare minimum required realizations. For instance, a reduction up to 10% for ensembles of 10 and 100 realizations is not the same: while 1 representative realization is clearly insufficient to represent the uncertainty characterized by an ensemble of 10, 10 representative realizations may be adequate to represent an ensemble of 100. *Figure 6* depicts the workflow used throughout the thesis to select representative models. For simplicity we refer to the process as "Repr. Select" from here on. Note that the terms representative realization and representative model mean the same here and are used interchangeably in the remaining of the text. Also note that, the representative ensemble is always a subset of the full ensemble.



Figure 6 Workflow for selection of representative models (Repr. Select)

We can split the Repr. Select process into four important steps, which are detailed in the following *subsections 3.1.1 Feature Selection - subsection 3.1.4 Weighting.*

3.1.1 Feature Selection

Feature selection is the first and perhaps the most important step in representative model selection. As the famous "Garbage In, Garbage Out" principle in computer science and related engineering domain, the inputs for further

operations are often the most important and should be meticulously chosen. Two main categories of features have been identified, namely static (e.g., permeability) and dynamic (e.g., oil saturation snapshots) properties. Static features do not require any simulation as the initial parameters of the reservoir models can be accessed directly. Dynamic features on the other hand require simulation and are unique to the well configuration and production strategy. Production strategy is kept constant to generate the dynamic features.

Dynamic flow features are generally preferred as they offer better distinction on relevant flow patterns rather than using all the differences in parameters. Because, not all permeability grids have the same influence in the model response, grids on the edge of the field are debatably not as important as grids in the middle of the field; dynamic feature like oil saturation snapshots are able to easily distinguish flow barriers and therefore able to more effectively differentiate between models. Single or many types of parameters can be used. However, the use of multiple types of parameters combined should be considered with care as different parameters may have different importance and different scales. Thus, weighting and normalization should be imposed on features that have multiple types of parameters.

3.1.2 Projection Method

Tensor decomposition and MDS are used as the projection methods for clustering. Both methods are fundamentally different but both are effective projection methods. Tensor decomposition method used in this thesis utilizes high order singular value decomposition (HOSVD) from Insuasty et al. (2015).

The number of dimensions to retain in the projection is decided to be determined by an automated cut-off criterion. For tensor decomposition, the cut-off criterion is determined to be 95% of the cumulative energy content from decomposition in the projected space dimension. Whereas MDS utilize the stress value, the dimension p is increased appropriately to a value where the stress is less than 5%. MDS is an optimization algorithm and the process is repeated 300 times to reduce the chances of local optima. As a result, the lowest dimension may be found without sacrificing a good fit between data and projection.

3.1.3 Clustering

K-means algorithm is applied to cluster the projected data. K-mean++ initial seed is used and the clustering process is also repeated 100 times to decrease the probability of having a local optimum clustering. Only one representative model is selected from each clusters. Therefore, the number of representative models, N_{repr} desired have to be determined before clustering. The representative model selected is the realization closest to the centroid of each clusters. If only two points exist in a clusters, the realization is randomly chosen between the two. As such, any number of representative models can be used to form a representative ensemble. Note, $N_{repr} = n$ where $n = 1, 2, ..., N_{full}$, which N_{full} is the number of realizations in the full ensemble is used when referring to the selected representative ensemble.

3.1.4 Weighting

As a consequence of having only one realization from each clusters and the face that each clusters does not necessarily have the same amount of samples, weighting is needed to better represent the full ensemble. Here we determine the weight for each representative realization as the normalized number of realizations within the respective clusters. The purpose of weighting is to make representative ensemble statistically more similar to the full ensemble.

3.1.5 SOM

As presented in *Section 2.4 Self-organizing Map (SOM)*, SOM can be viewed as a projection method or a clustering technique. This thesis seeks to utilize the clustering ability of SOM to select representative realizations from an ensemble. This method is presented as an alternative to cluster data explained in *subsection 3.1.2 Projection Method and 3.1.3 Clustering* can therefore be used to replace both steps and follows the workflow illustrated in *Figure 7*.

The amount of nodes is determined by the number of representative realizations desired. Due to two-dimensional SOM used in this thesis, the number of representative realizations is constrained by the multiple of two numbers (e.g., 2×2 , 2×3 , or 3×3 grid yields 4, 6, and 9 nodes respectively). Note that the number of nodes does not necessarily

produce the exact number of representative realizations as some nodes may be empty (e.g., 3×3 grid may have one empty node and result in only 8 representative models selected).



Figure 7 Representative model selection using SOM

Figure 7 illustrates how SOM is used in this thesis consists of 2 stages. This method starts with the same feature selection as previously mentioned. Next, the grid of nodes is defined. The first SOM is then applied and sub-ensembles are created from each of the nodes. After that, the second SOM is performed on each sub-ensemble and the representative realization is then randomly selected within the maximum node of the second SOM. Like for the clustering method, the weights for representative models selected by SOM are determined according to the normalized number of realizations in each one of the nodes of the primary SOM. It is important to note that, when utilizing SOM as a clustering algorithm, the amplitude of neighborhood adaptation needs to be set to zero as we want each node to be specialized to a different type of input (i.e., the realizations characteristic).

3.2 Validation

Validation is needed to study the delineation of original ensemble from representative ensemble. In *Figure 8*, two main stages can be identified, the first one being right after the representative models selection (i.e., clustering) and the second one being after optimizations (i.e., comparing the results).

Stage 1 validation may be the most important and the hardest to quantify the performance. Optimization such as robust optimization and history matching are computationally intensive. Therefore, if a measure of performance can be obtained during stage 1, we can avoid the computationally expensive part of the workflow and reselect representative models if needed. One method of stage 1 validation is the cluster validity analysis (i.e., assessing the clustering quality). It is often based on specific criteria, but these criteria are usually very subjective (Jain et al., 2009). Cluster validation is done by applying statistical methods and testing the statistical significance. There are three types of cluster validation: the first being an external assessment that compares the recovered structure into a priori structure. The second is an internal examination to determine if the structure is intrinsically appropriate for the data and lastly, a relative test that compares two structures and measures their relative quality (Jain et al., 2009). Although applicable to determine the quality of clusters, cluster validation does not offer performance prediction in our workflow. For more details please refer to Jain and Dubes (1988) and Dubes (1993).

Another method may be implemented with statistical tests. However, the huge difference in sample sizes and the fact that representative ensembles are a subset of full ensemble make many statistical tests unsuitable. One statistical test deemed applicable is the two-sample Kolmogorov-Smirnov (KS) test which provides the *p*-value of significance for comparison of cumulative distribution function (CDF) between two populations of different sample sizes. More information on KS-test is available in *Appendix B*. Further deliberation on stage 1 validation can also be found in *Chapter 5. Discussion*.



Figure 8 Stages of validation

The second stage where the results of both representative ensemble and full ensemble are compared allows for quantitative validation more naturally. The full ensemble results serve as the reference. In robust optimization, the results of both representative ensemble and full ensemble have the same number of samples thus making visual comparison easier. By plotting results such as NPV CDF of full ensemble and representative ensemble together, a qualitative validation is possible. Calculating the mean of final NPV of the ensemble also allows for a quantitative validation because it is the objective function of robust optimization.

Two sample KS test is also applicable in stage 2 validation but having the results based on hypothesis testing is regarded as inadequate to quantify the performance therefore abandoned. Since robust optimization is performed using the mean NPV of the ensemble, comparing the mean NPV is an acceptable metric for performance. Ideally, full ensemble robust optimization should yield the highest ensemble's mean NPV and matching it allows for a quantitative performance measure.

In history matching, visual inspection is harder to be performed. Owing to the fact that history matching does not only take matching the truth production rates into account but also the spread of uncertainty the ensemble offers. A good representative ensemble should be able to cover most, if not all the uncertainty variation in observed measurements while showing improvement from prior to posterior towards observed measurements. Visual inspection is still possible where the representative ensembles must, to an extent, covers the full ensemble's observed measurements spread.

Normalized Sum of squared errors (SSE) can be measured for a quantitative performance metric. Normalization is required because the number of realizations in representative and full ensembles are different. Representative ensembles should have a normalized SSE that is close to, or less than the squared observation error standard deviation of the full ensemble's normalized SSE. Although this measurement is far from ideal, it does however provide a measure of performance.

3.3 Workflow of Representative Robust Optimization



Figure 9 Workflow of robust optimization on full ensemble and representative ensemble. (Left) Unoptimized reference (Middle) Representative robust optimization (Right) Full robust optimization reference.

Figure 9 depicts three workflows on how the results of representative ensemble robust optimization are acquired and compared. Only the middle workflow is needed for using representative ensemble robust optimization. The left of *Figure 9* is the workflow for unoptimized full ensemble result whereas the right workflow is for robust optimized full ensemble result. Both results are the guidelines to analyze the performance of representative robust optimization.

3.4 Workflow of Representative History Matching



Figure 10 Workflow of history matching on full ensemble and representative ensemble. (Left) Full ensemble history matching (Right) Representative history matching

Figure 10 illustrates how the performance of representative ensemble compared to full ensemble are obtained. The first step is to create the representative ensemble using the same method as in *Chapter 3.1 Representative Model Selection*. Next, history matching is performed on both full and representative ensembles and the posterior ensemble is simulated to generate data for comparison. For the application of representative ensemble, only the workflow on right of *Figure 10* is carried out.

4. Examples and Results

This section examines the performance of representative ensemble against full ensemble, by applying the methodology described in *Chapter 3. Methodology* to a few examples. *Section 4.1 Case Studies* describes the synthetic models used to scrutinize the performance of selected representative ensemble. *Section 4.2 Robust Optimization Results* and *section 4.3 History Matching Results* present and analyze results obtained from the optimization experiments.

4.1 Case Studies

4.1.1 2D Model

The 2D synthetic models used for this work consist of 50 ensembles of 50 realizations each. The 2D models are simple two-dimensional reservoir models with an inverted five-spot well configuration, where one injector is in the middle with four producers at every corner of the field. The model has 21×21 grids of 700×700 m with a heterogeneous permeability and porosity fields. The parameters of the reservoir model are shown in *Table 1* and the configuration of the field is depicted in *Figure 11*.



Figure 11: 2D inverted five-spot configuration

Tuble 1. Reservoir Troperties for 2D mouel				
Reservoir Parameters			Economic Parameters	
Oil Density, ρ_o	800 k	g/m³	Oil Price	80 \$/bbl
Water Density, ρ_w	1,000 k	g/m³	Water Production cost	5 \$/bbl
Oil Viscosity, µ _o	0.5	сP	Water Injection cost	5 \$/bbl
Water Viscosity, μ_w	1	сP	Discount factor	0.15 [-]
Residual Oil Saturation, S _{or}	0.2	[-]		
Connate Water Saturation, S_{wc}	0.2	[-]		
Relative Permeability Oil Endpoint, $k_{ro,e}$	0.9	[-]		
Relative Permeability Water, Endpoint, $k_{rw,e}$	0.6	[-]		
Oil Corey Exponent, n _o	2	[-]		
Water Corey Exponent, n_w	2	[-]		
Initial Water Saturation, S_{wi}	0.2	[-]		
Initial Reservoir Pressure, $oldsymbol{p}_0$	300	bar		

Table 1: Reservoir Properties for 2D Model

The optimization was run for 1,500 days with well control updates every 150 days by changing bottom hole pressure with a range of 200-300 bar for producers and 300-500 bar for injectors. The experiments for production optimization were performed with the open-source reservoir simulator Matlab Reservoir Simulation Toolbox (MRST) (Lie et al., 2012) developed by SINTEF and some modifications to allow for robust optimization. And the history matching runs were carried out with the Automatic Differentiation General Purpose Research Simulator (AD-GPRS) (Voskov and Zhou, 2012) developed at Stanford University.

Simsim Model



Figure 12 Permeability image of sixteen randomly chosen realizations from simsim model ensemble 50

Simsim Model is the same example as the 2D five-spot model described in Barros et al. (2016a). *Figure 12* illustrates the permeability distribution of the model where very distinct patches of high and low permeability can be observed.

Channel Model



Figure 13 Permeability image of sixteen randomly chosen realizations from channel model ensemble 50

Distinct channel features of high permeability may be observed in *Figure 13*. Channel Model is created using SNEISIM algorithm (Strebelle, 2002) that utilizes multi-point geostatistical simulation based on training image used for modeling layer 3 of the Stanford VI reservoir model (Castro et al., 2015). The only differences between simsim model and channel model are the permeability and porosity fields. The rest of the parameters remains the same.

4.1.2 3D Model

<u>Egg Model</u>

The Egg Model is a synthetic reservoir model created to serve as a benchmark for water flooding optimization, closedloop reservoir management and computer-assisted history matching (Jansen et al., 2014). The model consists of 100 realizations of channelized reservoir with 60×60×7 grid cells of which only 18,553 cells are active thus having the shape of an egg as illustrated in *Figure 14*.



Figure 14 (Left) Reservoir model displaying the position of the injectors (blue) and producers (red). (Right) Six randomly chosen realizations. (from Jansen et al., 2014)

The Egg Model is a two-phase (oil and water) model with no aquifer and no gas cap. The porosity of the model is homogeneous with heterogeneous permeability giving the channelized geological characteristic. The reservoir model's properties are stated in *Table 2*. One notable difference from the 2D Models is that Egg Model includes water and oil compressibility although still missing an important element which is faulting that has major effect on the flow pattern. Egg Model has very distinct channelized high permeability streaks that contribute to different flow characteristic in all realizations.

Tuble 2 Reservoir properties for Egg Model				
Reservoir Parameters			Economic Parameters	
Oil Density, <i>c</i> _o	900 kg	∫m³	Oil Price	126 \$/m ³
Water Density, c_w	1000 k	g∕m³	Water Production cost	19 \$/m ³
Oil Viscosity, μ_o	5	сP	Water Injection cost	6 \$/m ³
Water Viscosity, μ_w	1	сP	Discount factor	0 [-]
Residual Oil Saturation, S _{or}	0.1	[-]		
Connate Water Saturation, S_{wc}	0.2	[-]		
Relative Permeability Oil Endpoint, $k_{ro,e}$	0.8	[-]		
Relative Permeability Water Endpoint, $k_{rw,e}$	0.75	[-]		
Oil Corey Exponent, n_o	4	[-]		
Water Corey Exponent, n_w	3	[-]		
Initial Reservoir Pressure, $m{p_0}$	400	bar		
Initial Water Saturation, S_{wi}	0.1	[-]		
Oil Compressibility, <i>c</i> _o	10-10	Pa-1		
Water Compressibility, c_w	10-10	Pa ⁻¹		
Porosity, φ	0.2	[-]		

Table 2 Reservoir properties for Egg Model

The robust optimization of production strategy consists of ten control intervals on water injection rates of all eight injector wells (blue) over a period of 3,600 days with control updates every 360 days. The bottom hole pressure of producers are kept constant at 395 bar. Further details on the Egg Model can be found at Jansen et al. (2014). For the Egg Model, both robust optimization and history matching experiments were carried out with the use of AD-GPRS only to obtain the required gradients.

4.2 Robust Optimization Results

Representative ensemble is first tested on robust optimization workflow to investigate the performance and at the same time explore critical criteria such as type of input feature and minimum amount of representative models required. The types of input features considered in this section are the permeability, the NPV time series (NPV taken at every control interval), and oil saturation snapshots (taken at every control interval). In order to objectively distinguish the realizations, the field production strategy is kept constant.

The other important criteria to be determined is the minimum number of representative models required to achieve a good performance. 3, 5 and 10 representative models from the full ensemble have been used to ascertain the necessary amount. The results are presented in sections of different models used.

The results are presented as final NPV CDF curve. The grey line indicates initial NPV before robust optimization, black line is the full robust optimized result and acts as the reference to be compared with. Dashed colored lines are the NPV CDF of representative ensemble after robust optimization utilizing 3, 5 and 10 representative realizations respectively. Visual inspection is used to evaluate the performance of each representative ensembles where high conformation of representative ensemble curve to full ensemble curve implies a good performance.

4.2.1 2D Model

Robust Optimization is performed on 10 sets of both 2D models. The results for the first four ensembles are shown in the following sections. The rest of the results is available in *Appendix A*. Figure 26 depicts the effect of history matching on the 2D models.

<u>Simsim Model</u>

Using permeability as feature input shows an acceptable performance on representative ensembles. Tensor decomposition has a noticeable poorer performance compared to MDS in this example $N_{repr} = 3$ (i.e., 5% of the full ensemble) seems to be insufficient to have a good robust optimization in most cases shown.



Figure 15 Simsim model NPV CDF of 4 ensembles using permeability as feature. (Left) MDS (Right) Tensor decomposition



The choice of NPV time series as selection feature results in a better performance of selected representative ensembles. In both cases MDS and tensor decomposition performed well in shown results. Once again, using $N_{repr} = 3$ seems inadequate for robust optimization.



Figure 17 Simsim model NPV CDF of 4 ensembles using oil saturation snapshots as feature. (Left) MDS (Right) Tensor decomposition

The NPV CDF results of representative ensembles selected using oil saturation snapshots in *Figure 17* shows good performance on most cases with the exception of $N_{repr} = 3$ from ensemble-3 using tensor decomposition having a very poor performance. The comparison of the results in *Figure 15, Figure 16 and Figure 17* gives us a notion that using 3 realizations is not sufficient to be used as a representative ensemble.



Figure 18 Simsim model NPV CDF of 4 ensembles using (Left) random selection (Right) oil saturation snapshots with SOM



Figure 18 (left) shows the performance of randomly selected representative ensembles, making clear that random selection is not a good method of choosing representative realizations. However, the results give an important insight to this example used. Firstly, $N_{repr} = 10$ (i.e., 20% of the full ensemble) seems to be enough representing the full ensemble. Secondly, some ensembles are easier to represent (ensemble-2) than others. Thus, the quantification of performance should have at least more than one example ensemble. *Figure 18 (right)* is the result from using SOM and

oil saturation snapshots for selecting representative models. The performance appears to be on par with the clustering methods.

Figure 19 gives the overall performance on the results presented in in *Figure 15, Figure 16 and Figure 17*. The mean NPV of ensemble is a good measure for performance since it is also the robust optimization's objective function. The black bar designates optimized mean NPV for the full ensemble and it is rather clear that $N_{repr} = 3$ is insufficient as a good representative ensemble. We also notice that NPV time series and oil saturation snapshots are better features for representative selection.





Figure 20 Channel model NPV CDF of 4 ensembles using permeability as feature. (Left) MDS (Right) Tensor decomposition



Figure 21 Channel model NPV CDF of 4 ensembles using NPV time series as feature. (Left) MDS (Right) Tensor decomposition

The results of channel model illustrated in *Figure 20* using permeability as feature shows similar performance for MDS and tensor decomposition. All representative ensembles performed very well in all the ensembles shown. However, upon further analysis on more results in *Appendix A* revealed that $N_{repr} = 3$ still is insufficient and performed poorly

in a few cases. Compared to *Figure 20*, results in *Figure 21* use the NPV time series as selection feature and noticeable poorer performance can be observed with both MDS and tensor decomposition, especially for ensemble-4. This may be due to realizations that have similar NPV time curves that are similar despite having very distinctive flow patterns.



Figure 22 Channel model NPV CDF of 4 ensembles using oil saturation snapshots as feature. (Left) MDS (Right) Tensor decomposition



Figure 23 Channel model NPV CDF of 4 ensembles using (Left) random selection (Right) oil saturation snapshots with SOM

Figure 22 shows the results obtained using oil saturation snapshots as the feature. All representative ensembles show good performance except for $N_{repr} = 3$ which showed slightly poorer performance than the rest. From *Figure 20*, *Figure 21* and *Figure 22*, the performances of MDS and tensor decomposition are hardly different. Further analysis on more results available in *Appendix A* showed the same, where performance differences between MDS and tensor decomposition are indistinguishable.

Random selection in channel model shown in *Figure 23 (left)* appears to indicate that channel model is a simpler model to represent compared to simsim model. Even though showing good performance on randomly selected representative ensembles, MDS and tensor decomposition selected representative ensembles still performed

noticeably better. *Figure 23 (right)* shows that SOM is able to be used as an alternative to the presented selection methods. However results with SOM performs noticeably poorer than the main representative selection method.



Figure 24 provides the comparison of mean NPV in *Figure 20, Figure 21* and *Figure 22*. Once again, $N_{repr} = 3$ have lower performance than $N_{repr} = 5$ and $N_{repr} = 10$ in almost all ensembles. The performances of MDS and tensor decomposition are very similar for the channel model. However, the feature using NPV time series showed a poorer performance, especially for ensemble-4. From the analysis of the results, representative ensembles are thought to be applicable in robust optimization at least in a simple 2D model.

4.2.3 3D Model

<u>Egg Model</u>

The same workflow of using representative workflow is implemented in Egg Model robust optimization. *Figure 25* indicates the final NPV CDF of Egg Model using ensembles of 5, 10 and 20 representative models for robust optimization. The results using representative ensembles are very promising with a few performing even better than the full ensemble. This may be due to a discovered problem in robust optimization carried out using AD-GPRS where, during robust optimization, a few realizations yield gradients that are orders of magnitude higher than the other realizations. Thus, resulting in an inferior robust optimization that is not optimized evenly for all realizations. This is reflected in the full(100) ensemble NPV curve which is lower than full(90) NPV curve where 10 identified realizations with unusually high gradient hae been removed from the ensemble. The problem remains unresolved at the time of writing.

The results showed that having $N_{repr} = 5$ is sufficient for a good representative ensemble. In representative selection using oil saturation snapshots for both MDS and tensor decomposition, we observe poor performing representative ensemble of 10 and 20 which may be due to the selection of realizations that have unusually high gradient. The results

also showed that permeability may be better selection feature in this case. However this may be due to the fact that the Egg model does not have other parameters that will dictates flow pattern.



4.3 History Matching Results

History matching is performed using field production rates of oil and water, the rates is studied to analyze the performance of the representative ensemble. AD-GPRS is used to perform history matching and because AD-GPRS only provide gradients for permeability, the history matching is done by updating the permeability parameters for the reservoir models.

4.3.1 2D Model



Figure 26 Permeability results of history matching using MDS. (Top) Simsim Model (Bottom) Channel Model

The porosity of 2D model is modified to be homogeneous as history matching is carried out on permeability only. History matching is performed on the measured water and oil field production for 2D models. The observation error is expressed in terms of standard deviation, $\sigma_{prod_rates} = 5 \text{ m}^3/\text{day}$ of the production rates measurement. Figure 26 depicts the effect of changes in permeability field in history matching using full ensemble and representative ensembles. For simsim model, representative ensembles has noticeably bigger changes compared to full ensemble.

Simsim Model

One of the simplest and illustrative ways to determine the performance of representative ensemble to full ensemble is by visual inspection on field's production profile of oil and water. The history matching time is arbitrarily selected at 1,500 days to standardize the comparison. Note that history matching is implemented strictly on the production rates measured at 1,500 days (yellow dots) and not the historical data until 1,500 days.



Figure 27 Simsim Model field production data of representative ensemble using all oil saturation snapshots at time 1,500 days. Red is oil and blue is water. (Left) The priori (Right) The posterior

At first glance, it is clear that the representative ensemble performs well in representing the full ensemble. The dotted line (blurred) depicts the full ensemble and the dashed line portrays the representative ensemble whereas the solid line is the truth realization. By comparing the dotted and dashed line, we noticed the representative ensemble is able to cover the spread of uncertainty in full ensemble. Although comparatively less scattered, the posterior of representative ensemble still has an acceptable spread.

Figure 27 also allows the study on the performance of using two different methods of projection mainly MDS and tensor decomposition, and the number of representative realizations needed. By visual inspection, $N_{repr} = 10$ have better representation of full ensemble's uncertainty over $N_{repr} = 5$. The comparison of MDS and tensor decomposition shows a little advantage in using MDS as the selected realizations have a better representation of the full ensemble. Although it is inconclusive from *Figure* 27, further analysis on more data available in *Appendix A* has showed similar interpretation that MDS performs slightly better.

Figure 28 is to study the effect of history matching at various times. Generally we notice again the representative ensemble performs satisfactory at all control interval time. Note that the representative selection is done considering all the snapshots of oil saturation until end of simulation time, which means that the same representative realizations have been selected for all the plots. One noticeable characteristic concerning the history matching time is a considerably poorer match at t = 450 days and t = 600 days, which is the onset of water breakthrough. The reason behind this may be due to the fact that history matching is done using the field production data rather than individual well production data. The contrast is that at the onset of water breakthrough in field production data resulted in more solutions as water breakthrough in specific well is not known as we are history matching on field production data.



Figure 28 Simsim Model field production data of posterior of 10 representative realizations using MDS and all oil saturation snapshots at various history matching time. Red is oil and blue is water



Figure 29 Simsim Model field production data of posterior of 10 representative realizations using MDS and oil saturation snapshots up to specified history matching time. Red is oil and blue is water

Figure 29 differs from *Figure 28*: here the selection of representative realizations in *Figure 29* is based on oil saturation snapshots up until the history matching time making the feature data sparser and at earlier t < 1,200 days the feature data may not be sufficient for a good representative model selection. The first apparent observation by comparing *Figure 28 and Figure 29* is that history matching at earlier times performs worse in *Figure 29*. This is most likely due to the fact that the saturation snapshots taken at earlier times have insufficient data to differentiate between realizations effectively. As the time to history match increases, the representative model selection shows improvement and even shares the same realizations as in *Figure 28*.



Figure 30 Simsim Model NPV comparison of 10 representative realizations using MDS. Repr prior all and repr post all are selected using all oil saturation snapshots whereas repr prior and repr post are selected using oil saturation snapshots up to the history matching time. (Left) The prior NPV CDF plot. (Right) The posterior NPV CDF plot.

Another way to determine performance is to analyze the NPV of the ensemble. In essence the representative ensemble's NPV distribution should favorably have a good representation of full ensemble's NPV distribution. For this comparison we study the prior and posterior NPV CDF plot of the ensembles in *Figure 30*. Note that no optimization is carried out on the ensemble thus shifting of the NPV CDF curves does not reflect performance. Also for that reason, the full ensemble's curve will be the reference and a closer match of representative ensemble on full ensemble's curve will be interpreted as good representation.

In *Figure 30(left)*, prior representative all have a good representation of the full ensemble's NPV CDF plot however repr prior shows a weaker representation. In *Figure 30(right)*, both representative ensembles have very similar curves to each other. However, they do not match to the curve of full ensemble. Almost all posterior NPV CDF shows smaller spread compared to the prior except for history matching done in the later time of which the spread of NPV grows. One possibility is that due to having higher water production rate and much lower oil production rate at the later part of the history matching time, which will greatly affect the NPV as it relies heavily on oil price and water injection/production cost.

<u>Channel Model</u>

In channel model, similar results as presented previously are shown in this section. *Figure 31* is similar to *Figure 27* where the comparison of prior and posterior production profiles is done. Again, the representative ensembles provide acceptable representation of the full ensemble's uncertainty in both cases. The posterior in *Figure 31(right)* shows noticeable reduction in spread of uncertainty especially on the water production rate. However the reduction in

spread is marginal. The difference between the performance of MDS and tensor decomposition is much less noticeable compared to *Figure 27*. However, $N_{repr} = 10$ seems to have a better representation than $N_{repr} = 5$.



Figure 31 Channel Model field production data of representative ensemble using all oil saturation snapshots at time 1,500 days. Red is oil and blue is water. (Left) The prior (Right) The posterior



Figure 32 Channel Model field production data of posterior of 10 representative realizations using MDS and all oil saturation snapshots at various history matching time. Red is oil and blue is water

Figure 32 shows that for channel model, choosing the history matching at different time does not have any real impact on the history matched results. It could be due to the fact that the truth is an outlier realization. Meaning that more

significant model updates are needed and the regularization term in history matching is preventing larger changes to preserve initial model channel location.



Figure 33 Channel Model field production data of posterior of 10 representative realizations using MDS and oil saturation snapshots up to specified history matching time. Red is oil and blue is water



Figure 34 Channel Model NPV comparison of 10 representative realizations using MDS. Repr prior all and repr post all are selected using all oil saturation snapshots whereas repr prior and repr post are selected using oil saturation snapshots up to the history matching time. (Left) The prior NPV CDF plot. (Right) The posterior NPV CDF plot.

Figure 33 shows the selection based on oil saturation of snapshot up to the history matching time. A comparison between *Figure 32 and Figure 33* and suggests that selecting representative realizations based on all oil saturation snapshots is more advantageous over only based on oil saturation snapshot up to certain history matching time. *Figure 34* compares the representative realizations selection based on NPV CDF plots. In terms of NPV, both selection based on different history matching time and all saturation snapshots shows similar result and they conform well to the full ensemble NPV CDF curve.

In all, the representative ensembles prove to be sufficient to replace full ensemble in history matching without compromising uncertainty quantification too much. The prior and posterior of the representative ensemble still cover adequate spread in uncertainty when compared to full ensemble's uncertainty spread although understandably less.

4.3.2 3D Model

<u>Eaa Model</u>

History matching of Egg Model is performed using the field oil and water production rates and each injectors' pressure. The observation error is expressed in terms of production rates standard deviation, $\sigma_{prod_rates} = 5 \text{ m}^3/\text{day}$ and injectors' pressure standard deviation, $\sigma_{inj_pressure} = 10$ bar of the measurement. For Egg Model, the time of history matching is arbitrarily chosen to be at 1,800 days.



Figure 35 Examples of history matched layer 4 of Egg Model permeability field using (Top) MDS (Bottom) Tensor decomposition.

Figure 35 shows the effect of history matching on the permeability field. Similar to what was observed in 2D model, the changes in permeability field of representative ensemble is bigger than of full ensemble. The results comparison of Egg Model are based on the field water and oil production data similar to the results of 2D models. Since the history matching of the Egg Model is carried out using the injectors' pressure as well, looking at the pressure profile of each injectors can shed some insight on the performance of the representative ensembles compared to the full ensemble.



Figure 36 Egg Model water (blue) and oil (red) production rates. Representative ensembles selected using MDS.



Figure 37 Egg Model injector rates of representative ensembles using MDS as projection method. (Left) Prior (Right) Posterior

The first comparison with full ensemble is the field water and oil production illustrated in *Figure 36 and injectors pressure Figure 37* where MDS is used to choose the representative ensembles. Both figures show similar results where a slight reduction in the spread of both rates are noticeable in the posterior results. In *Figure 36*, a few realizations of the full ensemble have showed larger spread in the posterior result which may due to improper changes in permeability field during history matching. During the history matching simulation, a few realizations have been noticed to require far greater time for simulation which suggested a poor update on the realizations' permeability

field. This problem was not encountered in the representative ensembles simulations. *Figure 37* depicts the injectors' pressure of representative ensembles against the full ensemble. The representative ensembles are able to have similar outline as the full ensemble although understandably less. $N_{repr} = 10$ showed a rather deficient coverage of the uncertainty and is considered rather insufficient to represent the full ensemble.



Figure 38 Egg Model water (blue) and oil (red) production rates. Representative ensembles selected using tensor decomposition.



Figure 39 Egg Model injector rates of representative ensembles using tensor decomposition as projection method. (Left) Prior (Right) Posterior

Similar to *Figure 36 and Figure 37, Figure 38 and Figure 39* compare the performance of representative ensembles using tensor decomposition. *Figure 38* showed a good representation on the field production rates. Tensor decomposition provides a larger spread in pressure profile when comparing *Figure 39 to Figure 37* of using MDS. In spite of that, it is hard to distinguish the performance difference between the two projection methods with visual inspection. However, it is rather clear that $N_{repr} = 10$ is not able to have a good representation from the injectors' pressure profile. This observation is noticeable in both projection methods used where, in a few cases, have highly deviated pressure profile. From the figures, the representative ensembles provide a relatively good matches to the curve and spread of full ensemble. Although, it is clear that $N_{repr} = 20$ represents the uncertainty spread much better than using $N_{repr} = 10$.



Figure 40 Normalized sum of square error at 1,800 days of field oil and water production rates to the truth.

Figure 40 depicts the normalized sum of squared error (SSE) of the water and oil production rates compared at 1,800 days. SSE provides an indication to the spread of the uncertainty to some extent. In our application case, closer match to full ensemble's normalized SSE value can be viewed as the reference since uncertainty covered is as important as accuracy. Both water and oil production rates' SSE are very similar, this may be due to the same weightage in history matching. A large increase in the posterior SSE of full ensemble's water production rates is most probably due to the poorly updated realizations as mentioned previously. Generally, there should be reduction in the normalized posterior SSE values which are observed in representative ensembles using MDS but not tensor decomposition. With considering $\sigma_{prod_rates} = 5 \text{ m}^3/\text{day}$ meaning the SSE may have an acceptable deviation of 25 m³/day, thus all the representative ensembles are within an acceptable range compared to the full ensemble.



Figure 41 Egg Model history matching prior and posterior final NPV CDF using full ensemble and representative ensemble selected using MDS and tensor decomposition.

Figure 41 depicts the final NPV CDF of prior and posterior in order to compare both MDS and tensor decomposition selected representative ensembles. The truth realization has a final NPV of 16 million and all the posterior NPV CDF have a noticeable shift towards the truth's NPV which indicate history matching is performing effectively. In terms of number of representative realizations, both $N_{repr} = 10$ and $N_{repr} = 20$ have similar curve thus does not offer clear performance difference among them. It is clear from *Figure 41* that tensor decomposition representative ensembles have better match with the full ensemble's NPV CDF compared to MDS selected representative ensembles especially in the posterior results.

5. Discussion

The discussion is separated into a few category of relevance. Firstly, the summary of the methodology, findings and possible improvements of this thesis are examined.

- 1. Features have been identified as a critical component in the selection of representative models. Therefore, various features were compared on simple 2D models and we have chosen to focus on permeability, NPV time series and oil saturation snapshots. From the results, permeability and oil saturation snapshots were found to be better. The reason NPV time series was not performing up to par is thought to be due to NPV relying on economic factors thus limiting the ability of differentiating between realizations. Although permeability was a good feature in our case, it is not recommended for models that contain other properties that can influence the flow characteristic of the models.
- 2. Considering projection or dimensionality reduction is important to the methodology, where two principally different techniques were implemented. However, we have found little to no noticeable difference in the effectiveness of tensor decomposition compared to MDS as a projection technique for clustering. In tensor decomposition, the cutoff criteria which used 95% of the cumulative energy content for projection dimension remained an unstudied problem. Nevertheless, case models used are admittedly rather simple where more complex model may show different results and may give better insight on the performance of MDS compared to tensor decomposition.
- 3. Although features are very important for the effectiveness of representative selection, MDS projection relies heavily on the pairwise distance between realizations. Therefore, the type of distance measures may have notable impact on the projected data. This thesis only applied Euclidean distance in using MDS and the effect of distance measure may be pursued in future works.
- 4. K-means clustering partitions data into groups with unique centroids which is the ideal representative point of each groups. The selected representative models have to be compromised to the one closest to each centroids thus introducing imperfection to the representative ensembles. This may explain why some representative ensemble has substandard performance in a few cases especially evident with $N_{repr} = 3$. Although not tested, the distance of selected realizations to the centroid may be a criteria for reselection of representative realizations.
- 5. SOM was introduced where the clustering is solely based on grouping by competition, consequently similar realizations will be grouped together. However, the clustering technique with SOM presented here is an exploratory attempt to provide an alternative to K-means clustering. Further studies are required for many parameters such as number of training iteration, stopping criteria, the competition function and the effect of amplitude of neighborhood to determine their appropriate value. In spite of that, preliminary use of SOM resulted in comparable performance with K-means clustering with using MDS or tensor decomposition as projection method.
- 6. Since the goal was to discover the bare minimum representative models needed to represent an ensemble, the amount of realizations has to be drastically reduced. Thus, in order to preserve the full ensemble statistically, weighting is needed for each representative models. We have investigated the importance of weight from simulation runs and concluded that weighting is essential and thus included in the methodology. Many types of weighting scheme may be applied. Therefore, there are rooms to improve the representative ensembles' performance by identifying a more suitable weighting scheme such as one that takes the distance of realizations to centroid into account.

Next, the results and recommendation for future works that may improve this methodology are discussed.

1. The goal of this thesis was to accelerate robust optimization and history matching by using far less realizations than the original ensemble. Based on the results obtained for our case studies, representative ensembles can undoubtedly be used to accelerate robust optimization workflows. The minimum representative realizations seems to be around 5 to 10 reservoir models, although more studies should be carried out to ascertain this statement.

- 2. The performance of using representative ensembles in history matching was not as well defined as in robust optimization although it was still able to capture most of the quantified uncertainties. Compared to full ensemble, the representative ensembles still cover big portion of the uncertainty spread. In 2D models, $N_{repr} = 10$ outperforms $N_{repr} = 5$ whereas in Egg Model, $N_{repr} = 20$ performs better than $N_{repr} = 10$. This may indicate that the minimum number of realizations required for history matching is related to the percentage representation of the full ensemble contrary to robust optimization. Understandably, an ensemble should provide UQ for the reservoir and naturally having less realizations in representative ensemble reduces the ability for UQ. It is important to note that each representative realizations has its own weightage and visual inspection on the spread was unable to take that into account.
- 3. The results have shown that proposed methodology can be implemented in robust optimization and history matching. Therefore, using this method should be applicable to CLRM without the need of extensive modifications to the workflow, provided that adjoint-based optimization is available (Barros et al., 2016b). However, there is more work to be done if we wish to apply this methodology in ensemble-based optimization.
- 4. The speedup obtained by using representative ensembles can be further increased with the incorporation of ROM techniques such as multiscale method. If proved attainable, this is a step in the right direction to realize the VOI workflow that has been unfeasible in terms of computational cost until now.

The validation of the results has faced many challenges, those challenges are presented in the following section.

- 1. Although not included in this report, various analyses were carried out to assure good selection of realizations before further optimizations, but the lack of samples made most statistical tests inadequate. Stage 1 validation (Figure 8) of the selected representative models remains a huge challenge. The main reason is that representative ensemble is a subset of the full ensemble and have much less realizations than the full ensemble. Therefore, classical statistical tests such as variance explained are not suitable as a measure because the results in variance explained are always lower than the full ensemble. The mean of ensembles has been tested but it does not reflect the quality of selected representative models when compared with stage 2 results. The covariance matrices of representative ensembles are of much lower rank compared to full ensemble. Thus, covariance suffers from the same problem as variance as well as the entropy of SVD of representative when compared to full ensemble. KS-test performed in stage 1 validation also proved ineffective. Although applicable to provide a quantitative measurement of fit to NPV of full ensemble, robust optimization changes the final NPV where conformity to the initial unoptimized NPV is thus meaningless.
- 2. Clustering validation provides a measure on how well the clustering algorithm performs based on criterion function. However, it does not provide a measure on how well the original dataset is clustered based on patterns of data. Jain and Dubes (1988) stated that "the validation of clustering structures is the most difficult and frustrating part of cluster analysis. Without a strong effort in this direction, cluster analysis will remain a black art accessible only to those true believers who have experience and great courage."

Lastly, we comment on the general understandings of the topic address in this thesis which are predominantly conjectural.

- 1. Ideally, there is only one strategy for robust optimization which maximizes the objective function while being constrained to all realizations. Intuitively, representative models can 'pull' the robust optimization towards that ideal strategy. However, ensembles that have less than 5 representative models proved to be difficult as having one more or one less realization affects the ability of the representative ensemble to 'gravitate' in the right direction. Example shown in Insuasty et al. (2015) which uses 50 realizations out of 1000 realizations should be possible to be reduced further to 10 or even less realizations and still provide a comparable optimal production strategy.
- 2. Currently, the comparison of history matching performance is carried out trying to match the spread of the full ensemble in terms of production rates or the NPV CDF curve or injectors' pressure (only for Egg Model). However, it is not clear whether this approach is considered better because, alternatively, it may also be desirable for the representative ensemble to have history matched results closer to the truth. Without a quantitative clarification on this matter, the performance of representative ensembles in history matching remains unquantifiable.

- 3. In the channel model example for history matching, all the posterior results are very similar. The reasoning was that the regularization term is preventing further changes in the permeability field. As the author's personal opinion, the geological accuracy in the model is somewhat overrated by having the conventional regularization term in history matching. If the observed data are clearly showing a very different measurements than expected, it may simply mean that the model is very different from the truth, thus, there is very little incentive in preserving geological accuracy in the first place. This thinking is also recounted by Kahrobaei (2016) using a simple 2D model. By removing the regularization term, his history matching procedure was able to reveal hidden geological features (e.g., barrier and high permeable streak) present in the true permeability field.
- 4. The method for selection for representative models presented in this thesis uses simulated feature data. In real reservoir management, measured data could be incorporated as a feature in choosing the representative models which are more likely to reflect the real reservoir. Therefore, a reselection of representative realizations may be beneficial at the first history matching interval.

6. Conclusion

The goal of the thesis was to accelerate workflow of robust optimization and history matching by using representative models to reduce the total computational cost. The thesis found that representative ensemble is highly effective in robust optimization. Having 5 or more representative models are sufficient in both 2D and 3D models tested to obtain a comparable optimal strategy to using a full ensemble. Nevertheless, the case studies we considered here are not as complex as real field models. Still, the results in robust optimization are very promising and should be tested in real field applications.

In history matching, presented results suggest that using 20 realizations (20% of the full ensemble) allows to quantify most of the uncertainties while achieving history matched models that are comparable to those obtained considering the full ensemble. Even so, using representative ensembles in history matching remains an open debate as there is a marginal decrease in the quality of UQ compared to using the full ensemble. Consequently, the choice of N_{repr} in history matching should be a compromise between the amount of acceleration required and the importance of UQ to the application purpose.

In terms of computational cost, our case studies showed that representative ensembles are capable of significantly reducing the number of simulations in robust optimization and history matching. The speedup is directly proportional to the percentage reduction in representative realizations used. When combining both robust optimization and history matching in CLRM, utilizing only 10% of full ensemble in one of the two processes is estimated to yield a speedup of two times assuming both have similar computational cost. Furthermore, by using 10% of full ensemble in both processes, an acceleration of one order of magnitude will be attainable in the CLRM workflow.

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Figure 43 Simsim model NPV CDF of 6 additional ensembles using NPV time series as feature. (Left) MDS (Right) Tensor decomposition



Figure 44 Simsim model NPV CDF of 6 additional ensembles using oil saturation snapshots as feature. (Left) MDS (Right) Tensor



Figure 45 Simsim model NPV CDF of 6 additional ensembles using (Left) Random selection (Right) oil saturation snapshots with SOM.



Figure 47 Channel model NPV CDF of 6 additional ensembles using NPV time series as feature. (Left) MDS (Right) Tensor decomposition



Figure 48 Channel model NPV CDF of 6 additional ensembles using oil saturation snapshots as feature. (Left) MDS (Right) Tensor decomposition



Figure 49 Channel model NPV CDF of 6 additional ensembles using (Left) Random selection (Right) oil saturation snapshots with SOM

Appendix B

The Kolmogorov–Smirnov test (KS test) is a nonparametric test of the equality of continuous, one-dimensional probability distributions that can be used to compare a sample with a reference probability distribution (one-sample KS test), or to compare two samples (two-sample KS test). Two-sample KS test is used to test whether two underlying distributions differ significantly. The KS-test has the advantage of making no assumption about the distribution of data. More information on KS test can be found at Chakravart, Laha, and Roy, (1967).