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The Development of a Hybrid Optimization Algorithm for the Evaluation and Optimization of the Asynchronous Pulse Unit

Eric Inclan
eincl001@fiu.edu

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THE DEVELOPMENT OF A HYBRID OPTIMIZATION ALGORITHM FOR THE
EVALUATION AND OPTIMIZATION OF THE ASYNCHRONOUS PULSE UNIT

A thesis submitted in partial fulfillment of
the requirements for the degree of
MASTER OF SCIENCE
in
MECHANICAL ENGINEERING
by
Eric Inclan

2014
To: Dean Amir Mirmiran  
College of Engineering and Computing

This thesis, written by Eric Inclan, and entitled The Development of a Hybrid Optimization Algorithm for the Evaluation and Optimization of the Asynchronous Pulse Unit, having been approved in respect to style and intellectual content, is referred to you for judgment.

We have read this thesis and recommend that it be approved.

_______________________________________  
Leonel Lagos

_______________________________________  
Igor Tsukanov

_______________________________________  
George S. Dulikravich, Major Professor

Date of Defense: September 12, 2014

The thesis of Eric Inclan is approved.

_______________________________________  
Dean Amir Mirmiran  
College of Engineering and Computing

_______________________________________  
Dean Lakshmi N. Reddi  
University Graduate School

Florida International University, 2014
DEDICATION

This thesis is dedicated to my wife, and my mother, as well as the Blake, Elviro, and
Babyak families.
ACKNOWLEDGMENTS

First, I would like to thank Dr. George S. Dulikravich, my major professor, for inspiring me, for challenging me to go farther than I have ever gone before, and for giving me so many opportunities to expand my learning. Also, thank you for guiding me toward optimization research, and for giving me the tools I needed to succeed.

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I would like to thank Dr. Marcelo Colaço for his help with OPTRAN. Muito obrigado!
ABSTRACT OF THE THESIS

THE DEVELOPMENT OF A HYBRID OPTIMIZATION ALGORITHM FOR THE EVALUATION AND OPTIMIZATION OF THE ASYNCHRONOUS PULSE UNIT

by

Eric Inclan

Florida International University, 2014

Miami, Florida

Professor George S. Dulikravich, Major Professor

The effectiveness of an optimization algorithm can be reduced to its ability to navigate an objective function’s topology. Hybrid optimization algorithms combine various optimization algorithms using a single meta-heuristic so that the hybrid algorithm is more robust, computationally efficient, and/or accurate than the individual algorithms it is made of. This thesis proposes a novel meta-heuristic that uses search vectors to select the constituent algorithm that is appropriate for a given objective function. The hybrid is shown to perform competitively against several existing hybrid and non-hybrid optimization algorithms over a set of three hundred test cases. This thesis also proposes a general framework for evaluating the effectiveness of hybrid optimization algorithms. Finally, this thesis presents an improved Method of Characteristics Code with novel boundary conditions, which better characterizes pipelines than previous codes. This code is coupled with the hybrid optimization algorithm in order to optimize the operation of real-world piston pumps.
TABLE OF CONTENTS

CHAPTER | PAGE
--- | ---
1. INTRODUCTION | 2
  1.1. Personal Contributions | 4

2. LITERATURE REVIEW | 6
  2.1. General Optimization Theory | 6
    2.1.1. Designs as Vectors | 6
    2.1.2. Objective Functions | 6
    2.1.3. Constraints | 7
    2.1.4. Performance Benchmarking | 8
  2.2. Passive Pure Optimization Algorithms | 9
    2.2.1. Local Optimization | 10
    2.2.2. Global Optimization | 10
  2.3. Hybrid Optimization Algorithms | 14
    2.3.1. No Free Lunch | 15
    2.3.2. Hybrid Algorithm Architectures | 16
    2.3.3. Switching Logic | 17

3. SEARCH VECTOR BASED HYBRID OPTIMIZATION ALGORITHM | 21
  3.1. A Global-Global Hybrid | 21
    3.1.1. Single Objective – Multiple Topologies | 21
    3.1.2. Search Vectors | 23
    3.1.3. Constituent Algorithm Search Vector | 29
  3.2. Search Vector Based Hybrid | 30
    3.2.1. Constituent Algorithm Selection Criteria | 32
    3.2.2. Hybrid Algorithm Architecture | 34
    3.2.3. Ramifications of Hybrid Algorithm Architecture | 39
  3.3. Proposed Multi-Objective Extension of Hybrid | 42
    3.3.1. Multi-Objective Search Vectors | 42

4. HYBRID ALGORITHM BENCHMARKING | 45
  4.1. Benchmark Case Set Up and Reporting | 46
    4.1.1. Establishing Statistical Significance | 46
    4.1.2. Impact of Floating Point Arithmetic | 48
    4.1.3. Computational Expense | 49
  4.2. Comparison of Blind Hybrid Optimization Algorithms | 50
    4.2.1. Comparison of Constituent Algorithm Selection Methods | 50
    4.2.2. Constituent Algorithm Selection Preference | 51
    4.2.3. Blind Hybrid Performance | 54
  4.3. Comparison of All Search Vector Hybrids | 57
    4.3.1. Revisiting Constituent Algorithm Selection Preference | 58
    4.3.2. Search Vector Hybrid Performance | 59
4.3.3. Impact of Problem Dimensionality ................................................................. 64
4.4. Comparison of Auto Hybrid to Non-Search Vector Hybrids ................................. 66
  4.4.1. Comparison of Execution Times ................................................................. 74
4.5. Discussion of Results ....................................................................................... 78
  4.5.1. Characterizing Hybrid Performance ......................................................... 78
  4.5.2. Search Vector Hybrid Characterization .................................................... 82
5. OPTIMIZATION OF THE ASYNCHRONOUS PULSE UNIT ................................. 86
  5.1. Pipeline Simulation Software ........................................................................ 87
    5.1.1. Existing Codes ......................................................................................... 87
    5.1.2. Updated Method of Characteristics ....................................................... 88
  5.2. Verification ..................................................................................................... 97
    5.2.1. Steady State Results ............................................................................. 97
    5.2.2. Transient State Results ...................................................................... 108
  5.3. Validation ...................................................................................................... 109
    5.3.1. Experimental Set Up and Boundary Conditions ................................... 113
    5.3.2. Simulation Results ............................................................................... 114
  5.4. Evaluating and Optimizing the APU .............................................................. 118
    5.4.1. Optimization Problem Set Up .............................................................. 118
    5.4.2. Optimization Results ........................................................................... 120
6. CONCLUSIONS .................................................................................................... 124
REFERENCES ........................................................................................................ 126
APPENDIX .............................................................................................................. 133
# LIST OF TABLES

<table>
<thead>
<tr>
<th>TABLE</th>
<th>PAGE</th>
</tr>
</thead>
<tbody>
<tr>
<td>Table 1: Number of Test Cases per Dimension Number</td>
<td>9</td>
</tr>
<tr>
<td>Table 2: Categories of Search Vector Based Hybrids</td>
<td>31</td>
</tr>
<tr>
<td>Table 3: Constituent Algorithm Selection Usage</td>
<td>34</td>
</tr>
<tr>
<td>Table 4: Legend of Acronyms</td>
<td>45</td>
</tr>
<tr>
<td>Table 5: Sample of Student’s t-test Results</td>
<td>46</td>
</tr>
<tr>
<td>Table 6: Total Number of Hypothesis Rejections</td>
<td>47</td>
</tr>
<tr>
<td>Table 7: Hypothesis Rejections – DE</td>
<td>47</td>
</tr>
<tr>
<td>Table 8: Sample of Student’s t-test Results</td>
<td>48</td>
</tr>
<tr>
<td>Table 9: Various Numerical Results by Operating System</td>
<td>48</td>
</tr>
<tr>
<td>Table 10: Probability of Violating Constraints</td>
<td>64</td>
</tr>
<tr>
<td>Table 11: Time Required for Convergence - OPTRAN, AUTO2, and PSO</td>
<td>76</td>
</tr>
<tr>
<td>Table 12: Probability of Correct Constituent Selection</td>
<td>84</td>
</tr>
<tr>
<td>Table 13: Errors for Steady State Pressure BC</td>
<td>100</td>
</tr>
<tr>
<td>Table 14: Errors for Steady State Volume Flow BC</td>
<td>104</td>
</tr>
<tr>
<td>Table 15: Optimal Piston Schedules</td>
<td>123</td>
</tr>
<tr>
<td>Table 16: Selection of Test Cases – Exceptionally Good Algorithm Performance</td>
<td>139</td>
</tr>
<tr>
<td>Table 17: Selection of Test Cases – Similar, Good Performance: DE Algorithms</td>
<td>150</td>
</tr>
<tr>
<td>Table 18: Selection of Test Cases – Similar, Bad Performance: DE Algorithms</td>
<td>151</td>
</tr>
<tr>
<td>Table 19: Selection of Test Cases – Varying Performance: DE Algorithms</td>
<td>153</td>
</tr>
<tr>
<td>Table 20: Selection of Test Cases – Varying Performance Among DE OAs</td>
<td>156</td>
</tr>
<tr>
<td>Table 21: Selection of Test Cases – Modifications to DN3</td>
<td>160</td>
</tr>
</tbody>
</table>
### LIST OF FIGURES

<table>
<thead>
<tr>
<th>FIGURE</th>
<th>PAGE</th>
</tr>
</thead>
<tbody>
<tr>
<td>Figure 1: Various Sections of Rastrigin Function</td>
<td>22</td>
</tr>
<tr>
<td>Figure 2: Example of Global Best Vector</td>
<td>24</td>
</tr>
<tr>
<td>Figure 3: Example of Population Weighted Average</td>
<td>25</td>
</tr>
<tr>
<td>Figure 4: Example of Negative of the Global Worst Vector</td>
<td>26</td>
</tr>
<tr>
<td>Figure 5: Example of Population Movements and their Accompanying Centroids Using Two Fictitious Algorithms CA1, and CA2</td>
<td>29</td>
</tr>
<tr>
<td>Figure 6: Utopia vector for two-objective minimization problem</td>
<td>43</td>
</tr>
<tr>
<td>Figure 7: Relative Performance of Blind Hybrids with Different Constituent Selection Methods</td>
<td>51</td>
</tr>
<tr>
<td>Figure 8: CKO Selection Preference of PWA Blind hybrid</td>
<td>52</td>
</tr>
<tr>
<td>Figure 9: CKO &amp; DN3 Selection Preference of IBWA Blind hybrid</td>
<td>53</td>
</tr>
<tr>
<td>Figure 10: Selection Preferences of GB and PGB Blind hybrids</td>
<td>53</td>
</tr>
<tr>
<td>Figure 11: Selection Preferences of NGW and AV1 Blind hybrids</td>
<td>54</td>
</tr>
<tr>
<td>Figure 12: Relative Performance of Blind hybrids and their Constituent algorithms</td>
<td>55</td>
</tr>
<tr>
<td>Figure 13: Convergence Histories for Modified BST on TC 110, and GB Blind hybrid Constituent Algorithm Selection</td>
<td>56</td>
</tr>
<tr>
<td>Figure 14: Convergence Histories for TC 110 – Blind hybrids</td>
<td>56</td>
</tr>
<tr>
<td>Figure 15: Relative Performance of Blind hybrids</td>
<td>57</td>
</tr>
<tr>
<td>Figure 16: Selection Preferences of AUTO1 and AUTO2 Hybrids</td>
<td>58</td>
</tr>
<tr>
<td>Figure 17: Selection Preferences of DIR Hybrids</td>
<td>58</td>
</tr>
<tr>
<td>Figure 18: Best Performance of DIR, AUTO1, AUTO2, and their Constituent Algorithms</td>
<td>59</td>
</tr>
<tr>
<td>Figure 19: Worst Performance of DIR, AUTO1, AUTO2, and their Constituent Algorithms</td>
<td>60</td>
</tr>
</tbody>
</table>
elements .......................................................................................................................... 115

Figure 61: Validation of MOC code – Pressure BC – Half Stoke of Air, 50 mesh elements .......................................................................................................................... 116

Figure 62: Validation of MOC code – Volume Flow BC – Full Stoke of Air, 50 mesh elements .......................................................................................................................... 116

Figure 63: Validation of MOC code – Pressure BC – Full Stoke of Air, 50 mesh elements .......................................................................................................................... 117

Figure 64: Optimization of Piston Operation - Fully Flooded Pipeline....................... 120

Figure 65: Optimization of Piston Operation – Pipeline with Air on Upstream Side of Plug .......................................................................................................................... 122

Figure 66: Optimization of Piston Operation – Asynchronous (a) Single Phase (b) Pipeline with air on Upstream Side of Plug................................................................. 122

Figure 67: Best Performance of Unmodified Optimization Algorithms .................. 138

Figure 68: Convergence Histories, Test Case 4............................................................. 141

Figure 69: Convergence Histories, Test Case 45........................................................... 141

Figure 70: Convergence Histories, Test Case 220.......................................................... 142

Figure 71: Convergence Histories, Test Case 234.......................................................... 143

Figure 72: Convergence Histories, Test Case 236.......................................................... 144

Figure 73: Convergence Histories, Test Case 238.......................................................... 145

Figure 74: Convergence Histories, Test Case 239.......................................................... 146

Figure 75: Convergence Histories, Test Case 242.......................................................... 147

Figure 76: Convergence Histories, Test Case 331.......................................................... 148

Figure 77: Convergence Histories, Test Case 378.......................................................... 149

Figure 78: Convergence Histories - Similar, Good Performance: DE Algorithms ....... 150

Figure 79: Convergence Histories –Curse of Dimensionality Among DE Algorithms.. 152
Figure 80: Convergence Histories – Varying Performance Among DE Algorithms ..... 154
Figure 81: Convergence Histories – Curse of Dimensionality Among PSO Algorithms 155
Figure 82: Convergence Histories – Varying Performance of PSO Algorithms .......... 156
Figure 83: Convergence Histories – Varying Performance of Yang/Simon Algorithms 157
Figure 84: Best Performance of Unmodified vs Modified Optimization Algorithms .... 158
Figure 85: Convergence Histories – Modified DN3 ................................................. 159
Figure 86: Convergence Histories – PSO and PRD ................................................ 160
Figure 87: Convergence Histories – Various Improvements Due to Modifications ..... 161
CHAPTER 1

1. INTRODUCTION

The past 30 years have seen a wave of novel global optimization algorithms published. Many of these algorithms draw inspiration from natural processes, while others combine successful features from different algorithms into a single, more robust algorithm through a process called hybridization. One powerful approach to hybridization is to develop a meta-heuristic that performs automatic switching among a collection of constituent optimization algorithms. This approach is attractive in part because it is modular. That is, new modular optimization algorithms can be added or removed from the hybrid algorithm at any time, making it easy to update the hybrid with the latest methods. Furthermore, the meta-heuristic itself can be modified without making changes to the constituent algorithms. In essence, this approach is one of the most customizable approaches to optimization algorithm development. This thesis proposes a new meta-heuristic scheme for hybridization based on search vectors that serve as guides to aid in the selection of a constituent algorithm that is appropriate for the given problem.

The selection of an optimization problem for a given problem, however, raises a variety of concerns regarding the quality of the criteria used. Specifically, there needs to be a mechanism that relates the topology of an objective function to some feature of the optimization algorithm such that the algorithm selected for the problem is likely to be the best performing algorithm of all constituent algorithms available to the hybrid. This thesis explores some of the theory surrounding this topic, and demonstrates how using search vectors provides a good first indication about which global search algorithm is appropriate for a specific problem.
In light of the sheer volume of optimization algorithms presented, this thesis also explores questions relating to how the performance of hybrid optimization algorithms ought to be evaluated. Many authors select a small set of test cases with which to benchmark the performance of their algorithms. While this approach is simple, a small set of test cases is prone to produce misleading results because it is possible to tailor an optimization algorithm to a class of problems. Specialization is an undesirable quality for hybrid algorithms, because the primary goal of hybridization is to increase the number and type of problems that can be solved by the algorithm. This thesis evaluates the proposed hybrid optimization algorithm using a set of nearly three hundred test cases, so that any specialization in the algorithm will be revealed. After evaluating the hybrid optimization algorithm using standard techniques, this thesis proposes a general framework for evaluating the performance of hybrids that use automatic switching meta-heuristics.

After benchmarking the hybrid algorithm against other popular algorithms, the hybrid algorithm is applied to a real-world problem. The Applied Research Center at Florida International University has been engaged in research for the Department of Energy including the evaluation of pipeline unplugging technologies. One such technology proposed by the Applied Research Center is called the Asynchronous Pulse Unit. The goal of this pipeline pressurization technology is to build up a large enough pressure to dislodge plugs that form within the pipeline. In order to assist in this effort, this thesis presents an improved Method of Characteristics code using novel boundary conditions that accurately model the experimental set ups used by the Applied Research Center over the past two years. The code models the propagation of pressure transients in
black-iron pipelines created by piston pumps connected to their inlet. By coupling the Method of Characteristics code to the hybrid optimization algorithm, this thesis demonstrates that the hybrid algorithm can predict optimal piston pump operation schedules that produce high pressures across the plug while simultaneously operating within the specified safety limitations of the pipeline. This information, generated inexpensively using widely available desktop computers, can be used to help guide experimental efforts in obtaining the best possible piston pump operation schedule for the pipelines at their disposal.

1.1. Personal Contributions

As a member of the DOE Fellows program at the Applied Research Center, I assisted in the construction of experimental pipelines, as well as gathering experimental data. Additionally, I performed computational fluid dynamics simulations of the pipeline using ANSYS Fluent in addition to writing an updated Method of Characteristics code based on [1]. As a member of the MAIDROC Laboratory, I proposed the hybrid presented in this thesis and wrote the computer code for the hybrid using C++ and OpenMPI. I translated 300 standard optimization test cases (used for benchmarking optimization algorithm performance) originally written in Fortran 77, into C++. I benchmarked the performance of the hybrid and other optimization algorithms (listed below) on MAIDROC’s 240-core cluster named Tesla, as well as a shared-memory architecture platform, and interfaced the hybrid with Method of Characteristics code for the optimization of the Asynchronous Pulse Unit. I wrote the post-processing code used to create most of the figures and tables in this thesis using MATLAB R2010a. I evaluated optimization algorithms developed by other authors including Biogeography-Based
Optimization, five variations of Differential Evolution, Particle Swarm, two variations of Quantum Particle Swarm, the Firefly Algorithm, Cuckoo Search, and the Bat Algorithm. I also proposed modifications to Differential Evolution and Particle Swarm that led to improvements in their performance. Those modified algorithms were subsequently incorporated into the hybrid optimization algorithm. I also compared the hybrid developed in this thesis to a hybrid named OPTRAN, which was previously developed by researchers at MAIDROC.
CHAPTER 2

2. LITERATURE REVIEW

2.1. General Optimization Theory

    Engineering problems are inherently optimization problems. The task is never merely to design a product, but to design a product that meets a specific set of goals in the best possible way. Therefore, it is possible to formulate engineering problems at a higher level of abstraction so that algorithms can generate a design that satisfies these goals.

2.1.1. Designs as Vectors

    In optimization literature, vectors are often used to represent designs. For example, the shape of a molecule can be represented by the collection of positions of each atom in three-dimensional space. For a diatomic molecule, this would require a six-dimensional vector. The vector representing the design is called the “design vector” [2], although it can take on names such as “habitat” [3], “individual” [4], “bat” [5], or “decision variable” [6] depending on the author. The space created by the set of all possible designs for a given problem is called the design space. This thesis only deals with designs that can be expressed as vectors, therefore, all design spaces are also assumed to be vector spaces. The goal is to locate a vector in this space that satisfies the goals of the design process to the greatest possible extent. This vector is called an “optimal” solution or design.

2.1.2. Objective Functions

    The goal of an engineering problem is associated with a metric (e.g. cost, or weight). These metrics are functions of the design parameters (e.g. a larger beam will
cost and weigh more than a smaller one). Thus, an engineering problem can be represented by a mathematical function called an “objective function,” that is designed to reflect the goal of the optimization process (e.g. minimize cost). By convention, the objective function is written such that the goal is always met by minimizing the function [7],

\[
\text{Minimize } U = U(\bar{x})
\]

(2.1)

where \( U \) is the objective function and \( \bar{x} \) is the design vector.

Although a simple engineering problem may only have a single objective function, that objective function can have more than one minimum. The lowest possible value of the objective function is called the global minimum, while other minima with higher values are called local minima. A function with a single global minimum is called “unimodal,” while a function with multiple global minima is called “multimodal” [8].

Real-world engineering problems are usually multi-objective. If the objective functions have distinct global minima (i.e. the objectives are conflicting) the problem has a cardinality greater than one, and there is no single solution to the problem [8]. Therefore, there exists a set of solutions called the “Pareto optimal set,” or “Pareto Front” that satisfies each objective such that no improvement can be made in one objective without diminishing performance in another objective [8].

The optimization algorithms considered in this thesis are designed for a single-objective optimization problem, but often have multi-objective extensions.

2.1.3. Constraints

Real-world engineering problems also have constraints. These constraints may be limitations on the value of an objective function (e.g. cost cannot exceed a given value),
the value of a design parameter (e.g. wing span must remain below a given value), or some combination thereof. Constraints can be expressed as inequalities or equalities, as follows,

\[ g(x) < 0 \] \hspace{1cm} (2.2) \\
\[ h(x) = 0 \] \hspace{1cm} (2.3)

For example, an equality constraint for the composition of a metal alloy would be that the sum of the percentages of all alloying elements must equal 100%. Due to precision issues that arise from floating point arithmetic, equality constraints are often recast as,

\[ |h(x)| < \epsilon \] \hspace{1cm} (2.3a)

where \( \epsilon \) is a small number (e.g. \( 10^{-7} \)).

Many optimization algorithms lack explicit mechanisms for handling constraints. Therefore, several methods have been developed including the Lagrangian and Augmented Lagrangian Methods [7], Exterior and Interior Penalty Functions [7], Rosen’s Projection [9] [10] and others. This thesis utilizes a modified exterior penalty function, which is discussed in Chapter 3.

2.1.4. Performance Benchmarking

When developing optimization algorithms, it is desirable to compare its performance to that of other algorithms because this provides an indication of the algorithm’s relative speed, accuracy, and robustness. It is common practice to use a handful of standard test cases for performance benchmarking. Some authors have pointed out that benchmarking in this way can be misleading because many standard functions contain symmetries or other features (convexity, unimodality, etc.) that can be taken
advantage of by a specialized algorithm [11]. Such algorithms perform well for the problem at hand but poorly for other types of problems. The algorithm presented in this thesis is intended for simply-connected-domain, black-box problems, and so must not be specialized.

The Schittkowski & Hock test cases [12] [13] is a set of over 300 test cases ranging from unconstrained, smooth and continuous objective functions to heavily constrained, discontinuous objective functions. The dimensionality of the problems ranges from two to one hundred, as shown in Table 1 below.

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<tr>
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</tr>
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<tbody>
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<td>Dimension</td>
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This set is large and diverse enough to be considered useful and representative of a wide range of real-world problems. Therefore, this set of test cases can reveal when an optimization algorithm is tailored excessively in favor of one class of problems.

2.2. Passive Pure Optimization Algorithms

Most optimization algorithms converge to a solution in an iterative fashion based on some mathematical formula. This systematic procedure will be referred to as the search logic of the algorithm. In this thesis, a passive optimization algorithm denotes an algorithm whose search logic is static for every iteration. A dynamic algorithm changes logic based on information from the objective function, or some other scheme. A pure algorithm (here, a heuristic, and sometimes also referred to as a metaheuristic) will refer to an algorithm that uses a single logic, while a hybrid combines more than one search
logic. Hybrids are classified as metaheuristics.

2.2.1. Local Optimization

For well over 80 years, myriad algorithms have been devised for finding the local minimum of a function. Most of these methods begin with a point assumed to be near a minimum (the “initial guess”), formulate a search direction and a step size, and search along the resulting line toward the minimum. Two of the most widely used methods in this category, the DFP Conjugate Gradient Method [14] and BFGS Quasi-Newton Method [15], are called Gradient-Based Methods because they use gradient information to determine the search direction and step size. Local optimization methods suffer the drawback that they are only guaranteed to find the global optimum when the objective function is convex, because in this case the sole local optimum is also the global optimum. Otherwise, local optimization algorithms simply converge to the nearest minimum and stop searching once they locate it. Gradient-based methods have the additional drawback that the function must be differentiable.

2.2.2. Global Optimization

Global optimization algorithms are designed to search large portions of the design space in order to locate the region containing the global minimum. This is usually achieved using a set of design vectors (often called a “population” of design vectors). Many of these algorithms are inspired by observations from biology. One of the first of these algorithms to be developed, Genetic Algorithms (GA) [16], is inspired by evolutionary changes in DNA and is a combinatorial algorithm. Other algorithms are designed for continuous domains and use linear combinations of design vectors in their search logic. Two very popular methods used in the hybrid developed for this thesis are
presented below. The remaining algorithms considered for this thesis are briefly presented in the Appendix.

### 2.2.2.1. Particle Swarm Optimization

Particle Swarm Optimization (PSO) has become very popular due to its simplicity and speed. It is based on the social behavior of various species and uses linear combination of design vectors to form a new design [17]. Going forward, the equation used to modify the designs will be referred to as the update equation. The basic PSO algorithm is given below.

1) Create initial set (population) of design vectors.
2) Evaluate objective function(s) for each design vector.
3) Store copy of initial population to serve as individual best vectors and store the global best.
4) Begin main loop:
   a. For each solution in the population,
      i. Apply update equation (2.4).
      ii. Evaluate objective function(s) using new design vector.
      iii. Replace individual best with new solution if new solution is superior.
   b. Replace global best if best new solution is superior to previous global best.
5) End main loop once population converges or maximum number of iterations is reached.

The update equations are,

\[
\tilde{X}_i^{g+1} = \tilde{X}_i^g + \tilde{V}_i^g \tag{2.4}
\]

\[
\tilde{V}_i^g = \alpha \tilde{V}_i^{g-1} + \beta R_1 (\tilde{X}_{best,i} - \tilde{X}_i) + \beta R_2 (\tilde{X}_{best,G} - \tilde{X}_i) \tag{2.5}
\]

where \( \tilde{V} \) is the so-called “velocity vector,” \( \alpha \) and \( \beta \) (0.5 and 2, respectively) are user defined scalars, \( g \) is the iteration number, and \( R_1 \) and \( R_2 \) are uniformly distributed random numbers \( \epsilon [0,1] \). The vector \( \tilde{X}_{best,i} \) corresponds to the best value ever held by the \( i^{th} \)
design vector (referred to here as the “individual best”, and \( \vec{X}_{best,G} \) is the best solution ever found (also known as the “global best”). The first term on the right of Eq. 2.5 is the “inertia,” which is effectively a scalar multiple of the velocity from the previous iterations. In this thesis, the velocity is initially set to zero.

### 2.2.2.2. Differential Evolution

Differential Evolution (DE) utilizes an update equation in order to generate a new design vector, and replaces an existing design vector with the new one if the new design vector is superior. The standard DE algorithm can be described as follows:

1) Create initial population of candidate design vectors.
2) Evaluate objective function(s) for each design vector.
3) Begin main loop:
   a) Copy original population to temporary population.
   b) For each design vector in the temporary population,
      i. Create a new design vector:
         1. Randomly select one dimension, \( j \), of design vector and apply update equation.
         2. For each dimension (excluding \( j \)) of the current design vector,
            a. If \( R < CR \), apply update equation, otherwise, leave unchanged.
      ii. Evaluate objective function(s) for new design vector.
     iii. Compare new design vector to corresponding design vector from original population
     iv. If new design vector is superior to original design vector, replace the original with the new design vector.
4) End main loop once population converges or maximum number of iterations is reached.

In the above algorithm, \( CR \) is a user-defined scalar \( \epsilon [0,1] \) known as the “crossover rate,” and \( R \) is a uniformly distributed, random number \( \epsilon [0,1] \).

There are many forms of DE currently in use, several of which vary only by the update equation used. Three particularly successful forms are the so-called rand/1/bin, and best/2/bin proposed in [18] as well as Donor3 proposed in [19]. Their respective update equations are as follows,
where $Y$ is the resulting coordinate in the $k^{th}$ dimension of the new design vector, and $F$ is a weighting factor (a user-defined scalar $\epsilon [0,2]$). The variable $X$ denotes a coordinate from an existing design vector vector, and the subscript $r$ indicates that it was randomly selected. Therefore, the component $Y$ in Eq. 2.6, for rand/1/bin, is a linear combination of components from three distinct, randomly selected design vectors, while Eq. 2.7, for best/2/bin is the linear combination of four distinct, randomly selected design vectors and the global best design vector. The Donor3 method utilizes a weighted average of three components, where $\lambda_1$, $\lambda_2$, and $\lambda_3$ are uniformly distributed, random numbers $\epsilon [0,1]$.

The fourth variation of DE is called Trigonometric DE, or TDE [4]. Its mutation equation also contains an average, but here it is the arithmetic mean. The right-hand difference term is replaced with three differences, whose scalar coefficients are determined by the objective function values corresponding to the respective design vectors. The update equation is,

$$
\hat{X}_k = \frac{\hat{X}_{r_1} + \hat{X}_{r_2} + \hat{X}_{r_3}}{3} + \left(p_2 - p_1\right)(\hat{X}_{r_1} - \hat{X}_{r_2}) + \left(p_3 - p_2\right)(\hat{X}_{r_2} - \hat{X}_{r_3}) + \left(p_1 - p_3\right)(\hat{X}_{r_3} - \hat{X}_{r_1})
$$

(2.9)

where,

$$
p' = \left|U(\hat{X}_{r_1})\right| + \left|U(\hat{X}_{r_2})\right| + \left|U(\hat{X}_{r_3})\right|
$$

(2.10)
\[ p_1 = \frac{U(\tilde{x}_{r1})}{p'} \]
\[ p_2 = \frac{U(\tilde{x}_{r2})}{p'} \]
\[ p_3 = \frac{U(\tilde{x}_{r3})}{p'} \]  

(2.11)

Clearly, if all three \( p \) values equal zero, this method will yield an undefined value. Therefore, as a precaution, the following condition was added to this algorithm,

\[ if(p' = 0) \quad p' = 10^{40} \]  

(2.12)

This condition was implemented in lieu of the more common practice of adding a small constant to \( p' \) so that the algorithm’s performance will not be biased when the values of \( p \) are small. Apart from a distinct update equation, this method differs from the previous method in that it also contains an additional condition,

a) For each dimension (excluding j) of the current design vector,
   i. If \( R_1 < CR \),
      1. If \( R_2 < M_t \) use Eq. 2.9 to create mutant, else use Eq. 2.6

where \( R_1 \) and \( R_2 \) are distinct, random numbers \( \epsilon [0,1] \), and \( M_t \) is a user-defined scalar \( \epsilon [0,1] \). The method’s developers suggested that \( M_t \) be set to 0.05, because TDE is a “greedy” search method [4]. Lampinen and Fan also remarked that if \( M_t \) were set to zero, this method would reduce to DE rand/1/bin.

Global optimization algorithms often have the drawback that they require many function evaluations (sometimes thousands or more) to locate the region containing the global minimum, and can require just as many evaluations to converge to the solution itself once within that region.

2.3. Hybrid Optimization Algorithms

Depending on the objective function topology, some algorithms perform better
than others. Thus, it is desirable to combine algorithms into a more complex, hybrid algorithm so that their strengths can be leveraged and their weaknesses mitigated. Algorithms that make up the hybrid are called “constituent algorithms.” For example, using a global optimization algorithm to locate the region containing the global minimum and switching to a local optimization algorithm dramatically improves the likelihood of reaching the global minimum in a computationally efficient manner. Such a hybrid is classified as a “global-local” hybrid (for example, see [20]).

2.3.1. No Free Lunch

The question then becomes, is it possible to create a black-box optimization algorithm (hybrid or not) with competitive or even superior performance against all other algorithms for all possible optimization problems? In other words, is it possible to create an effective general-purpose optimization algorithm? Some authors have said no (given certain assumptions). Wolpert and Macready [21] claim that “the average performance of any pair of algorithms across all possible problems is identical.” Their “No Free Lunch” (NFL) theorem indicates that if one algorithm is superior to another for a given set of optimization problems, it must then be inferior over another set of optimization problems. Droste, Jansen, and Wegener, [22] show that while this is true under some circumstances, it is not absolutely applicable because there are instances where an algorithm can perform in an above-average sense. They propose an “Almost No Free Lunch” (ANFL) theorem, which states that an algorithm is efficient at solving a certain class of problems because it implicitly utilizes information about the structure of the function. Therefore, “it is possible to describe other simple functions which are closely related to functions easy for [the algorithm] and which, nevertheless, are hard for [the algorithm].” Yang [23] points
out that Wolpert and Macready’s NFL theorem is based on assumptions that do not always apply: (1) the design space is countable and finite, and (2) the algorithm does not revisit the same region. It has been shown that if the problem domain is continuous (uncountable), or not closed under permutation (revisiting), that the NFL theorem does not hold [24] [25] [26]. Therefore, under certain real-world conditions it appears possible to develop a black-box optimization algorithm with above-average performance.

2.3.2. Hybrid Algorithm Architectures

There currently exist a great many optimization algorithms in literature, and there are untold thousands of ways to combine them. Talbi [27] developed a taxonomy, referenced below, to categorize hybrid algorithm architectures, which was later expanded upon by authors including Raidl [28]. For the purpose of this thesis it suffices to say that, in general, there are three noteworthy approaches to hybridization, each based on the way in which the population of design vectors is operated on. A global-local hybrid can be created from any one of these architectures simply by passing some or all of the population to a local optimization algorithm at some stage of the optimization process.

2.3.2.1 Competitive

Competitive hybrids (high-level, relay [27]) switch between constituent algorithms such that only a single constituent algorithm operates on the entire population at a time. For example, the competitive hybrid might begin with PSO, and then switch to DE once some criterion is met. The hybrid presented in this thesis is of this type. Additional examples will be discussed in greater detail below.

2.3.2.2 Cooperative

Cooperative hybrids (high-level teamwork [27]) allow multiple constituent
algorithms to operate on subsets of the population simultaneously during each iteration. Unlike competitive hybrids, the switching logic pertains to the size of the population passed to each constituent algorithm, which may be constant. An example of such hybrids can be found in [29].

2.3.2.3 Merged

Merged hybrids (low-level, relay [27]) combine the essential components of different constituent algorithms into a single algorithm. In this way, the basic operations of all constituents are executed on the entire population during each iteration of the hybrid algorithm. Thus, the population is updated multiple times during a single iteration. This is different from a competitive hybrid in that competitive hybrids may not execute a given constituent at all, and it is different from the cooperative hybrid in that the entire population is operated on. Examples of such hybrids can be found in [30] [31] [32].

2.3.3. Switching Logic

Within the scope of a competitive hybrid, the essence of the algorithm is the way it switches from one constituent algorithm to another. Since each constituent algorithm has its own search logic, the switching mechanism may override that logic by interrupting the sequence of events that would naturally follow. Over several years, researchers affiliated with MAIDROC have developed and tested several hybrid algorithms with automatic switching that do not override the constituent algorithm’s search logic [33] [34] [35]. Rather, these hybrids allow the constituent algorithm to proceed until it triggers some failure mode or convergence criterion, and then switch to another algorithm in order to perform an efficient local search, or perform a new global search. The interested
reader is referred to [36] [37] [38] for a discussion of multi-objective hybrids developed by MAIDROC.

The first in the series was simply a global-local algorithm made up of GA and DFP [33]. Once GA’s convergence rate slowed to a certain value, the hybrid would switch to DFP and converge to the nearest minimum. Once DFP converged, the hybrid would restart GA and this cycle would repeat a number of times in order to increase the likelihood of finding the global minimum.

The second in this series [34], combined three local optimization algorithms [DFP, the Nelder-Mead (NM) simplex method, and simulated annealing (SA)], and used GA to perform a global search. It enforced constraints using Rosen’s projection, feasible searching, and random design generation. The additional local search algorithms enabled the hybrid to switch from GA to NM in the event of a failure (a bad mutation or lost generation). If the objective function variance was small, the hybrid would switch from GA to SA. If the design vector variance was small, the hybrid would switch from GA to DFP. NM and SA would switch to DFP in the event of a stall or insufficient energy, respectively. SA would switch to NM if it exceeded a predetermined number of iterations.

The third and fourth hybrids in the series were very similar. They each included DFP, GA, NM, and sequential quadratic programming (SQP). The fourth generation [39] added a second global optimization algorithm (DE), and replaced SA with a Quasi-Newton algorithm by Pshenichny-Danilin (LM). The hybrid would begin with GA and cycle through NM, or SQP. SQP’s convergence would trigger LM, which would similarly trigger DFP. DFP’s convergence would trigger the activation of DE. Once DE’s
convergence stalled, the algorithm would switch back to GA and repeat for a given number of loops.

The fifth and sixth hybrids in the series utilized only three constituent algorithms (PSO, DE, and BFGS), where the sixth included the use of response surfaces (surrogate models that approximate the value of the objective function but are computationally faster to execute) in order to improve its overall speed [33]. The global optimization search would begin with PSO, and would switch to DE once a certain percentage of the population appeared to converge. If the search executed by DE produced an improvement in the current minimum, the hybrid would switch back to PSO. Otherwise, it would switch to BFGS to rapidly locate the nearest minimum. Once BFGS converged, the hybrid would switch back to PSO and loop as the previous hybrids did.

Each of these hybrids calls their constituent optimization algorithms in a predetermined sequence based on a set of rules (triggers). This style of hybrid architecture development uses inductive reasoning that proceeds roughly as follows:

If algorithm X behaves in manner A, this implies B.

Given B, use algorithm Y.

Otherwise, use algorithm Z.

All global-local hybrids are based on this logic (if the global optimization algorithm appears to converge, the hybrid switches to the local optimization algorithm). In cases like these, the architect of the hybrid has assumed certain characteristics about the nature of the design space, the objective function space, and the search logic of the constituent algorithms, and developed rules intended to capitalize on these characteristics. Thus, it follows that if any of the assumptions are wrong (A does not imply B), or if the rules are
inadequate (e.g. Y should not always be used given B), the hybrid risks converging to a local minimum. While the additional search logic of hybrid algorithms may increase the risk of poor convergence beyond those of its constituent algorithms, hybrid algorithms possess the greatest potential in obtaining as close to a free lunch as possible. The hybrids discussed above have been shown to outperform their constituent algorithms [33]. Therefore, a properly crafted switching mechanism is the key to a competitive hybrid’s success.
CHAPTER 3

3. SEARCH VECTOR BASED HYBRID OPTIMIZATION ALGORITHM

3.1. A Global-Global Hybrid

The motivation behind the hybrid developed in this thesis is based on two observations. The first comes from the statement in [22]: “Each search heuristic which is able to optimize some functions efficiently follows some idea about the structure of the considered functions.” Therefore, a hybrid that intelligently switches between global optimization algorithms (a global-global hybrid) can efficiently optimize a wider range of objective functions without requiring input from the user. That is, it serves as a better black-box algorithm than its constituent algorithms (in the global optimization sense).

3.1.1. Single Objective – Multiple Topologies

It is easy to read the above statement and mistakenly assume that a single objective function must have a single structure. While this is true from a strictly theoretical standpoint, this is not really the case in practice when no a priori knowledge of the problem is available. Global optimization is an inherently statistical process that begins with a sample of designs. Determinations regarding the topology of an objective function depend entirely on the distribution and size of the sample, which can be misleading because it is incomplete. The second motivation for the hybrid presented here, and the stronger statement made for it in this thesis is that, from the “perspective” of the optimization algorithm, the topology of most objective functions appears to change throughout the optimization process. That is, most objective functions have multiple “effective” topologies. Therefore, for the broader set of optimization problems there should be no expectation that any single algorithm will perform in an above-average
sense. A hybrid, on the other hand, has the potential to do so (within the scope of real-world scenarios discussed in Chapter 2).

For example, consider the 2-dimensional Rastrigin Function shown in Figure 1.

![Figure 1: Various Sections of Rastrigin Function](image)

A sample taken over the square domain bounded by either $x_1, x_2 \epsilon [-50,50]$ or $[-0.5,0.5]$ would likely indicate that the Rastrigin Function is convex. If the sample was taken over the domain $[-5,5]$, however, the function would appear to be noisy with a vaguely convex trend. Finally, a sample taken over $[-1.5,1.5]$ would result in a function that appears to be periodic in all directions. Assuming the algorithm begins with the largest domain, and assuming it is designed for convex functions, it will converge rapidly from the $[-50,50]$ region into the $[-5,5]$ region, and then slow down dramatically and probably converge to a local minimum. This is because the convexity assumption is approximately valid so
long as the population is spaced very far apart. Once the population converges to a smaller space (i.e. the distribution of the sample decreases), the assumption is no longer valid and the algorithm becomes ill-suited for the effective topology it is navigating.

Therefore, the hybrid presented here, and previously briefly introduced in [33], does not wait for a constituent algorithm to converge. Rather, it uses other indicators to determine if, and when, to use a constituent algorithm.

3.1.2. Search Vectors

In local optimization, many algorithms fall into the category of line search algorithms, whose basic equation takes the form,

$$\bar{x}_{\text{final}} = \bar{x}_{\text{initial}} + \alpha \bar{s}$$

where \( \bar{s} \) is the search direction, and \( \alpha \) is the scalar step size that enables the algorithm to “step” from the initial design to a final design. The hybrid presented here utilizes a population-based analogy of the search direction called a “search vector” with the following general form,

$$\vec{v}_{\text{search}} = \vec{v}_{\text{sample}} - \vec{v}_{\text{centroid}}$$

(3.2)

Where \( \vec{v}_{\text{search}} \) is the search vector, and \( \vec{v}_{\text{centroid}} \) is the arithmetic mean of the population. The sample point (also a vector), \( \vec{v}_{\text{sample}} \), is any vector taken from the population or calculated from some formula that is used to guide the search. In this analogy, rather than moving a single design from one location to another within the design space, the entire population is moved from one region in space to another. This movement is represented by the displacement of the population’s centroid, as follows,

$$\vec{v}_{\text{centroid,final}} = \vec{v}_{\text{centroid,initial}} + \vec{v}_{\text{search}}$$

(3.3)
The distribution of the population is not considered in this calculation, indicating that the population can disperse from or converge toward its centroid without affecting the hybrid’s switching logic. The following subsections discuss the formulas for the eight sample points used to create search directions in this thesis:

### 3.1.2.1. Global Best Vector

The Global Best vector (GB) is the fittest design vector of the population (i.e. has the lowest objective function value). An example of this vector is shown in Figure 2 below in red.

![Figure 2: Example of Global Best Vector](image)

### 3.1.2.2. Population Weighted Average

To calculate the Population Weighted Average (PWA), the population is ranked from best to worst, with the best receiving a rank equal to the population size, and the worst having a rank of one. This operation is given by (3.4) below.

\[
y_{PWA,i} = \frac{1}{\sum_{j=1}^{N} \sum_{i=1}^{\text{dim}} r_j x_{i,j} \hat{e}_i}
\]

(3.4)

where \(y_{PWA,i}\) represents the \(i^{th}\) coordinate of the PWA, \(\text{dim}\) represents the dimensionality of
the design space, \( N \) is the population size, \( r_j \) represents the rank of the \( j^{th} \) design vector in the population, \( \hat{e}_i \) is the unit vector in the \( i^{th} \) direction, and \( x_{i,j} \) is the \( i^{th} \) coordinate of the \( j^{th} \) design vector. This vector is not originally a part of the population.

### 3.1.2.3. Individual Best Weighted Average

Another vector called the Individual Best Weighted Average (IBWA) is conceptually identical to the PWA but uses the individual best population stored for Particle Swarm. In the first iteration, this vector is equal to the PWA. An example of the PWA is shown in Figure 3 below in red. For alternative (conceptually similar) combinations of individual best vectors, the interested reader is referred to [40].

![Figure 3: Example of Population Weighted Average](image)

### 3.1.2.4. Negative of the Global Worst Vector

The Negative of the Global Worst vector (NGW) is constructed by reflecting the global worst vector (the design with the highest objective function value) across the center of the domain, according to the equation below:

\[
\bar{y}_{NGW} = \bar{d}_L + \bar{d}_U - \bar{x}_{GW}
\]

(3.5)

where \( \bar{y}_{NGW} \) represents the NGW, \( \bar{d}_L \) represents the lower limit of the domain, \( \bar{d}_U \) is
the upper limit of the domain, and $\bar{x}_{GW}$ is the global worst vector. In Figure 4 below, the blue vector is the global worst, and the vector in red is the “negative of the global worst.” This vector is not originally a part of the population.

Figure 4: Example of Negative of the Global Worst Vector

3.1.2.5. Population Centroid

As discussed above, the Population Centroid (PC) is the arithmetic mean of the population. This vector is not originally a part of the population.

3.1.2.6. Individual Best Population Centroid

Another vector called the Individual Best Population Centroid (IBPC) is conceptually identical to the PC but uses the individual best population stored for Particle Swarm. In the first iteration, this vector is equal to the PC. This vector is not originally a part of the population.

3.1.2.7. Projected Global Best

The definition of the Projected Global Best (PGB) borrows from classical dynamics. The PGB is based on the formula for projectile motion under constant acceleration, as shown below.

$$x = x_0 + v_0 t + \frac{a}{2} t^2$$  \hspace{1cm} (3.6)
where \( X \) represents position, \( x_0 \) is the initial position, \( v_0 \) is the initial velocity, \( a \) is the acceleration, and \( t \) is time. Time, here, is understood to be the number of iterations. This concept was implemented due to its mathematical simplicity and because it strives to predict the location of the next GB, rather than relying strictly on population values from the current iteration. Given the GBs for the previous two iterations and the current iteration, \( x_0, x_1, \) and \( x_2 \), respectively, the PGB, represented by \( x_3 \), can be derived from the following linear system of equations:

\[
x_0 = x_0 \\
x_1 = x_0 + v_0 + \frac{a}{2} \\
x_2 = x_0 + 2v_0 + 2a
\]

This leads to a system of two equations with two unknowns, namely the initial velocity and acceleration. Using matrix notation and combining (3.6b) with (3.6c),

\[
\begin{bmatrix}
1 & 0.5 \\
2 & 2
\end{bmatrix}
\begin{bmatrix}
v_0 \\
a
\end{bmatrix} =
\begin{bmatrix}
x_1 - x_0 \\
x_2 - x_0
\end{bmatrix}
\]

(3.6d)

Inverting the coefficient matrix yields,

\[
\begin{bmatrix}
v_0 \\
a
\end{bmatrix} =
\begin{bmatrix}
2 & -0.5 \\
-2 & 1
\end{bmatrix}
\begin{bmatrix}
x_1 - x_0 \\
x_2 - x_0
\end{bmatrix}
\]

(3.6e)

Thus, \( v_0 \) and \( a \) can be expressed in terms of the previous positions.

\[
v_0 = -0.5x_2 + 2x_1 - 1.5x_0 \]

(3.6f)

\[
a = x_2 - 2x_1 + x_0
\]

(3.6g)

At the third iteration, the position equation becomes,
\[ x_3 = x_0 + 3v_0 + \frac{9}{2}a \quad (3.6h) \]

Substituting (3.6f) and (3.6g) into (3.6h) and simplifying yields the final result,

\[ x_3 = 3x_2 - 3x_1 + x_0 \quad (3.6i) \]

This equation is computed separately for each component of the PGB, as though each dimension were under its own constant acceleration. This vector is not originally a part of the population.

3.1.2.8. Average Vector 1

The Average Vector 1 (AV1) is the arithmetic mean of the GB, PWA and NGW, and was found to be advantageous in certain objective function topologies after some experimentation.

During early development of this hybrid, only a single sample point was used throughout the entire optimization process. This way, the sample point could be calculated without being evaluated (unless it was already part of the population), reducing the total number of objective function evaluations required by the algorithm. After some experimentation, it was decided that all sample points should be evaluated and compared. The sample point with the lowest objective function value was then selected for use by the hybrid. It is clear from Equation 3.3 that,

\[ \bar{v}_{\text{centroid, final}} = \bar{v}_{\text{sample}} \quad (3.3a) \]

Therefore, for each iteration, the sample point is the desired final destination for the population centroid. The sample point is one of the indicators that guides the global search (during early development, it was the only indicator). Each iteration, the goal of the hybrid switching mechanism is to shift the population toward the sample point.
because this point is assumed the correct direction for the population to move to in order to locate the global minimum. Therefore, the hybrid must select a constituent algorithm that is most likely to produce this result.

3.1.3. Constituent Algorithm Search Vector

Each constituent algorithm has some mechanism (an equation, etc.) to produce new candidate design vectors for the next iteration. The set of new candidate designs shall be called the “temporary population.” The objective function is not executed for these points (i.e. the temporary population is not evaluated), and contains as many design vectors as there are in the primary population used by the hybrid. The arithmetic mean of the temporary population shall be called the “temporary centroid.” In order to select a constituent algorithm, the hybrid will execute a constituent algorithm ten times, so that ten temporary populations are created. The temporary centroid for each temporary population is computed, and then averaged to produce one final vector (a centroid of all ten temporary populations).

Figure 5: Example of Population Movements and their Accompanying Centroids Using Two Fictitious Algorithms CA1, and CA2
This final result is called the “centroid of the constituent algorithm,” or simply the $C_{CA}$. For example, suppose the hybrid has two constituent algorithms called CA1 and CA2 (see Figure 5). It is possible that CA1 will update the current population (black dots) in one direction (red dots), while CA2 will update the population in another direction (blue dots). The vectors in Figure 5 represent the centroids of each population. Since each constituent algorithm contains random parameters, it is executed ten times so that its $C_{CA}$ has some statistical significance without excessively increasing the method’s computational cost. The $C_{CA}$ represents the location that the population centroid would be moved to if that constituent algorithm were used. This process is repeated for every constituent algorithm so that a $C_{CA}$ is computed for each (e.g. a $C_{PSO}$, $C_{DE}$, etc.). During early stages of development, the $C_{CA}$’s were compared to the sample point using some equation (discussed below) in order to determine which constituent algorithm to use, but never evaluated. Dr. Dulikravich suggested that the $C_{CA}$’s be evaluated and used as the sole indicator for constituent algorithm switching. After more experimentation, it was later decided that both $C_{CA}$’s and sample points be evaluated and used as indicators, and a more sophisticated switching logic was devised. In order to facilitate certain comparisons, the “constituent algorithm search vector” was defined as follows,

$$\vec{v}_{CA} = \vec{c}_{CA} - \vec{v}_{centro}\text{r}$$

where $\vec{c}_{CA}$ is the $C_{CA}$, and $\vec{v}_{CA}$ is the $C_{CA}$ translated so that it originates at the centroid of the population. When needed, the $C_{CA}$ can be compared directly to the sample point, and the search vector can be compared directly to the constituent algorithm search vector.

### 3.2. Search Vector Based Hybrid

One of the goals in developing the Search Vector Based Hybrid was to minimize
the number of additional objective function evaluations required by the hybrid beyond those performed by the constituent algorithms (minimize overhead).

<table>
<thead>
<tr>
<th>Category</th>
<th>Evaluate &amp; Compare Search Vector</th>
<th>Evaluate &amp; Compare $C_{CA}$</th>
</tr>
</thead>
<tbody>
<tr>
<td>Blind</td>
<td>No</td>
<td>No</td>
</tr>
<tr>
<td>Direct</td>
<td>N/A</td>
<td>Yes</td>
</tr>
<tr>
<td>Auto</td>
<td>Yes</td>
<td>Yes</td>
</tr>
</tbody>
</table>

Therefore, during the early stages of development, none of the search vectors or $C_{CAS}$ were evaluated unless they were already part of the population. Since these vectors guide the population without any information about the objective function at that location, this category of search vector hybrid is called a “Blind Hybrid Optimization Algorithm.” The “Direct Hybrid Optimization Algorithm” does not compute search vectors at all, but at the suggestion of Dr. Dulikravich, computes and evaluates the $C_{CAS}$ and uses their values to compare each constituent algorithm directly. Both the Blind and Direct hybrids have an automatic constituent algorithm switching mechanism, but do not automatically switch between search vectors. That is, the Blind and Direct hybrids use a single search vector throughout the entire optimization process. The “Auto Hybrid Optimization Algorithm” evaluates the sample points so that it can automatically switch between search vectors and constituent algorithms each iteration.

In order to achieve this functionality, the search vectors and constituent algorithms are implemented as modules. This provides the added benefit that the hybrid can utilize any set of algorithms, whether it is the same algorithm tailored for different problems (e.g. multiple modules of Differential Evolution with different parameters for
different classes of problems), or a variety of different methods including newly published algorithms.

3.2.1. Constituent Algorithm Selection Criteria

Although the hybrid’s search logic assumes that the sample point is the best direction to move in, there are several ways to use this information when selecting a constituent algorithm. The following five selection methods have been developed for this thesis:

3.2.1.1. Lowest Distance

This method calculates the Euclidean distance between the $C_{CA}$ and the sample point. This distance is calculated in the design space; therefore, the objective function value is not taken into consideration.

$$d = \sqrt{\sum_{i=1}^{\dim}(\vec{v}_{\text{sample},i} - \vec{C}_{CA,i})^2}$$  \hspace{1em} (3.8)

The constituent algorithm whose $C_{CA}$ is closest to the sample point will be selected.

3.2.1.2. Highest Dot Product

This method calculates the dot product between the search vector and the constituent algorithm search vector. That is,

$$\frac{\vec{v}_{\text{search}} \cdot \vec{v}_{CA}}{\|\vec{v}_{\text{search}}\| \|\vec{v}_{CA}\|} = \frac{\left(\vec{v}_{\text{sample}} - \vec{v}_{\text{centroid}}\right) \cdot \left(\vec{c}_{CA} - \vec{v}_{\text{centroid}}\right)}{\|\vec{v}_{\text{sample}} - \vec{v}_{\text{centroid}}\| \|\vec{C}_{CA} - \vec{v}_{\text{centroid}}\|}$$  \hspace{1em} (3.9)

Again, the objective function value is not taken into consideration. The constituent algorithm whose centroid is closest to parallel to the search vector will be selected. This formulation cannot be used when the sample point is the population centroid (the lowest distance can be used instead).
3.2.1.3. Lowest Scaled Distance

Similar to the lowest distance, this method uses the following equation below,

$$d_s = (d + 1)^\left[\left(\frac{\sigma_{\text{temp}}}{\sigma_{\text{pop}}} - \frac{1}{2}\left(\frac{\sigma_{\text{temp}}}{\sigma_{\text{pop}}} - 1\right)\right)\right]$$  \hspace{1cm} (3.10)

where $d_s$ is the “scaled” distance between the sample point and the CCA, while $d$ is the Euclidean distance from Eq. 3.8. The term in brackets is the exponent of the $d+1$ term, and contains the ratio between the standard deviation of the temporary population, $\sigma_{\text{temp}}$, and the current population standard deviation, $\sigma_{\text{pop}}$. The standard deviation are based on the distances between the populations under consideration and their respective centroids. The exponent rewards constituent algorithms whose standard deviation is less than the existing population’s standard deviation, or greater than 50% of that value. This was developed in an effort to penalize methods that converge too quickly or cause the population to spread out further. The constituent algorithm with the lowest scaled distance is selected.

3.2.1.4. Direct CCA Comparison

This method, proposed by Dr. George Dulikravich, ignores the objective function values of sample points. Instead, it evaluates the CCA's generated by the constituent algorithms and selects the algorithm with the minimum CCA.

3.2.1.5. Pareto Ranking Method

This method treats the Lowest Distance and Direct Comparison methods as two distinct objectives. The constituent algorithms are ranked using the Pareto dominance scheme in [8]. The constituent algorithm is then randomly selected from the Pareto optimal set (the constituent algorithms that are non-dominated in terms of lowest distance
and minimum $C_{CA}$. Although developed independently, and applied differently, the concept of using multi-objective processes in a single objective problem appears in [41], and its references.

The various search vector based hybrids developed for this thesis use one of these constituent algorithm selection methods depending on the information available. During early development, three methods were tested with the Blind hybrid. The results are discussed in Chapter 4.

<table>
<thead>
<tr>
<th>Category</th>
<th>Selection Methods Used</th>
</tr>
</thead>
<tbody>
<tr>
<td>Blind</td>
<td>Lowest Distance</td>
</tr>
<tr>
<td></td>
<td>Highest Dot Product</td>
</tr>
<tr>
<td></td>
<td>Lowest Scaled Distance</td>
</tr>
<tr>
<td>Direct</td>
<td>Direct $C_{CA}$ Comparison</td>
</tr>
<tr>
<td>Auto</td>
<td>Pareto Ranking Method</td>
</tr>
</tbody>
</table>

The latest version of the hybrid (Auto) currently utilizes six constituent algorithms: PSO, PSO with Random Differences (PRD), Modified Quantum Particle Swarm (MQP), DE best/2/bin with randomized parameters (BST), DE Donor3 with randomized parameters (DN3), and Cuckoo Search (CKO). See the Appendix for descriptions of each algorithm.

Early versions of the Blind hybrid used the Firefly Algorithm (FFA) in lieu of DN3 in order to diversify the collection, but it was replaced due to its high computational expense (overhead).

3.2.2. Hybrid Algorithm Architecture

It is now possible to describe the search vector based hybrid algorithm in detail. The basic steps (automatic switching) performed by this hybrid during each iteration can be summarized as follows:
I) Calculate each sample point  
   i) Evaluate each sample point (Auto only)  
II) Calculate each CCA  
   i) Evaluate each CCA (Direct and Auto only)  
III) Perform constituent algorithm selection  
IV) Operate on population using constituent algorithm

Therefore, this hybrid does not have any rules or equations enabling it to generate new designs of its own. Rather, the initial population is created using a random number generator, and the constituent algorithms generate every subsequent population. The hybrid architecture “wraps around” the constituent algorithms.

3.2.2.1. Random Number Generator

The hybrid is equipped with two random number generators: the ran2 function given in [42], as well as Sobol’s Algorithm [43] downloaded from [44]. Ran2 is the hybrid’s default random number generator due to its speed.

3.2.2.2. Population Size

A typical global optimization algorithm only operates on its population once per iteration, but the search vector hybrid executes its constituent algorithms a total of $10M + 1$ times per iteration, where $M$ is the number of constituent algorithms. This means that the current version of the hybrid loops over the population 61 times per iteration (the first 60 times are simply to generate temporary populations). For this reason, and due to the high computational expense typical of objective function evaluations, it is desirable to reduce the population size.

One popular approach to determining the population size is to vary it linearly with the dimensionality of the problem. For example, the following might be used,

$$N = 10 \dim$$ (3.11)
where \( \text{dim} \) is the dimensionality of the problem, and \( N \) is the population size. In order to reduce the size of the population for problems with high dimensionality, the following nonlinear function was developed after several tests,

\[
N = \text{ceil} \left( \frac{12.59\text{dim}^{2/3} + 14.142\sqrt{\text{dim}} + 65.5\log(\text{dim})}{3} \right)
\]  

(3.12)

where ceil is a function that rounds a number up to the nearest integer. Both (3.11) and (3.12) result in a population size of 20 for a 2D problem.

### 3.2.2.3. Constraint Enforcement

For simplicity, the hybrid uses a modified objective function based on a type of exterior penalty function. Since the global minima of the Schittkowski and Hock test cases are published, the objective function is first modified to reflect the error \( (E) \), that is the difference between the current value and the published minimum value \( (U^*) \), as follows,

\[
E(\bar{x}) = U(\bar{x}) - U^*
\]  

(3.13)

This is then combined with a penalty function. The standard form given for an exterior penalty function given in [7] is,

\[
p(\bar{x}) = \sum_{i=1}^{m} \max[0, g_i(\bar{x})]^2 + \sum_{j=1}^{l} [h_j(\bar{x})]^2
\]  

(3.14)

where \( m \) is the number of inequality constraints, and \( l \) is the number of equality constraints. This penalty function modifies the objective (error) function as follows,

\[
W(\bar{x}) = E(\bar{x}) + r_p p(\bar{x})
\]  

(3.15)

where \( W \) is the modified objective function, and \( r_p \) is a user-defined scaling factor. The scaling factor serves to increase or decrease the magnitude of the penalty generated by
the penalty function. If the penalty is too small relative to the magnitude of the objective function, this form of constraint enforcement can easily yield infeasible results, but if it is too large, it may yield suboptimal results. The greater concern, however, is feasibility. Because the scaling factor is user-defined its value often depends on the user’s intuition, or is tailored to suit a specific class of problems or algorithms. To alleviate this issue (letting \( r_p = 1 \)), the following penalty function is used,

\[
p(\vec{x}) = n|E(\vec{x})| + \sum_{i=1}^{m} \max[0, g_i(\vec{x})] + \sum_{j=1}^{l} |h_j(\vec{x})| \tag{3.16}
\]

where \( n \) is the total number of constraints violated. This formula ensures that the penalty function always has a magnitude at least as large as that of the objective function. It is important to note that this form of constraint enforcement changes the effective topology of the objective function, often making a function non-differentiable.

### 3.2.2.4. Blind Hybrid Algorithm Steps

The earliest of the search vector based hybrids developed, this hybrid requires the fewest additional objective function evaluations. Its primary advantage, however, is not in its computational cost, but its utility as a research tool. By forcing the search to proceed in a particular direction, it is possible to gain insights into the relationship between the effective topology of the objective function and the optimization algorithm that is ultimately selected. The algorithm is as follows,

I) Initialize population of design vectors  
II) Evaluate objective function for all design vectors  
III) Begin Hybrid Loop  
   i) Calculate sample point  
   ii) Calculate all \( C_{CAS} \)  
   iii) Perform constituent algorithm selection  
   iv) Operate on population using constituent algorithm  
   v) If convergence detected, end loop, otherwise return to step i)
IV) End algorithm

As demonstrated in the algorithm, the objective function is only evaluated following initialization and during step (iv).

3.2.2.5. Direct Hybrid Algorithm Steps

The second of the search vector based hybrids presented, this hybrid uses CCAs in lieu of search vectors and evaluates them. Therefore, it requires additional objective function evaluations (equal to the number of constituent algorithms used).

I) Initialize population of design vectors
II) Evaluate objective function for all design vectors
III) Begin Hybrid Loop
   i) Calculate all $C_{CA}$
   ii) Evaluate all $C_{CA}$
   iii) Select constituent algorithm with minimum $C_{CA}$
   iv) Operate on population using constituent algorithm
   v) If convergence detected, end loop, otherwise return to step i)
IV) End algorithm

Apart from the direct comparison in step (iii), the $C_{CA}$s are never used again. Future versions of the hybrid may incorporate these vectors into the population to improve the diversity of the population, or to replace inferior design vectors.

3.2.2.6. Auto Hybrid Algorithm Steps

The latest version of the search vector hybrid has the greatest overhead cost, but also the greatest amount of information with which to make a decision. The algorithm is as follows,

I) Initialize population of design vectors
II) Evaluate objective function for all design vectors
III) Begin Hybrid Loop
   i) Calculate all sample points
   ii) Evaluate all sample points
   iii) Select minimum sample point
   iv) Calculate all $C_{CA}$

v) Evaluate all CCAs
vi) Select constituent algorithm using Pareto Ranking Method
vii) Operate on population using constituent algorithm
viii) If convergence detected, end loop, otherwise return to step i)
IV) End algorithm

The first version of the Auto hybrid did not incorporate the sample points or CCAs into the population. Certain constituent algorithms utilize the “global best” vector in their search (e.g. DE and PSO). Since the selected sample point may be superior to the best design vector in the population, the second version of the hybrid passed the sample point to these constituent algorithms. The results are discussed in Chapter 4.

3.2.3. Ramifications of Hybrid Algorithm Architecture

This section will discuss the implications of the structure of the Auto hybrid optimization algorithm. The Auto hybrid will simply be referred to as “the hybrid.”

3.2.3.1. Search Logic

The hybrids discussed in Section 2.3.3. allow their global optimization constituent algorithms to converge, which suggests that these hybrids assume there is a “good match” between the constituent algorithm and the objective function topology. The search vector based hybrid switching mechanism often prohibits any one constituent algorithm from running completely to convergence. Rather, the hybrid compares the constituent algorithm to the search vector and assumes that a good match between the CCA and the search vector indicates that the constituent algorithm is a good match for the effective topology (which changes each iteration). Therefore, the search logic inherent in the constituent algorithm is overridden by the search logic of the hybrid.

Like all algorithms, the hybrid’s structure is well suited for certain classes of problems. For example, the hybrid assumes that the sample point with the lowest
objective function value is the best. Therefore, if there exists a class of “deceptive”
functions for which this assumption is invalid, this approach may impair the hybrid. An
improvement on this hybrid could include the addition of statistical metrics that could
warn against misleading functions, and give the hybrid a means by which to select other,
more appropriate sample points. For the purpose of this thesis, however, a random
parameter was introduced in the algorithm. Each iteration after the first, there is a 10%
chance that the hybrid will reselect the constituent algorithm used in the previous
iteration. Another way to implement a similar feature could be to introduce a probability
that the hybrid will select a random constituent algorithm rather than the previous one,
but that is beyond the scope of this thesis

3.2.3.2. Constituent Algorithm Implementation

Although the hybrid algorithm is simple, the proper incorporation of constituent
algorithms and maintenance of auxiliary vector populations, such as the velocity of PSO,
is not trivial. Algorithms like BST, DN3, and CKO do not utilize information from
previous iterations and can be treated as independent modules. The PSO and PRD
algorithms, however, utilize a velocity vector population that is a function of the previous
iteration’s velocity. As long as one of these modules is called sequentially, the velocity
vector can be computed normally. However, if the hybrid switches from PSO or PRD to
another method and then back to PSO or PRD, the velocity must be reinitialized. Since
the inertia term is a vital part of PSO, the manner in which this is done dramatically
affects its performance. As a first attempt, the velocity is re-initialized according to the
following equation,
where $V_j$ is the $j^{th}$ coordinate of the velocity vector, $D_{L,j}$ and $D_{U,j}$ are the lower and upper limits of the search domain in the $j^{th}$ direction respectively, $R$ is a uniformly distributed random number $\epsilon [0,1]$ and $g$ is the generation (iteration) number.

Additionally, algorithms with DE-style comparisons have an implicit form of elitism. These comparisons prohibit the superior design vectors like the global best design vector from being replaced by inferior design vectors. This feature is not present in algorithms like PSO. Therefore, if the hybrid algorithm switches from DE to PSO and back to DE it is possible to lose superior design vectors. Therefore, the hybrid stores the global best design vector separately, which partially remedies this problem. Nevertheless, the hybrid allows the other vectors to be overwritten by constituent algorithms.

3.2.3.3. Computational Expense

Since many real-world problems have objective functions that require hours to evaluate, generating the search vectors and $C_{CAS}$ is relatively computationally inexpensive. Evaluating them, in these cases, is not. However, if the objective function is inexpensive to evaluate (such as when surrogate models are used) the hybrid’s overhead becomes an important contributor to the overall computational expense. Accordingly, the hybrid was written using OpenMPI in order to benefit from a parallel computing environment.

The global optimization algorithms utilized in the hybrid are embarrassingly parallel. Therefore, in parallel environments, the populations are subdivided among nodes, reducing the time required for generating temporary populations. For problems of
low dimensionality, communication costs between nodes can exceed the costs related to the population size. When run on Tesla (in serial), the hybrid completes the average 2D problem (200 iterations) in roughly 2.4 seconds. On Tesla, in parallel (20 nodes), the communication costs increase this time to 6 seconds. One hundred dimensional problems, however, take on the order of hours to days in serial, but only minutes to hours in parallel.

3.3. Proposed Multi-Objective Extension of Hybrid

The notion of a search direction along a given topology is already well established. When the problem becomes multi-objective, and the objective function space grows in cardinality, the notion of a single search direction must change to accommodate the principles of Pareto optimality [8].

3.3.1. Multi-Objective Search Vectors

Some search vectors present in this hybrid, such as the population centroid, can be used in a multi-objective context without modification. Others, such as the population weighted average and negative of the global worst, must take on a new meaning and new formulation. In the context of multi-objective optimization, rankings no longer refer to single vectors, but rather, refer to sets of vectors. Non-dominated vectors have a rank of one (the Pareto-optimal set). Vectors dominated only by those in the Pareto Front have a rank of two, and the ranks increase in a similar fashion for inferior vectors. The negative of the global worst, therefore, would originate at the centroid of the worst set of vectors and pass through the centroid of the population. The formula for the population weighted average could take the form,
\[
\tilde{y}_{PWA} = \frac{1}{\sum_{i=1}^{B} \left( \frac{1}{i} \right)} \sum_{i=1}^{B} \left( \frac{1}{iA_i} \sum_{j=1}^{A_i} \tilde{x}_j \right) 
\]

where \( \tilde{y}_{PWA} \) represents the population weighted average, \( B \) represents the number of distinct ranked sets in the population (as well as the value of the rank itself), \( A_i \) represents the number of vectors in each set, and \( \tilde{x}_j \) is a design vector. Equation (3.17) is roughly equivalent to finding the centroid of each set, and then obtaining a weighted average of those vectors. The global best vector can be randomly selected from the Pareto Front or defined as the centroid of that set.

The projected global best vector can remain the same. However, it is also possible to capitalize on a feature unique to multi-objective optimization: the utopia vector (depicted in Figure 6 below).

![Figure 6: Utopia vector for two-objective minimization problem](image)

The utopia vector is a projection of the Pareto Front vectors whose values are the current global minimum for one of the objectives. Determining \( \tilde{x}_{\text{utopia}} \) from \( \tilde{x}_1 \) and \( \tilde{x}_2 \) is non-trivial. One way to approximate it is to identify the global worst vector (most dominated
point), and assume $\tilde{x}_{\text{utopia}}$ is located at the point that forms a parallelogram with the global worst, $\tilde{x}_1$, and $\tilde{x}_2$. Alternatively if a surface is constructed from the Pareto Front, as in [45], the utopia vector can be projected onto this surface, and that new point can be used as the projected global best vector.
CHAPTER 4

4. HYBRID ALGORITHM BENCHMARKING

As previously stated, the search vector hybrids presented in this thesis utilize six constituent algorithms. These six were selected from a pool of twenty possible algorithms: 11 distinct algorithms, and 9 modifications to those algorithms. All algorithms evaluated for this thesis are given in Table 4 below.

Table 4: Legend of Acronyms

<table>
<thead>
<tr>
<th>Acronym / Symbol</th>
<th>Algorithm</th>
<th>Modification</th>
</tr>
</thead>
<tbody>
<tr>
<td>PSO</td>
<td>Particle Swarm</td>
<td>n/a</td>
</tr>
<tr>
<td>PRD</td>
<td>Particle Swarm</td>
<td>With Random Differencing</td>
</tr>
<tr>
<td>QPS</td>
<td>Quantum Particle Swarm</td>
<td>n/a</td>
</tr>
<tr>
<td>MOP</td>
<td>Modified QPS</td>
<td>n/a</td>
</tr>
<tr>
<td>FFA</td>
<td>Firefly Algorithm</td>
<td>n/a</td>
</tr>
<tr>
<td>BAT</td>
<td>Bat Inspired Metaheuristic</td>
<td>n/a</td>
</tr>
<tr>
<td>CKO</td>
<td>Cuckoo Search</td>
<td>n/a</td>
</tr>
<tr>
<td>STD</td>
<td>DE - rand/1/bin</td>
<td>With randomized parameters</td>
</tr>
<tr>
<td>STD – R</td>
<td>DE - rand/1/bin</td>
<td>With randomized parameters, and sorted comparisons</td>
</tr>
<tr>
<td>STD – R,S</td>
<td>DE - rand/1/bin</td>
<td>With randomized parameters, and sorted comparisons</td>
</tr>
<tr>
<td>BST</td>
<td>DE - best/1/bin</td>
<td>n/a</td>
</tr>
<tr>
<td>BST – R</td>
<td>DE - best/1/bin</td>
<td>With randomized parameters</td>
</tr>
<tr>
<td>BST – R,S</td>
<td>DE - best/1/bin</td>
<td>With randomized parameters, and sorted comparisons</td>
</tr>
<tr>
<td>BST – SPC</td>
<td>DE - best/1/bin</td>
<td>With randomized parameters, sorted comparisons, special vectors and dynamic mutation</td>
</tr>
<tr>
<td>DN3</td>
<td>DE - Donor3</td>
<td>n/a</td>
</tr>
<tr>
<td>DN3 – R</td>
<td>DE - Donor3</td>
<td>With randomized parameters</td>
</tr>
<tr>
<td>DN3 – R,S</td>
<td>DE - Donor3</td>
<td>With randomized parameters, and sorted comparisons</td>
</tr>
<tr>
<td>TDE</td>
<td>Trigonometric DE</td>
<td>n/a</td>
</tr>
<tr>
<td>TDE – R</td>
<td>Trigonometric DE</td>
<td>With randomized parameters</td>
</tr>
<tr>
<td>GB</td>
<td>Blind Hybrid</td>
<td>n/a</td>
</tr>
<tr>
<td>PGB</td>
<td>Blind Hybrid</td>
<td>n/a</td>
</tr>
<tr>
<td>IBWA</td>
<td>Blind Hybrid</td>
<td>n/a</td>
</tr>
<tr>
<td>AV1</td>
<td>Blind Hybrid</td>
<td>n/a</td>
</tr>
<tr>
<td>PWA</td>
<td>Blind Hybrid</td>
<td>n/a</td>
</tr>
<tr>
<td>NGW</td>
<td>Blind Hybrid</td>
<td>n/a</td>
</tr>
<tr>
<td>DIR</td>
<td>Direct Comparison Hybrid</td>
<td>n/a</td>
</tr>
<tr>
<td>AUTO1</td>
<td>Automatic Switching Hybrid</td>
<td>n/a</td>
</tr>
<tr>
<td>AUTO2</td>
<td>Automatic Switching Hybrid</td>
<td>Passes sample point to constituent algorithms rather than global best</td>
</tr>
</tbody>
</table>

The Blind hybrids are named after the sample points that they use (see Chapter 3). Prior
to selection, the algorithms listed above were compared, and their relative performance is reported in the Appendix. Two-hundred and ninety-six of the Schittkowski & Hock test cases were used to benchmark the candidate algorithms, as well as the hybrids presented below.

4.1. Benchmark Case Set Up and Reporting

In order to compare the algorithms, each algorithm was executed for a certain number of iterations, and the final value of the objective function was used as the measure of performance. In order for this approach to be statistically meaningful, these numerical experiments must be repeated a certain number of times. Each numerical experiment is called a “trial.” For each trial, the optimization algorithm was executed for 200 iterations on a given Schittkowski & Hock test case.

4.1.1. Establishing Statistical Significance

The number of trials needed to establish a statistically significant result is not known a priori. In order to establish a guideline, a search vector hybrid was executed on the full set of test cases for 50, 60, 70, 80, and 90 trials. The mean accuracy and standard deviations were calculated and used to perform Student’s t-tests on each pair of trials. The table below contains a small sample of the data from the Student’s t-tests.

<table>
<thead>
<tr>
<th>TC</th>
<th>Dim</th>
<th>50 vs. 60</th>
<th>60 vs. 70</th>
<th>70 vs. 80</th>
<th>80 vs. 90</th>
</tr>
</thead>
<tbody>
<tr>
<td>1</td>
<td>2</td>
<td>0</td>
<td>0</td>
<td>0</td>
<td>0</td>
</tr>
<tr>
<td>2</td>
<td>2</td>
<td>0</td>
<td>0</td>
<td>0</td>
<td>0</td>
</tr>
<tr>
<td>3</td>
<td>2</td>
<td>1</td>
<td>0</td>
<td>0</td>
<td>0</td>
</tr>
<tr>
<td>4</td>
<td>2</td>
<td>1</td>
<td>0</td>
<td>0</td>
<td>0</td>
</tr>
<tr>
<td>5</td>
<td>2</td>
<td>1</td>
<td>0</td>
<td>0</td>
<td>0</td>
</tr>
<tr>
<td>6</td>
<td>2</td>
<td>0</td>
<td>0</td>
<td>0</td>
<td>0</td>
</tr>
<tr>
<td>7</td>
<td>2</td>
<td>1</td>
<td>1</td>
<td>0</td>
<td>0</td>
</tr>
<tr>
<td>8</td>
<td>2</td>
<td>1</td>
<td>0</td>
<td>0</td>
<td>0</td>
</tr>
</tbody>
</table>

Table 5: Sample of Student’s t-test Results
Table 5 shows the test case number (TC), the dimensionality of the problem (Dim), and the result of the t-tests for each pair of successive trial numbers. A zero represents cases when the null hypothesis is accepted at the 5% significance level, and a one represents cases when the null hypothesis is rejected. The hybrid performs the same in TCs 1, 2, and 6 for any number of trials. On TCs 3, 4, 5, and 8, however, the hybrid requires 60 trials before its performance converges statistically. For TC 7, the hybrid requires 70 trials before its performance converges. Since each trial is computationally expensive, we seek a trade-off between the statistical significance of the results, and the time required to generate them. Table 6 summarizes the results of the Student’s t-tests.

Table 6: Total Number of Hypothesis Rejections

<table>
<thead>
<tr>
<th></th>
<th>50 vs. 60</th>
<th>60 vs. 70</th>
<th>70 vs. 80</th>
<th>80 vs. 90</th>
</tr>
</thead>
<tbody>
<tr>
<td>50 vs. 60</td>
<td>61</td>
<td>16</td>
<td>16</td>
<td>17</td>
</tr>
</tbody>
</table>

There is a very clear gain in statistical significance by increasing the number of trials from 50 to 60. After that, however, there are no more significant gains. Therefore, this thesis will use 60 as the number of trials for benchmarking results. Oddly, the number of rejections does not monotonically decrease for the hybrid. Therefore, the tests were repeated using DE rand/1/bin with randomized parameters.

Table 7: Hypothesis Rejections – DE

<table>
<thead>
<tr>
<th></th>
<th>50 vs. 60</th>
<th>60 vs. 70</th>
<th>70 vs. 80</th>
<th>80 vs. 90</th>
</tr>
</thead>
<tbody>
<tr>
<td>50 vs. 60</td>
<td>43</td>
<td>13</td>
<td>15</td>
<td>13</td>
</tr>
</tbody>
</table>

Table 7 shows a similar trend, which suggests that the algorithm architecture does not significantly impact the result of the Student’s t-test.
Table 8: Sample of Student’s t-test Results

<table>
<thead>
<tr>
<th>TC</th>
<th>Dim</th>
<th>50 vs. 60</th>
<th>60 vs. 70</th>
<th>70 vs. 80</th>
<th>80 vs. 90</th>
</tr>
</thead>
<tbody>
<tr>
<td>18</td>
<td>2</td>
<td>1</td>
<td>0</td>
<td>0</td>
<td>0</td>
</tr>
<tr>
<td>19</td>
<td>2</td>
<td>0</td>
<td>0</td>
<td>0</td>
<td>0</td>
</tr>
<tr>
<td>20</td>
<td>2</td>
<td>0</td>
<td>1</td>
<td>0</td>
<td>0</td>
</tr>
<tr>
<td>21</td>
<td>2</td>
<td>1</td>
<td>0</td>
<td>0</td>
<td>0</td>
</tr>
<tr>
<td>22</td>
<td>2</td>
<td>0</td>
<td>0</td>
<td>0</td>
<td>0</td>
</tr>
<tr>
<td>23</td>
<td>2</td>
<td>0</td>
<td>0</td>
<td>0</td>
<td>1</td>
</tr>
<tr>
<td>18</td>
<td>2</td>
<td>1</td>
<td>0</td>
<td>0</td>
<td>0</td>
</tr>
<tr>
<td>19</td>
<td>2</td>
<td>0</td>
<td>0</td>
<td>0</td>
<td>0</td>
</tr>
</tbody>
</table>

Table 8 shows there is no clear trend from one pair of trials to another for certain test cases. This is true for DE as well as the hybrid used above. The common connection between algorithms is their random number generator. The random number generators create near-uniform distributions of points, and there is no guarantee that the distribution of points of the converged, final result will be normal.

Given that there is no monotonic trend in terms of population size, and inconsistent results from one pair of trials to another, it is likely that the normal distribution assumption of the Student’s t-test is being violated in some cases. Nevertheless, it appears that 60 trials are enough to establish a trend in performance.

4.1.2. Impact of Floating Point Arithmetic

Despite the fact that the global minimum is known, floating point arithmetic (FPA) introduces errors to the evaluation of the objective function. FPA error varies by operating system, and by possibly compiler code optimization. For example,

Table 9: Various Numerical Results by Operating System

<table>
<thead>
<tr>
<th>TC</th>
<th>Design Vector</th>
<th>Result</th>
<th>Operating System</th>
</tr>
</thead>
<tbody>
<tr>
<td>4</td>
<td>$X_1 = 0, X_2 = 1$</td>
<td>0</td>
<td>Windows 7 – 64 Bit</td>
</tr>
<tr>
<td>4</td>
<td>$X_1 = 0, X_2 = 1$</td>
<td>0</td>
<td>Ubuntu 12 – 32 Bit</td>
</tr>
<tr>
<td>4</td>
<td>$X_1 = 0, X_2 = 1$</td>
<td>-1.48102e-16</td>
<td>Rocks 5.1</td>
</tr>
</tbody>
</table>
While the result in Table 9 might seem trivial, such small differences in computational accuracy can determine whether or not a constraint violation penalty is applied, which ultimately affects the apparent accuracy of the algorithm. Complete FPA optimization is beyond the scope of this thesis. Nevertheless, steps were taken during post processing to partially address these issues. For example, any final result less than $10^{-20}$ or greater than $-1 \times 10^{15}$ was set equal to $10^{-20}$. Final results less than $-1 \times 10^{15}$ were considered erroneous if there was a difference between the C++ code and the published test cases. If not, they were considered to be an FPA error.

### 4.1.3. Computational Expense

Since objective function evaluations are typically the most computationally expensive part of the optimization process, they will be used as the metric of an algorithm’s speed. The number of objective function evaluations performed by an algorithm is fixed each iteration, therefore, the following multipliers can be used to convert from algorithm iterations to evaluations:

\[
\begin{align*}
\text{BST} & : N + 2 \text{ evaluations} \\
\text{CKO} & : 2*N \text{ evaluations} \\
\text{Blind Hybrids} & : N*(1 + \text{CKO call}) \text{ evaluations} + 2*\text{BST call} \\
\text{Direct Hybrid} & : N*(1 + \text{CKO call}) \text{ evaluations} + 2*\text{BST call} + 6 \\
\text{Auto Hybrids} & : N*(1 + \text{CKO call}) \text{ evaluations} + 2*\text{BST call} + 13 \\
\text{All Other Algorithms} & : N \text{ evaluations}
\end{align*}
\]

where $N$ is the population size, “CKO call” stands for the number of times the hybrid calls CKO to update the population, and, similarly, “BST call” stands for the number of times BST is called. Since CKO requires double the evaluations and yet performs like
PSO in many cases (see Appendix), Blind hybrids that use the minimum distance constituent algorithm selection method will check to see if there is a tie between PSO and CKO, and select PSO as the tie breaker.

4.2. Comparison of Blind Hybrid Optimization Algorithms

The Blind hybrids are the least computationally expensive hybrids created for this thesis. This benefit is balanced by the fact that they gather the least information from the search domain.

4.2.1. Comparison of Constituent Algorithm Selection Methods

Recall from Chapter 3 that for each sample point Blind hybrids used, there are three possible formulations for constituent algorithm selection: minimum distance, maximum dot product, and minimum, scaled distance. The goal of the Blind hybrid is to utilize search vectors to enhance the optimization process without requiring additional objective function evaluations. A comparison of Blind hybrid performance for each constituent algorithm selection type is given in Figure 7. The Blind hybrids are scored based on having the best accuracy and fastest convergence rate (calculated using objective function evaluations) for each test case. The minimum distance selection method performs at least 50% better than other methods for every sample point. Since it is also the simplest method to compute, it is used again for the Auto hybrids. It is worth noting that the minimum distance method works best with sample points that are not based on the main population (PGB, AV1, IBWA, and NGW).
This suggests that search directions derived directly from the population (using its members or an average of its members) provide a slightly better indication of the direction in which to search, whereas the more exotic sample points provide better information regarding the local neighborhood in which to search, relative to the search vector in use.

4.2.2. Constituent Algorithm Selection Preference

The Blind hybrid strategy provides useful insight into potential correlations.
between search vectors, optimization algorithms, and objective function topologies. For example, the PWA Blind hybrid selects CKO at least one order of magnitude more frequently than any other algorithm (see Figure 8). Furthermore, in 33 test cases CKO search is the only constituent algorithm selected by the BWA Blind hybrid.

CKO-Only Test Cases:

**Figure 8: CKO Selection Preference of PWA Blind hybrid**

There is an approximate pattern in the location of the global minima of the above-referenced test cases. They tend to be near/at the edge of the domain, the center of the domain, or in some pattern (such as a sequence of integers that increase with dimension), which suggests CKO seeks symmetries in the problem. However, as a reminder that correlation is not causation, it must be noted that there is no clear pattern in the topologies of the above-mentioned test cases. Therefore, the interaction between CKO and the topologies represented in this set is more involved than mere symmetries.

Similar to the PWA Blind hybrid, the IBWA Blind hybrid also selects CKO over other methods. However, it selects DN3-R with similar frequency (FFA is not used here to reduce computational expense). Furthermore, the IBWA Blind hybrid selects only CKO and DN3-R in 43 test cases (see Figure 9).
CKO & DN3-R Test Cases:

Figure 9: CKO & DN3 Selection Preference of IBWA Blind hybrid

Therefore, it appears that in a wide range of test cases, CKO and DN3 tend to search near the weighted average of the population or weighted average of the individual best population. Particularly in the test cases 101, 102, 103, 104, 234, 236, 239, 292, 293, 298, 299, 301, 302, 303, 304, 305, 379, 380, 381, 382, 383, 384, 388, 391, and 394, which make up the intersection of the two sets presented above.

Figure 10: Selection Preferences of GB and PGB Blind hybrids

The GB and PGB Blind hybrids clearly prefer BST-SPC over other methods. This is no coincidence since all forms of BST use the global best as the design vector to be perturbed. After BST-SPC, the hybrids select PSO, which also uses the global best design vector in its formulation. Unlike IBWA, or PWA, the PGB Blind hybrid rarely relies on
one or two optimization algorithms for a given topology. In fact, there are only four cases where the PGB Blind hybrid predominantly selects a subset of constituent algorithms: TC 45, 234, 236, and 239. In these cases, the hybrid selects two or more of BST-SPC, PSO, PRD, and MQP.

Similar to the PGB Blind hybrid, the NGW and AV1 Blind hybrids rarely select a subset of constituent algorithms for a given topology. The NGW and AV1 blind show only a marginal preference for PSO (Figure 11).

Recall that the AV1 sample point is an average that includes the NGW. The lack of selection preference suggests the NGW frequently moves across the domain so that there is no clear trend. The six sample points discussed here were also used in the Auto hybrid.

4.2.3. Blind Hybrid Performance

Although their primary purpose is to examine relationships between search vectors and constituent algorithms, these hybrids were compared to their constituent algorithms in order to determine the effectiveness of their switching logic.
4.2.3.1. Hybrid vs. Constituent Algorithms

Figure 12 shows the results of comparing Blind hybrids to one-another and their constituent algorithms. Interestingly, the GB hybrid is the most accurate and fastest in nearly 18% of the test cases. If speed is not considered, it is as accurate as or better than other optimization algorithms in over 23% of the test cases. From this, and the overwhelming success of BST-SPC, it can be concluded that a strategy as simple as moving the population toward the global best each iteration is enough to perform well in one-fifth of the Schittkowski & Hock test cases.

![Best Speed and Accuracy](image)

Figure 12: Relative Performance of Blind hybrids and their Constituent algorithms

None of the Blind hybrids outperform BST-SPC in any metric (speed, robustness, etc.) over the set of test cases because the Blind hybrid switching strategy is too simplistic to be robust. However, there are specific cases, such as TC 110, where the hybrid outperforms its constituent algorithms.
The GB Blind hybrid, which selects BST-SPC over 10 times more frequently than any other method, converges faster than all other optimization algorithms considered, although it is only slightly faster than BST-SPC (Figure 13).

The IBWA Blind hybrid, which only alternates between CKO and DN3, converges faster than CKO alone (Figure 14), but much slower than DN3-R, which converges in a manner similar to BST-SPC but requires 9,000 objective function evaluations to do so.

Special cases such as these raise the question: under what conditions can a hybrid be considered “good” when compared to its constituent algorithms? This topic will be discussed in Section 4.5.
4.2.3.2. Hybrid vs. Hybrid

Figure 15 shows the results of comparing Blind hybrids to one-another, using the minimum distance formulation.

![Figure 15: Relative Performance of Blind hybrids](image)

It is clear that the two most computationally efficient search methods, which account for the best results in roughly 82% of all test cases, are the GB and PGB search vectors. Once again, the simpler formulations are usually the best. It is worth noting that the class of test cases for which the PGB Blind hybrid is the best does not intersect with other hybrids. That is, if speed is not taken into consideration, it will still perform well in exactly 35.14% of test cases.

4.3. Comparison of All Search Vector Hybrids

The three remaining hybrids are DIR, AUTO1, and AUTO2. Unlike AUTO1, the AUTO2 hybrid passes the sample point to the constituent algorithm in lieu of the global best vector (Table 4). These hybrids were designed to improve upon the performance of the Blind hybrids by using additional information about the objective function topology.
4.3.1. Revisiting Constituent Algorithm Selection Preference

Like the GB and PGB Blind hybrids, AUTO1 and AUTO2 select BST-SPC more frequently than other constituent algorithms, but MQP is selected more often than PSO.

AUTO2 utilizes a single constituent algorithm in only two test cases: TC 234, and 239. In fifteen other test cases, it may rely on three or four out of the six constituent algorithms. These test cases are: TC 4, 17, 33, 45, 95, 96, 221, 226, 236, 238, 262, 278, 279, 353, and 366. For the remaining test cases, however, AUTO2 uses all six constituent algorithms.

The DIR hybrid has a trend very similar to the Auto hybrids, but calls BST-SPC less frequently. The striking similarity of the selection preferences is clearly due to the inclusion of $C_{CAS}$, and the repeated use of the GB sample point by the Auto hybrids.
4.3.2. Search Vector Hybrid Performance

4.3.2.1. Hybrid vs. Constituent Algorithms

Comparing DIR, AUTO1, and AUTO2 against their constituent algorithms, AUTO2 is clearly the dominant hybrid, but BST-SPC is still the most robust algorithm.

![Best Speed and Accuracy](image)

AUTO2 and BST-SPC could be considered ideal for different classes of test cases, despite the fact that AUTO2 utilizes BST-SPC. See section 4.5 for a more detailed discussion of this topic.

In addition to accuracy and speed, another indicator of an algorithms’ performance is the frequency with which it converges to a local minimum. Since all of the algorithms presented here converge to a local minimum in one problem or another, it is instructive to count the number of times an algorithm produces the single worst result for a given test case.
Figure 19: Worst Performance of DIR, AUTO1, AUTO2, and their Constituent Algorithms

Figure 19 shows that the AUTO2 hybrid and BST-SPC are least likely to produce the worst result. Since AUTO2 incorporates constituents including BST-SPC, this demonstrates that the AUTO2 switching logic dramatically reduces the likelihood of producing a terrible result in spite of its constituents. On the other hand, the DIR hybrid still possesses a fairly high likelihood of yielding a poor result, indicating that switching search directions and constituents is a better strategy than switching based only on $C_{CAS}$.

If the convergence rate is calculated based on iterations rather than objective function evaluations, BST-SPC and AUTO2 switch roles. Situations like this might arise when surrogate models are used in lieu of objective function evaluations, greatly reducing the computational expense of the optimization process. In these cases, AUTO2 is the more robust algorithm.
4.3.2.2. Hybrid vs. Hybrid

Compared to the Blind hybrids, the DIR, AUTO1, and AUTO2 hybrids converge faster, are more accurate, and more robust (Figure 21).

AUTO2, in particular, is faster and more accurate than any other hybrid in over half of the test cases. AUTO2 obtains the single worst answer in 6.08% of test cases, superseded only by the AV1 Min Dist hybrid at 4.73%.
Comparing DIR, AUTO1, and AUTO2 exclusively, AUTO2 again significantly outperforms the other hybrids, reinforcing the conclusion that automatically switching search directions and constituents is better than switching based solely on $C_{\text{CAS}}$.

4.3.2.3. Hybrid vs. All

Figure 23 shows the relative performance of every optimization algorithm used in this thesis except Blind hybrids using the maximum dot product and minimum scaled
distance formulations. The four dominant methods in accuracy and speed are BST-R, BST-R,S, BST-SPC and AUTO2. AUTO2 obtains the single worst result in only 1.35% of test cases. The remaining 25 methods are well suited to very small percentages of test cases.

If we again relax the assumption that the objective function evaluation dominates the computational expense of the optimization process, then the AUTO2 hybrid becomes the most robust algorithm. If speed is not taken into consideration, then within 200 iterations AUTO2 obtains the best accuracy in 46.6% of test cases.

4.3.2.4. Feasibility of Results

Due to the use of an exterior penalty function, the accuracy of the hybrid includes the penalty of the constraint violation. Since the penalty function alters the topology of the objective function, it can mislead the hybrid (as well as the other optimization algorithms described in this thesis), causing it to converge to a local minimum outside of the feasible space. The results in Table 10 reflect the probability that the solution will be
infeasible after 200 iterations.

<table>
<thead>
<tr>
<th>Number of Constraints</th>
<th>Number of Test Cases</th>
<th>Mean Probability of Constraint Violation</th>
<th>Standard Deviation</th>
</tr>
</thead>
<tbody>
<tr>
<td>1</td>
<td>67</td>
<td>55.97%</td>
<td>49.65%</td>
</tr>
<tr>
<td>2</td>
<td>49</td>
<td>33.63%</td>
<td>36.42%</td>
</tr>
<tr>
<td>3</td>
<td>30</td>
<td>51.40%</td>
<td>39.23%</td>
</tr>
<tr>
<td>4</td>
<td>18</td>
<td>39.59%</td>
<td>34.58%</td>
</tr>
<tr>
<td>5</td>
<td>10</td>
<td>32.04%</td>
<td>28.79%</td>
</tr>
<tr>
<td>6</td>
<td>10</td>
<td>35.83%</td>
<td>37.11%</td>
</tr>
<tr>
<td>≥ 8</td>
<td>26</td>
<td>31.25%</td>
<td>26.27%</td>
</tr>
</tbody>
</table>

The standard deviations tend to be large because in some test cases the converged solution is infeasible for all trials, while in others the solution is always feasible (particularly for cases with a low number of constraints). The convergence trend suggests that 200 iterations may not be enough to locate a feasible solution for some test cases. Over the set of all trials and all test cases, the hybrid has a 39.01% chance of obtaining a feasible solution (no constraint violations). This performance is typical of unconstrained methods [7], and strongly indicates that a better approach to constraint enforcement is required.

4.3.3. Impact of Problem Dimensionality

AUTO2 clearly outperforms other optimization algorithms in higher dimensional test cases. Like other algorithms, the number of objective function evaluations required to locate the global minimum increases, but to a lesser degree than other algorithms. In the plots below, HOA stands for “Hybrid Optimization Algorithm.” The rapid drop in error shown in Figure 25 during the first few objective function evaluations is due to the use of the PC sample point.
Although AUTO2 does not locate the global minimum of TC 302 or TC 305, it outperforms all other optimization algorithms in accuracy and speed (see Appendix). This suggests one of three things: (1) new search vectors may be needed to speed up convergence, (2) new constituent algorithms may be useful, or (3) it might be necessary to incorporate a new operation (such as the random generation of new points, or a new selection method) to improve the hybrid’s robustness with increasing problem dimensionality.
Figure 26 shows the change in hybrid performance with increasing dimensionality. It clearly outperforms all other optimization algorithms in this thesis (see Appendix), but again, this is partially due to the hybrid’s ability to utilize the PC as a sample point. Problems lacking this form of symmetry might show an inferior trend.

4.4. Comparison of Auto Hybrid to Non-Search Vector Hybrids

Researchers at MAIDROC have compared previously developed hybrids to the commercial optimization program IOSO [46]. Figures 27 - Figure 34 present the performance of the AUTO2 hybrid (its convergence history and constituent algorithm selection histograms) alongside the convergence history of these hybrids on a set of six test cases. The figures showing performance for hybrids other than AUTO2 are taken from [46]. The number of trials used to generate the results given in [46], and whether or
not these plots are an average of those trials, or simply the best result from all trials is unknown. The AUTO2 results (excluding Figure 29) are the arithmetic mean of 60 convergence histories.

Results for the 4-dimensional Levy9 test case are shown in Figure 27. The AUTO2 hybrid roughly matches the Dulikravich, Martin, Colaço (DMC) hybrid’s performance at the first 2,000 objective function evaluations, and then rapidly surpasses it. The Colaço BFGS-based optimizer outperforms both in terms of speed, but appears to converge to a local minimum. IOSO is much more efficient, but also incorporates highly advanced surrogate modeling techniques. The surrogate model evaluations are not included in IOSO’s convergence history.
Only 17 of the 60 trials AUTO2 hybrid trials converged to the global minimum (Figure 28), and the DMC hybrid converges to similar local minima. In these 17 trials (Figure 29), the hybrid locates the global minimum to single-precision accuracy within roughly 2,250 objective function evaluations. The performance of AUTO2 relative to the other optimizers is approximately unchanged. Figure 29 uses the “average of the log” of the convergence histories (rather than the “log of the average,” or, logarithm of the arithmetic mean), given by Eq. 4.7, below.
\[ \tilde{H}^* = \frac{1}{N} \sum_{t=1}^{N} \log_{10} (\tilde{H}_t) \]  

(4.7)

where \( \tilde{H}_t \) is the vector representing the convergence history for a given trial, \( t \), and \( \tilde{H}^* \) is the “average of the log” of the \( N \) convergence histories (\( N \) trials). This representation was selected because it reflects the change in order of magnitude of several convergence histories better than a simple arithmetic mean.

Figure 30: Convergence History Comparison – Levy

The AUTO2 hybrid converges to the global minimum of the Levy function (Figure 30) in less than 3,000 objective function evaluations. No results were reported for IOSO. This Levy function is 7-dimensional, and varies slightly in form from the Levy9 function. This is an excellent illustration of the ANFL theorem’s claim that a small change in an objective function can cause a big change in an algorithm’s performance.
The AUTO2 hybrid converges to the global minimum of the Rosen function (Figure 31) within 300 objective function evaluations, which is on par with the MAIDROC hybrids.
Figure 32: Convergence History Comparison – Schwefel

Figure 32 illustrates a case in which the AUTO2 hybrid, and coincidentally IOSO, converges to a local minimum. Only 34 of the 60 trials converged to the global minimum. As with the Griewank function, this is probably due to AUTO2’s aggressive search strategy (see discussion in Section 4.5.2). When AUTO2 does converge to the global minimum, it generally does so in fewer than 500 objective function evaluations, as shown in Figure 33.
In Figure 34, the AUTO2 hybrid converges to the global minimum in less than 2,000 objective function evaluations, and does so monotonically. While monotonic convergence can be practical because it is predictable, it is no guarantee of superiority (recall the ANFL theorem). In fact, algorithms like SA and GA (without elitism) rely on hill-climbing mechanisms to perform global design space exploration. Nevertheless, AUTO2’s convergence rate is on par with the other MAIDROC hybrids, and retains its order-of-magnitude difference with IOSO typical of the test cases presented here.
AUTO2 was also compared with a hybrid developed at MAIDROC called OPTRAN. Different versions of OPTRAN exist. The one used here is the 4th generation of MAIDROC hybrids discussed in Section 2.3.3. The following steps were taken to minimize the bias in their comparison:

1. OPTRAN does not use a penalty function for constraint enforcement, therefore, the unconstrained subset of the Schittkowski & Hock test cases were used.
2. Each optimizer has a different population initialization technique. OPTRAN possesses the built-in capability to read in a single design, while AUTO2 can read in an entire population. Therefore, OPTRAN was provided with the global best design from the same initial population used by AUTO2.

On an objective-function-evaluation basis, AUTO2 dramatically outperforms OPTRAN (Figure 35). This is important because, unlike AUTO2, OPTRAN is a global-local hybrid. Therefore, this performance suggests that OPTRAN often converges to local minima that the AUTO2 hybrid can avoid.
4.4.1. Comparison of Execution Times

Although using objective function evaluations is a practical, implementation-agnostic, platform-agnostic measurement of an algorithm’s speed, another common practice is to compare their execution times. This method provides some indication of how an algorithm performs when the computational cost of evaluating the objective function is relatively low. Unfortunately, this approach can be biased by a large number of factors. Some biasing factors relevant to this thesis include:

- Bias due to the method of implementation
  - OPTRAN, and the Auto hybrid have different input/output (I/O) subroutines. They store different numbers of files, and different amounts of data within those files. Furthermore, the Auto hybrid performs some basic file clean up tasks (compressing and deleting files) once the optimization is complete. Also, OPTRAN prints a substantial amount of text to standard output, while the Auto hybrid does not. For the purpose of this thesis, OPTRAN’s standard output was redirected to a file (these files reached hundreds of megabytes for 100-dimensional problems).
  - All search vector hybrids presented in this thesis run in parallel. In order to minimize bias, the time comparisons are performed using a single processor which causes the MPI function calls to unnecessarily increase the Auto hybrid’s computational expense.
  - OPTRAN was written in FORTRAN, while the other algorithms were written in C++. The two languages handle tasks such as file I/O and passing...
variables to functions in very different manners, all of which impact performance.

- OPTRAN is used in conjunction with a C++ version of the Schittkowski & Hock test cases, and so must be cross compiled (FORTRAN and C++). Furthermore, the Auto hybrid and the PSO implementation used for this section use a compiler containing wrappers for MPI. Therefore, OPTRAN uses gfortran and g++, while the Auto hybrid and PSO use mpic++ (a wrapper for g++).

- Bias due to time measurement approach
  - In addition to differences between languages, within C++ there are several ways to measure time, each with significant consequences. For the purpose of this thesis it suffices to say that each optimization algorithm executable is timed from the beginning of its execution until the end using the high resolution clock in the <chrono> library with millisecond accuracy. The interested reader can learn more about timing from [47] [48] [49].
  - Although OPTRAN contains a timing subroutine, it is not active in the version used for this thesis, therefore, the method stated above is used.

An algorithm’s computational expense also depends on the skill of the programmer, and the number of processes being executed in the background by the operating system at the time of testing. Setting all these factors aside, an algorithm’s execution time will also vary from system to system. In this thesis, the algorithms were executed on an Ubuntu 13.10 virtual machine (Parallels 9) with 8GB of dedicated memory and 4 available cores. The virtual machine is hosted on a MacBook Pro (OS version 10.8.5) with a 2.8GHz Intel
Core i7 processor, 16GB 1600MHz DDR3 RAM, and a solid-state hard drive. Recall that the algorithms were executed on a single processor.

Table 11: Time Required for Convergence - OPTRAN, AUTO2, and PSO

<table>
<thead>
<tr>
<th>TC</th>
<th>Dim</th>
<th>Algorithm</th>
<th>Times (s)</th>
</tr>
</thead>
<tbody>
<tr>
<td></td>
<td></td>
<td></td>
<td>Trial 1</td>
</tr>
<tr>
<td>300</td>
<td>20</td>
<td>OPTRAN</td>
<td>4.065</td>
</tr>
<tr>
<td></td>
<td></td>
<td>AUTO2</td>
<td>8.578</td>
</tr>
<tr>
<td></td>
<td></td>
<td>PSO</td>
<td>0.289</td>
</tr>
<tr>
<td>301</td>
<td>50</td>
<td>OPTRAN</td>
<td>54.858</td>
</tr>
<tr>
<td></td>
<td></td>
<td>AUTO2</td>
<td>34.660</td>
</tr>
<tr>
<td></td>
<td></td>
<td>PSO</td>
<td>0.973</td>
</tr>
<tr>
<td>302</td>
<td>100</td>
<td>OPTRAN</td>
<td>495.488</td>
</tr>
<tr>
<td></td>
<td></td>
<td>AUTO2</td>
<td>94.454</td>
</tr>
<tr>
<td></td>
<td></td>
<td>PSO</td>
<td>2.585</td>
</tr>
</tbody>
</table>

Table 11 shows that OPTRAN’s required computational time increases by roughly one order of magnitude with each increase in problem dimensionality. This is likely due to the vast amounts of output OPTRAN generates. OPTRAN’s results vary significantly due to its various convergence criteria, which do not trigger at the same time: (1) reaching the maximum number of iterations, (2) reaching the maximum number of objective evaluations, (3) failing to decrease the objective function after executing every constituent algorithm [50].
Figure 36: Convergence Histories over Time (a) AUTO2 (b) OPTRAN
AUTO2, on the other hand, runs for a fixed number of iterations (here, 200). AUTO2 clearly outperforms OPTRAN for test cases with higher dimensionality in terms of execution time. The slight variability comes in the number of objective function evaluations per iteration as discussed in Section 4.1.3, Eq. 4.5. PSO’s computational cost is roughly one-thirtieth that of AUTO2 due to its simplicity.

Figure 36 shows the convergence history of OPTRAN and AUTO2 over time. In this test case (as well as TC300 and TC302) OPTRAN converges to a result more accurate than AUTO2. The smooth convergence profile occurring after the first ten seconds is probably the result of its use of the DFP constituent algorithm. This is an example of the advantage of global-local hybridization.

4.5. Discussion of Results

4.5.1. Characterizing Hybrid Performance

Having benchmarked the algorithms, the Auto hybrid is clearly a competitive alternative to other optimization algorithms. While this may be sufficient to establish a case for its use, there is one question that benchmarking alone cannot answer:

What makes a hybrid “good”?

Recall from [22] that any algorithm that efficiently solves a problem does so because it implicitly contains a heuristic that is compatible with the objective function’s topology. In the case of a pure optimization algorithm, metrics such as speed and accuracy provide an adequate characterization of this compatibility. Since a single algorithm cannot be ideally suited for all problems, an implicit goal of hybrids with switching mechanisms is to correctly match an objective function topology to a compatible constituent algorithm. Therefore, the true measure of a hybrid’s quality is not merely its speed and accuracy, but
also its ability to identify the constituent algorithm required to best solve a problem. In light of the “effective topology” concept presented earlier, the quality of the search vector hybrids is determined by its ability to perform this match during each iteration of the optimization process. Thus, a new metric for quality is introduced: the probability of selecting the most compatible constituent algorithm for a given objective function.

This new metric naturally raises another question: Over the set of all optimization problems, what is the upper limit of a hybrid’s performance? In order to answer this question, let us say that there are two optimization algorithms C₁ and C₂ that are ideally suited for two disjoint sets of objective functions A and B, respectively. The sets of objective function topologies represented by A and B are denoted Tₐ and Tₐ respectively. Let set D be a set of objective functions disjoint with respect to A and B, containing objective function topologies, Tᵋ, that are distinct from, and not reducible to combinations of Tₐ and Tₐ. An example of a function whose topology is reducible to a combination of others is the sine function, which can be thought of as an infinite series of piecewise convex and concave sections. Another (impractical) example would be to consider all continuous functions to be an infinite series of infinitesimal linear sections. Let U be the set of all objective functions, as shown in Figure 37 below:
Now say there is a hybrid optimization algorithm $H$ that automatically switches between $C_1$ and $C_2$ and is ideally suited to solve the set of objective functions $X$. In the worst case scenario, the set $X$ will be a proper subset of $A$ and/or $B$ (Figure 38). It is likely, however, that $X$ will be a subset of $A$, $B$, and $D$ (Figure 39), as a natural consequence of the ANFL theorem. This will be referred to as “typical” performance.

The typical case is drawn deliberately to suggest that the hybrid converges as quickly and accurately as $C_1$ and $C_2$ wherever $X$ intersects $A$ and $B$ respectively, because the term “ideally suited” should not be construed to mean that an algorithm’s superiority in one class of problems is mutually exclusive with all other algorithms. In fact, in practice there would be no guarantee that $C_1$ and $C_2$ are ideally suited for disjoint sets, which could adversely affect the hybrid’s performance.
A hybrid with “satisfactory” performance is one in which X is the union of A and B. Now, consider a set of objective functions, E, whose topologies can be decomposed into some combination of topologies in $T_A$ and $T_B$. A hybrid is considered to have “ideal” performance if X is the union of A, B, and E as shown in Figure 40.

![Figure 40: “Ideal” Performance](image)

A hybrid with “utopian” performance is one in which X is the union of A, B, D, and E.

Although it is trivial to stipulate that such levels of performance are attainable, in practice it may not be possible to create an ideal or utopian hybrid. It is obvious that a simple global-local hybrid, which allows its global algorithm to converge and then passes the best solution to the local algorithm, can have satisfactory performance. The precise condition utilized in such hybrids could cause its performance to deviate from satisfactory, but this is easy to remedy with some experimentation. The higher levels of performance, however, are more difficult to obtain.
4.5.2. Search Vector Hybrid Characterization

The premise of this thesis is that given the existence of a set E (described above), there exists a hybrid that can match its constituents to a series of effective topologies such that its performance is ideal. Underlying this statement is the assumption that the sequence of effective topologies the hybrid encounters during the optimization process can correctly represent the topologies into which the problem should be decomposed. Since the quality of the effective topology depends on the population size and distribution, it is easy to imagine circumstances in which the assumption may be invalid. Nevertheless, the validity of the premise alone is not sufficient to guarantee that the hybrid’s performance will be ideal. Consider a case where PSO is the single best constituent algorithm for a given problem, and the Auto hybrid consistently selects PSO. Although the hybrid is making the correct decision, it will always converge more slowly than PSO, on an objective function evaluation basis, due to the additional computational expense of evaluating the sample points and $C_{CAS}$.

In this sense, the Auto hybrids have typical performance by definition, except that here typical performance includes a subset of E. If the Auto hybrid utilized $C_1$ and $C_2$, one could say it exchanges convergence speed in subsets of A and B for improved overall robustness (the ability to solve problems in D and E). In order for a hybrid to have ideal performance, it would have to switch between algorithms without requiring any additional objective function evaluations. Alternatively, if the hybrid algorithm uses surrogate models to perform its constituent selection operations, and if the surrogate model adequately represents the objective function, and if the computational expense of evaluating the surrogate is negligible compared to that of the objective function, then
ideal performance is also attainable. Given the strict requirements for ideal performance, utopian performance is clearly challenging to obtain, even if D only contains one objective function.

The statement that a particular algorithm is ideally suited for a test case is relative. A hybrid’s performance is relative to its constituents, and its constituent’s performance is relative to all other optimization algorithms. Since only some optimization algorithms were examined for this thesis, there is epistemic uncertainty in the evaluation of a hybrid using test cases. Consider a hybrid that replaced its first constituent algorithm C₁ with a new algorithm C₃. It is then possible that objective function topologies in A could move to E, and the hybrid’s performance would appear to change relative to C₁.

Figure 41 below, shows the percentage of problems for which the AUTO2 hybrid and its constituents are ideally suited in terms of performance. Based on this, one can say that the AUTO2 hybrid correctly selects among constituents in 38.46% of the test cases.

![Best Speed and Accuracy](image)

Figure 41: Relative Performance of Auto2 vs. its Constituents

However, it is important to analyze the quality of the Auto hybrid’s search logic independent of its performance. Therefore, Table 12 below shows the probability that the
hybrid will select the most compatible constituent algorithm in the test cases for which it is not ideally suited (i.e. the remaining 61.54% of cases in which it is not the fastest). In Table 12, the hybrid is assumed to have selected the correct constituent algorithm (i.e. its search logic is valid) if it selects the algorithm identified in Figure 41 as being ideally suited for that given test case.

Table 12: Probability of Correct Constituent Selection

<table>
<thead>
<tr>
<th>Correct Constituent</th>
<th>Mean Probability of Selection</th>
<th>Standard Deviation</th>
</tr>
</thead>
<tbody>
<tr>
<td>BST-SPC</td>
<td>90.28%</td>
<td>2.96%</td>
</tr>
<tr>
<td>DN3</td>
<td>1.74%</td>
<td>0.37%</td>
</tr>
<tr>
<td>PSO</td>
<td>1.49%</td>
<td>1.03%</td>
</tr>
<tr>
<td>PRD</td>
<td>1.96%</td>
<td>1.04%</td>
</tr>
<tr>
<td>MQP</td>
<td>2.79%</td>
<td>1.08%</td>
</tr>
<tr>
<td>CKO</td>
<td>2.60%</td>
<td>1.78%</td>
</tr>
</tbody>
</table>

Overall, there is a 79.86% chance that the hybrid will select the correct constituent algorithm. However, there is a strong selection preference for BST-SPC, even in problems where other constituent algorithms are the correct choice. Although this is likely due to AUTO2 passing the search vector to BST-SPC for use as its global best, it also suggests one or both of the following:

1. Search vectors alone do not provide enough information to characterize the effective topology.

2. The aggressive search strategy employed by this hybrid favors local exploitation over global exploration.

Recall that the aggressive search strategy (see Section 3.2.3.1) stipulates that a good match between $C_{CA}$ and the minimum sample point implies a good match between the constituent algorithm and the effective topology. This is reasonable for functions that have clear global
trends, but can be misleading for objective functions that have large, apparently favorable neighborhoods located far from one another within the design space.

Given the high probability of correct constituent algorithm selection and the hybrid’s competitive performance, automatic switching between search vectors appears to be the beginning of a good foundation for the development of a hybrid that operates on continuous domains. Including statistical measures of the objective function’s distribution, as well as the population’s distribution over the course of the hybrid’s convergence, could help it detect the presence of misleading neighborhoods and other important topology features. This information could be used to probabilistically select sample points and constituent algorithms. Furthermore, it is possible to stop evaluating C_{CAS}, and simply use their proximity to the sample point as part of the selection mechanism. This would reduce the hybrid’s computational expense and bring its performance closer to ideal.
CHAPTER 5
5. OPTIMIZATION OF THE ASYNCHRONOUS PULSE UNIT

During World War II and the Cold War, the United States built a number of nuclear weapon research and manufacturing facilities across the country. These facilities produced massive amounts of high-level waste (HLW), which were stored in large single and double-shelled tanks. One such site, called the Hanford site, [51] contains:

“more than 53 million gallons of radioactive and chemically hazardous waste in 177 underground storage tanks, 2,300 tons (2,100 metric tons) of spent nuclear fuel, 9 tons (8 metric tons) of plutonium in various forms, about 25 million cubic feet (750,000 cubic meters) of buried or stored solid waste, and groundwater contaminated above drinking water standards, spread out over about 80 square miles (208 square kilometers), more than 1,700 waste sites, and about 500 contaminated facilities” [52].

As part of the cleanup process, some of this waste is transported through underground pipelines that can run over 8 miles in length [53]. Sometimes these pipelines become clogged, which leads to costly delays in waste removal [54] as well as environmental concerns.

Since 2007, the Applied Research Center (ARC) at Florida International University (FIU) has been testing and evaluating the efficacy of several pipeline-unplugging technologies including an erosive-wave-action technology by NuVision Engineering, and sonic resonance system by AIMM Technologies [55]. Additionally, FIU has proposed and developed two technologies: a pneumatic crawler [56], and a pressure-pulse system called the Asynchronous Pulse Unit (APU) [57].
In order to test the APU, the researchers at ARC have constructed a number of small-scale pipelines, and clogged them using non-radioactive simulant blockages made from a variety of materials. These experiments are time-consuming to set up, and limited in the data that can be extrapolated from them. Therefore, ARC has chosen to supplement these experiments with numerical simulations using software developed in-house [53] as well as commercially available computational fluid dynamics (CFD) software [58]. Although efforts to simulate the APU using the commercial program ANSYS Fluent at ARC are ongoing, they are beyond the scope of this thesis.

This chapter briefly reviews one pipeline simulation program developed at ARC, as well as a similar program presented in [59]. It then presents a new pipeline analysis program developed for this thesis. The software is verified, and validated. The operation of the APU is then cast as an optimization problem. Finally, the new software is coupled to the hybrid optimization algorithm presented in Chapter 3, and the optimization problem is solved.

5.1. Pipeline Simulation Software

5.1.1. Existing Codes

In 2012, researchers at ARC modified the Method of Characteristics (MOC) described in [60] to serve as a simplified computer model of the experimental set up. This method reduces the governing partial differential equations to a set of ordinary differential equations, whose solution can then be approximated. The interested reader is referred to [53] and [60] for a complete discussion. The so-called “Modified Method of Characteristics” (MMOC) is used in [53] to characterize pipelines, model their transient behavior, and optimize the operation of the piston pumps acting on the pipeline. Going
forward, this code will be referred to as “Wood’s code,” after its author. It will be used to verify the results of the code developed for this thesis.

Recently, Dr. Mambretti of the State University of Campinas in Brazil published a textbook on the topic of water hammer simulations [59]. The computer codes provided with this textbook will also be used to verify the code developed for this thesis. Going forward, this will be referred to as the “WHS code,” which stands for “Water Hammer Simulation” after the title of the book.

5.1.2. Updated Method of Characteristics

An MOC code based on the updated version of Wylie’s textbook [1] was written for this thesis. The new textbook contains several changes to the numerical procedure in [53]. The most notable changes are:

- The use of a staggered grid: nodes are now evaluated every other time step, beginning with odd numbered nodes during the first time step. This requires that each pipeline have an even number of “reaches” (mesh elements).
- The final forms of the equations for various pipeline elements have changed.

Furthermore, the code developed for this thesis differs from the code in [53] in a number of ways:

- It does not use interpolations in the numerical solution.
- It automatically adjusts the time step such that the Courant Condition is always satisfied.
- It is capable of modeling tee fittings, with and without air.
- It does not utilize geometric friction coefficients.
• New boundary conditions are proposed (see Section 5.1.2.5). Furthermore, it can
utilize either mass-flow or pressure boundary conditions.

Going forward, the MOC code developed for this thesis will be referred to as “the MOC
code.” The derivations of various equations used in the MOC code are presented in the
subsections that follow.

5.1.2.1. Material Properties

The pipeline is divided into an even number of pipe sections of length $dx$. Although the pressurization of the pipeline does cause the pipes to expand/contract radially, a quick calculation of the hoop stress in the pipeline shows that the deformation of the pipe wall is negligible. The hoop stress, $\sigma_\theta$, for a thin-walled cylinder is given by the equation in [61],

$$\sigma_\theta = \frac{Pr}{e}$$  \hspace{1cm} (5.1)

where $P$ is the internal pressure, $r$ is the pipe radius, and $e$ is the pipe wall thickness. Using Hook’s Law, the hoop strain, $\varepsilon_\theta$, for a pipe (open-ended cylinder) is,

$$\varepsilon_\theta = \frac{Pr}{eE}$$  \hspace{1cm} (5.2)

where $E$ is the Young’s Modulus of the pipe. Given that $E = 29,002,666$ psi [62], the internal radius is 1.534 inches (ANSI Sch. 40), the pipe wall thickness is .216 inches (ANSI Sch. 40), and the maximum allowed internal pressure is 300psi, the maximum hoop strain is $7.35 \times 10^{-5}$. The maximum change in internal diameter is $7.17 \times 10^{-5}$ inches. Therefore, the pipe cross-sectional area is assumed constant throughout the simulation.

Pipeline elasticity and the manner in which the pipeline is supported have a significant impact on the wave speed of a pressure transient. This effect is captured in the
modified wave speed equation [1],

\[
a = \sqrt{\frac{K}{\rho}} \frac{1}{\sqrt{1 + c_1(K/E)(D/e)}}
\]  

(5.3)

where \( a \) is the wave speed, \( \rho \) is the fluid density, \( K \) is the bulk modulus of the fluid, \( D \) is the pipeline inner diameter, and \( c_1 \) is a scalar that accounts for the effect of the pipeline’s anchoring. For a pipeline anchored throughout against longitudinal movement [1],

\[
c_1 = \frac{2ez}{D(1 + \mu)} + \frac{D(1 - \mu^2)}{D + e}
\]  

(5.4)

where \( \mu \) is the Poisson’s Ratio for the black-iron pipe (here assumed to be 0.31).

5.1.2.2. Straight Pipe

Among the equations in [1] that are different than those in [60] are the equations for a straight section of pipe. The derivation below fills in steps left out from [1]. The pipeline characteristic impedance, \( B \), is given as,

\[
B = \frac{a}{gA}
\]  

(5.5)

where \( A \) is the pipe’s cross-sectional area, and \( g \) is the acceleration due to gravity (here, 32.2 ft/s\(^2\)). The pipeline characteristic resistance, \( R \), is given as,

\[
R = \frac{\int dx}{2gDA}
\]  

(5.6)

The \( C^+ \) and \( C^- \) characteristic equations are,

\[
H_i = c_{r} - B_{r}Q_i
\]  

(5.7a)

\[
H_i = c_{m} + B_{m}Q_i
\]  

(5.7b)

where,
\[ B_p = B + R|Q_{i-1}| \quad (5.8a) \]
\[ C_p = H_{i-1} + BQ_{i-1} \quad (5.8b) \]
\[ B_m = B + R|Q_{i+1}| \quad (5.8c) \]
\[ C_m = H_{i+1} - BQ_{i+1} \quad (5.8d) \]

In the above equations, \(H\) represents the pressure head, \(Q\) represents the volumetric flow rate, and the subscripts \(i\), \(i-1\), and \(i+1\) refer to the given pipe segment, the upstream pipe segment, and the downstream pipe segment respectively.

Subtracting equation 5.7a from 5.7b yields,
\[
\left( I - P + Q + B \right) = 0 \quad (5.9a)
\]
Thus the flow rate can be solved for, resulting in,
\[
Q_i = \frac{C_p - C_m}{B_p + B_m} \quad (5.9b)
\]

Adding equation 5.7a and 5.7b yields,
\[
2H_i = C_p + C_m + (B_m - B_p)Q_i \quad (5.10a)
\]
Substituting in equation 5.9b yields,
\[
2H_i = C_p + C_m + \left( B_m - B_p \right) \frac{C_p - C_m}{B_p + B_m} \quad (5.10b)
\]
which simplifies to,
\[
H_i = \frac{C_p B_m + C_m B_p}{B_p + B_m} \quad (5.10c)
\]

It is clear from the above equations that flows and pressures in straight pipe sections depend only on the flow rates of adjacent sections from the previous time step.
5.1.2.3. Flooded Tee

The derivation of the equations for a flooded tee, based on the governing equations in [1] is now presented. For the sake of simplicity, flooded tees and tees with air bubbles were assumed to be located at the midpoint of the pipe element whose length they fell within. Therefore, the pipeline characteristic resistance was adjusted to,

\[ R_{half} = \frac{1}{2} R \]  \hspace{1cm} (5.11)

The \( C^+ \) and \( C^- \) equations remain the same. Here, the subscripts are changed to up (upstream) and down (downstream) for simplicity.

\[ C_p = H_{o,up} + BQ_{o,up} \]  \hspace{1cm} (5.12a)

\[ C_m = H_{o,down} - BQ_{o,down} \]  \hspace{1cm} (5.12b)

\[ B_p = B + R_{half} |Q_{o,up}| \]  \hspace{1cm} (5.12c)

\[ B_m = B + R_{half} |Q_{o,down}| \]  \hspace{1cm} (5.12d)

When under \( H \) or \( Q \), subscript \( o \) indicates the value at the beginning of the time step, while \( p \) indicates the value at the end of the time step. The upstream flow is split between the tee and the downstream section of pipe. This separation is captured in the following junction equation,

\[ Q_{p,up} = Q_{p,T} + Q_{p,down} \]  \hspace{1cm} (5.13)

where the subscript \( T \) refers to the tee. The tee volume expands as given by,
\[ \Delta V = (Q_{o,t} + Q_{p,t})dt \]  

(5.14)

where \( \Delta V \) is the change in volume, and \( dt \) is the change in time (size of time step).

Per [1] the tee is modeled as a lumped capacitor, meaning that friction and inertia effects can be neglected, and elastic effects dominate. Therefore, the liquid and tee wall elasticity are combined, and the change in pressure within the tee is given by,

\[ H_{p,t} = H_{o,t} + \frac{K_T dt}{\rho g V_T} (Q_{o,t} + Q_{p,t}) \]  

(5.15)

where \( V_T \) is the volume of the tee. The bulk modulus of the tee, \( K_T \), is given by,

\[ K_T = \frac{K_w}{1 + c_1(K_w/E)(D/e)} \]  

(5.16)

where \( K_w \) is the bulk modulus of the water in the tee. The pressure in the tee is assumed equal to that pipe section it is connected to, therefore,

\[ H_{p,up} = H_{p,t} = H_{p,down} = H_p \]  

(5.17)

This results in a system of four equations and four unknowns. The pressure head and flows at the end of the time step are given by,

\[ H_p = \left( \frac{C_p}{B_p} + \frac{C_m}{B_m} + \frac{\rho g V_T}{K_T dt} H_o + Q_{o,t} \right) \left( \frac{1}{B_p} + \frac{1}{B_m} + \frac{\rho g V_T}{K_T dt} \right) \]  

(5.18)

\[ Q_{p,up} = \frac{C_p - H_{p,up}}{B_p} \]  

(5.19)

\[ Q_{p,down} = \frac{-C_m + H_{p,down}}{B_m} \]  

(5.20)

\[ Q_{p,t} = \frac{\rho g V_T}{K_T dt} (H_{p,t} - H_{o,t}) - Q_{o,t} \]  

(5.21)
The volume and tee compressibility are updated each iteration. The compressibility of the water in the tee is varied as a function of pressure (temperature is assumed constant) based on the empirical relation given in [63],

\[
K_w = \frac{V^0(B - A_1P^2)}{V^p(B + A_1P + A_2P^2)^2} \tag{5.22}
\]

where \( B = 21504.59, A_1 = 3.314, A_2 = -7.388 \times 10^{-6} \), \( P \) is gage pressure in bars, \( V^0 \) is the specific volume of water at atmospheric pressure, and \( V^p \) is the specific volume of the pressurized fluid. For ease of calculation, \( V^p \) is assumed equal to \( V^0 \), which introduces a small error.

### 5.1.2.4. Tee With Air

A tee with an air bubble (modeled as an air chamber in [1]) utilizes the same junction and characteristic equations. However, the compressibility of the water in the tee is assumed negligible compared to that of the air bubble. Thus, the change of pressure within the tee is modeled using the reversible polytropic relation,

\[
H_{p,t}V^n_p = H_{atm}V^n_{atm} = C_A \tag{5.23}
\]

where the subscript \( atm \) refers to ambient conditions (here, sea level at standard temperature). Per [1], the change in volume is now negative, rather than positive, to reflect that the air is compressed when water enters the tee.

\[
\Delta V = -(Q_{v,t} + Q_{p,t})dt \tag{5.24}
\]

Substituting the above equation into the polytropic relation yields,
\[
(H_{p,T} + H_{am} - z)[V - dt(Q_{o,T} + Q_{o,T})]^n = C_A 
\]
(5.25)

where \(z\) represents the head of the air bubble above or below the ambient. From the characteristic equations, the flows in the adjoining pipe sections are given by Eq. 5.19 and 5.20, as before. Using the junction equation (5.13), the flow in the tee is,

\[
Q_{p,T} = - \left( \frac{1}{B_p} + \frac{1}{B_m} \right) H_{p,T} + \frac{C_p}{B_p} + \frac{C_m}{B_m} 
\]
(5.26)

Thus the polytropic relation (5.23) can be expressed as,

\[
(H_{p,T} + H_{am} - z)[V - dt\left(\frac{C_p}{B_p} + \frac{C_m}{B_m} + Q_{o,T}\right) + dt\left(\frac{1}{B_p} + \frac{1}{B_m}\right)H_{p,T}]^n - C_A = 0 
\]
(5.27)

This equation must be solved iteratively for \(H_{p,T}\). The method of choice in the textbook is Newton’s Method. Therefore, the equation is recast as,

\[
F = (H_{p,T} + H_{am} - z)[V - dt\left(\frac{C_p}{B_p} + \frac{C_m}{B_m} + Q_{o,T}\right) + dt\left(\frac{1}{B_p} + \frac{1}{B_m}\right)H_{p,T}]^n - C_A 
\]
(5.28)

Collecting terms,

\[
V_p^n = [V - dt\left(\frac{C_p}{B_p} + \frac{C_m}{B_m} + Q_{o,T}\right) + dt\left(\frac{1}{B_p} + \frac{1}{B_m}\right)H_{p,T}]^n 
\]
(5.29)

Thus, the derivative can be written as,

\[
\frac{dF}{dH} = V_p^n + ndtC_A \left( \frac{1}{B_p} + \frac{1}{B_m} \right) 
\]
(5.30)

Therefore, the equation,

\[
F + \frac{dF}{dH} \Delta H = 0 
\]
(5.31)

can now be solved iteratively for the pressure head of the tee. As before, the pressure
head is assumed the same in the tee and the pipe section it is connected to. The air
volume and $z$ are updated each iteration.

5.1.2.5. Boundary Condition Equations

For the experiments discussed below, there is only one applicable downstream
boundary condition, while there are two applicable upstream conditions. The downstream
condition is a wall, which mimics the behavior of a rigid plug. In this case, the flow rate
at the wall is zero and the $C^+$ characteristic equation, Eq. 5.7a, reduces to $H_{p,wall} = C_p$. If
there is an air pocket at the wall, the reduced $C^+$ characteristic equation is combined with
Eq. 5.23 as shown in [53] and [1].

The first applicable inlet condition is a pressure variation at the inlet, also known
as a “reservoir at upstream end” [1]. Here, the inlet pressure is specified as a function of
time, and the pressure head is solved for using the $C^-$ characteristic Eq. 5.7b. Like many
other standard approaches, this formulation is valid for an idealized case. It assumes the
flow rate at the inlet can take on any value resulting from Eq. 5.7b. It will be shown in
Section 5.3, however, this is not the case for the experimental set up. Therefore, the
following pressure inlet boundary condition is proposed for this thesis,

$$H_0 = H(t)$$

$$Q_0 = 0$$

where the $0$ subscript represents the inlet. It is shown in Section 5.3 that this boundary
condition provides a good approximation of the flow regime in the experimental pipeline.

The second applicable inlet condition is a volume flow inlet, also known as a
velocity inlet, or “discharge at the upstream end” [1]. In this case, the flow rate is
specified as a function of time, and the pressure head is solved for using the $C^-$
characteristic equation 5.7b. Since Eq. 5.7b only uses information at the inlet as well as downstream information, it does not take the pressure of the incoming fluid into account. The development of an improved volume-flow inlet condition is a topic of future research. Nevertheless, a fair approximation of the experimental boundary condition is,

\[ H_0 = H_1 \]  \hspace{1cm} (5.32c)
\[ Q_{inlet} = Q(t) \]  \hspace{1cm} (5.32d)

where the subscript \( i \) denotes the next element downstream of the inlet.

5.2. Verification

Since the MOC code is more versatile than Wood’s code and the WHS code, the performance of each code will first be established by comparing them to steady-state cases whose results are easy to compute. Afterwards, the MOC code will be compared to Wood’s code for a transient case, and as well as cases for which Wood’s code had to be modified. In each case presented in this section, the geometric friction factor in Wood’s code was set to zero.

5.2.1. Steady State Results

The first two cases are fully flooded pipelines. If the bulk modulus of the fluid in the pipeline is assumed constant, which is valid for small changes, the change in pressure due to the injection of a certain volume of fluid is given by the following equation [1],

\[ \Delta p = \frac{\Delta V}{V} K \]  \hspace{1cm} (5.33)

Here the change in volume is positive because an inflow (positive flow) of fluid is equal to a decrease in the total volume of the fluid.
It is a little more difficult to relate pressure changes to inflows for multiphase flows. However, the solution is trivial for steady pressure boundary conditions,

$$\lim_{t \to \infty} \Delta p = \Delta p_{inlet}$$  \hspace{1cm} (5.34)$$

This relation can be used for pipelines with air, as well as fully flooded pipelines.

5.2.1.1 Fully Flooded Straight Pipe

Assume a fully flooded pipeline that is 278ft long, is 3 inches in diameter. Assume the fluid has a wave speed of 4,512.17ft/s, and the pipeline has a Darcy friction factor of 0.021. Now apply a constant pressure head of 164.04ft of water. The response of the pipeline according to each code is shown in Figure 44 below.
Figure 44: Comparison of Steady-State Simulations with Pressure BC – Flooded Pipeline
(a) results from the MOC code with BC computed from Eq. 5.7b, (b) results from the MMOC code with BC computed from Eq. 5.7b, (c) the set up menu for the WHS code, (d) results from the WHS code

In this figure and those that follow, the acronym BC will be used for “boundary conditions.” Each code used a mesh size of eight elements. The MOC and Wood’s code clearly demonstrate a large unsteady response, centered about and converging to the analytical solution. The WHS code, however, shows a perfectly steady response. This is because the WHS code was set up so that a negligible amount of water was allowed to escape through the inlet (it was impossible to set the steady flow discharge to zero).
Using the proposed boundary conditions in Eq. 5.32, the MOC code and Wood’s code produced results nearly identical to those of the WHS code, as shown in Figure 45.

**Table 13: Errors for Steady State Pressure BC**

<table>
<thead>
<tr>
<th></th>
<th>Time-Averaged Pressure (psi)</th>
<th>Relative Percent Error</th>
</tr>
</thead>
<tbody>
<tr>
<td>Analytical Result</td>
<td>71.173</td>
<td></td>
</tr>
<tr>
<td>MOC Code (Eq. 5.7b)</td>
<td>71.173</td>
<td>0%</td>
</tr>
<tr>
<td>MOC Code (Eq. 5.32)</td>
<td>71.173</td>
<td>0%</td>
</tr>
<tr>
<td>Wood’s Code (Eq. 5.7b)</td>
<td>71.714</td>
<td>0.76%</td>
</tr>
<tr>
<td>Wood’s Code (Eq. 5.32)</td>
<td>71.173</td>
<td>0%</td>
</tr>
<tr>
<td>WHS Code</td>
<td>71.173</td>
<td>0%</td>
</tr>
</tbody>
</table>

The one difference between results was that both codes show the pressure rising within the first few time steps to the final solution, whereas the pressure rise in the WHS code is instantaneous. An instantaneous rise is only valid for incompressible flow; therefore, the MOC code and Wood’s code are consistent with the behavior of compressible flows. The error in of the average pressure given by each code is shown in Table 13. It should be noted that the time averaging excludes the initial transient portion of the simulation. The apparent accuracy of Wood’s code (using Eq. 5.7b) changes depending on which time
step is used to begin the time averaging. As the initial time step is moved closer to maximum time, error decreases monotonically, becoming more and more negative.

![Pressure vs. Time](image1.png)

![Pressure vs. Time](image2.png)

![Pressure vs. Time (Wood)](image3.png)

Figure 46: Comparison of Steady-State Simulations with Volume Flow BC – Flooded Pipeline
(a) MOC code with BC computed from Eq. 5.7b using 8 mesh elements, (b) MOC code with BC computed from Eq. 5.7b using 50 mesh elements, (c) Wood’s code with BC computed from Eq. 5.7b using 8 mesh elements

This suggests that the upper peaks are being decreasing faster than the lower limits are increasing. In other words, if left to run longer, Wood’s code would ultimately underestimate the steady-state solution.

For the same pipeline, assume the inlet boundary condition is now a volumetric flow condition. A volume 6.335in³ of water is added to the pipeline (total volume of 23,580.79in³) originally at atmospheric pressure, over the course of 0.63 seconds. As
shown in Figure 46 (a) and (b), the range of pressures calculated by the MOC code using the textbook boundary condition formula changes very little with increased mesh size. The average pressure, however, changes substantially (see Table 14). In addition, it shows very little dampening of the pressure transient. Wood’s code, on the other hand, shows relatively significant dampening, and a much higher initial peak pressure.

No results are reported for the WHS code because it does not include a means of specifying a volume flow inlet boundary condition. Note also that Wood’s code was modified to use the volume flow inlet boundary condition. Figure 47 shows the results of the MOC code with the proposed boundary condition (Eq. 5.32). Although the MOC code consistently underestimates the steady state solution, it converges to it monotonically with increasing mesh size. The results are very poor when the number of mesh elements is small because the assumption given in Eq. 5.32a is inaccurate for large sections of pipe (i.e. large mesh elements).
Figure 47: Comparison of Steady-State Simulations with Volume Flow BC, MOC – Flooded Pipeline

(a) BC computed from Eq. 5.32 using 8 mesh elements, (b) BC computed from Eq. 5.32 using 50 mesh elements, (c) BC computed from Eq. 5.32 using 500 mesh elements

Assuming an equivalent bulk modulus (including the effect of the pipe walls) equal to 271,331.3psi, the errors of the simulations in Figure 46 and Figure 47 are,
Therefore, the MOC has been verified for a straight pipeline.

5.2.1.2. Fully Flooded Pipeline with Tees

Flooded tees behave in a manner analogous to capacitors in a circuit. As the pressure around it increases, the pressure in the tee rises effectively storing the excess energy. Once the pressure in the connecting pipeline decreases, the tee releases energy into the adjacent fluid causing its pressure to rise again. With regard to steady-state solutions, however, the only difference between a straight pipe and a pipe with tees is the time required for the transient to vanish. Since there is no way to calculate this using Wood’s code or the WHS code, the results from the MOC code will be compared against the results from Eq. 5.33 and 5.34.

![Figure 48: MOC code with BC computed from Eq. 5.32 – Flooded Pipeline w/ Tees](a) Pressure BC, (b) Volume Flow BC, mesh of 50 elements

<table>
<thead>
<tr>
<th></th>
<th>Time-Averaged Pressure (psi)</th>
<th>Relative Percent Error</th>
</tr>
</thead>
<tbody>
<tr>
<td>Analytical Result</td>
<td>73.681</td>
<td></td>
</tr>
<tr>
<td>MOC Code Fig. Figure 46 (a)</td>
<td>59.071</td>
<td>-19.83%</td>
</tr>
<tr>
<td>MOC Code Fig. Figure 46 (b)</td>
<td>70.698</td>
<td>-4.05%</td>
</tr>
<tr>
<td>Wood’s Code Fig. Figure 46 (c)</td>
<td>79.808</td>
<td>8.32%</td>
</tr>
<tr>
<td>MOC Code Fig. Figure 47 (a)</td>
<td>49.242</td>
<td>-33.17%</td>
</tr>
<tr>
<td>MOC Code Fig. Figure 47 (b)</td>
<td>67.950</td>
<td>-7.78%</td>
</tr>
<tr>
<td>MOC Code Fig. Figure 47 (c)</td>
<td>72.929</td>
<td>-1.02%</td>
</tr>
</tbody>
</table>
Assume a pipeline with a total volume of 7952.156in$^3$ (including six tees with a total volume of 277.8221in$^3$). A constant inlet pressure head of 50ft of water results in a theoretical final pressure of 21.694psi. The MOC code yields a final pressure of 21.694psi (see Figure 48). The addition of 2in$^3$ of water to the pipeline results in a theoretical final pressure of 54.189psi. The MOC code yields a final pressure of 45.818psi (see Figure 48). Although the relative error (-15.45%) is higher than its straight-pipe counterpart, the general behavior is correct and the under-predicted result is consistent with the behavior observed for straight pipelines. Thus, the code is considered verified for a fully flooded pipeline with tee connections.

5.2.1.3. Pipeline with Air

Assume the same pipeline from Section 5.2.1.1, with an air bubble of volume 1,159.45in$^3$ located at the downstream end of the pipeline. As before, apply a constant pressure head of 164.04ft of water and assume isentropic compression.

![Figure 49: Comparison of Steady-State Simulations with Pressure BC – Pipeline with Air](image)

(a) MOC code with BC computed from Eq. 5.7b using 8 mesh elements, (b) Wood’s code with BC computed from Eq. 5.7b using 8 mesh elements

No results are reported for the WHS code because it does not provide for a downstream
air chamber condition. Wood’s code predicts a much higher pressure rise than the MOC code. For isothermal compression, Wood’s code predicts a peak pressure over 600psi while the MOC code predicts a rise just over 300psi. Nevertheless, both codes clearly converge to the inlet pressure as expected. The oscillations are due to the variation of the inlet flow rate.

By applying the proposed pressure boundary conditions, however, the MOC code demonstrates the nonlinear pressure rise expected of steady compression, and converges to the analytical result (Figure 50). Wood’s code does not show the same trend, possibly due to the fact that it does not update the air bubble head, z, each iteration (see Eq. 5.25).
Figure 51: Comparison of Steady-State Simulations with Volume Flow BC – Pipeline with Air
(a) MOC code with BC computed from Eq. 5.7b using 8 mesh elements, (b) Wood’s code with BC computed from Eq. 5.7b using 8 mesh elements

When the same case is executed with the volume flow boundary condition from Eq. 5.7b, both the MOC and Wood’s code produce very noisy responses of roughly the same magnitude.
Wood’s code yields a pressure at the plug that oscillates about 0.108psi, while the MOC code yields a fairly steady plug pressure of 0.403psi. After applying the proposed boundary condition (Eq. 5.32) to the MOC code the plug pressure holds steady at 0.370psi as shown in Figure 52. Furthermore, the pressure oscillations near the inlet dampen much faster. Given the general behavior of the MOC code, there is ample evidence to conclude the MOC code has been verified for single-phase flows and flows with air pockets.

5.2.2. Transient State Results

Although no verified transient solution has been provided for the MOC code, or Wood’s code, the two codes will be compared using the 285ft case presented Section 5.5.1 of [53].

5.2.2.1. Pipeline with Air

The MOC code and Wood’s code were run using the pressure boundary conditions and initial conditions given in Wood’s code. Note that this includes using a polytropic compression exponent of $n = 1.2$, as suggested in [60]. The differences between the two codes are:

- The mesh elements in the MOC code consists of eight equidistant pipe segments, whereas the mesh elements in Wood’s code have lengths matching the experimental set up described in Section 5.5.1 of [53].
• The time step size in the MOC code is 0.00123s. The time step size in Wood’s code is 0.1s.

As before, Wood’s code predicts a pressure rise significantly higher than the MOC code. Although the pipeline in the MOC code appears to be much more responsive (contains many more pressure waves) than Wood’s code, it also contains the main features predicted by Wood’s code.

5.3. Validation

Before proceeding to show the simulation results, it is important to review the specifics of the experimental set up in order to justify the use of the new boundary conditions given in Section 5.1.2.5.

It is impossible to replicate the pipeline layout of a DOE site because the exact layout of pipelines and the chemical composition of plugs is not known [64]. Therefore, researchers at ARC have constructed several pipelines of varying lengths in order to study the performance of pipeline unplugging technologies under different conditions. One such pipeline was a black iron pipeline 72ft in length (Figure 54, below).
The pipeline consists of four, 3in diameter black iron pipes and one, 3in diameter black iron bend connected by six tee fittings. All connections were threaded. These tee fittings are each equipped with a pressure transducer and a pressure release valve (Figure 55).

The tee fitting at the inlet is connected to a piston pump through a 1.25” pipe via two reducers, as shown in Figure 56 below.
Although tests in 2012 incorporated the use of a solid aluminum plug and a potassium-magnesium-sulfate (K-mag) plug [65], the experimental results will focus on the behavior of the fluid in the pipeline and disregard any solid-fluid interaction. Therefore, the “plug” at the end of the pipeline is small extension terminated with a pressure release valve as shown in Figure 57 below.

Thus, it is clear that the physical control volume (the geometric shape of the fluid volume) begins at the face of the piston cylinder and terminates at the closed pressure release valve. More importantly, the water in the physical experiment is not free to flow.
Rather, it is merely compressed and expanded by the piston. Therefore, the physics of this experimental set up resemble the dynamics of a pressure vessel with a moving boundary much more than that of a large network of pipes with freely flowing fluid.

The computational domain excludes the volume of fluid just before the “inlet tee” and just after the last tee before the plug. That is, the computational domain (in the case of this pipeline) includes roughly 99.754% of the volume of the physical domain. Typical boundary conditions for a pipeline assume that flows entering the pipeline are fully developed. Even if the flows are not fully developed, the boundary conditions in [60] and [1] clearly assume that if one value (inlet pressure or velocity) is specified, the other is free to take on any value that results from the computation (e.g. the upstream reservoir can absorb any flow leaving the pipeline). Given the proximity of the computational inlet to the piston head, it is impossible for the flow rate to assume an arbitrary value when pressure is specified. When air is present and the wave speed is reduced, this proximity also means that the unsteady changes in density and pressure occurring at the piston face cannot be neglected. Furthermore, when the speed of sound is high and the pipeline is short (as is the case with the fully flooded experimental pipeline) the pressure and density of the fluid upstream of the inlet cannot be assumed independent of the downstream flow field. Therefore, typical boundary conditions such as those in [1] are inappropriate for this experimental set up. Instead, new boundary conditions that mimic the physics of a pressure vessel must be developed. Although that research is outside the scope of this thesis, it will be shown in Section 5.3.2 that under the right conditions, the simplified boundary conditions proposed in Section 5.1.2.5 approximate the behavior of the physical pipeline with sufficient accuracy.
5.3.1. Experimental Set Up and Boundary Conditions

The data used in the following section for validation is based on a pipeline similar to that of the previous section. In this case, the pipeline is 94ft long. The specified volume flow, $Q(t)$, will be the time derivative of two third-order polynomials fitted through the measured piston displacement. The pressure head, $H(t)$, will be created using a higher-order polynomial fitted through pressure measurements taken by the “inlet” pressure transducer. The pressure transducers were numbered based on their order downstream of the piston, therefore, the inlet sensor is “P1.” In these experiments, air was only added to the fourth tee, therefore, the pressure of the air was measured by sensor “P4.”

The system described above contains several features that are not modeled by the MOC code:

- Air/water leaks: These can be safely neglected.
- Errors in the measurement of the air volume in the pipeline: They are assumed below the tolerance specified by ARC.
- Piston/Pipeline Vibrations
- Unsmooth piston movement: While establishing an experimental control case, it was observed that the piston velocity was not smooth due to an interaction between the control unit’s signal and the piston. This is assumed negligible.
- Heat Transfer: In previous experiments, the temperature variation measured in the pipeline was less than 1°C and located close to the piston, therefore, the pipeline can be assumed isothermal.
Pipeline Assembly: The pipeline has many small components that connect pipes, bends, and sensors together. Each component contains a contact surface that, if not installed exactly, can create a small gap in the pipeline. This gap will be filled with water or air when the pipeline is flooded and create reflection waves when the water is pressurized. Since the pipeline is assembled by hand, and the components themselves have imperfections, this cannot be avoided.

Behavior of Threaded Joints: It has been observed that threaded joints do not behave like rigid connections [62]. If the pipeline deflection is large enough, this behavior can have a significant impact on the accuracy of the model. Nevertheless, this behavior is neglected in this thesis.

5.3.2. Simulation Results

In the case of a fully flooded pipeline the proposed volume boundary condition predicts a peak at the correct time, but underestimates the peak pressure (Figure 58).

The underestimation of the pipeline transient (even with a larger mesh) is consistent with the behavior observed during verification.
The proposed pressure boundary condition is much more accurate than the volume flow boundary condition (Figure 59).

For a pipeline with 0.327 in$^3$ of air, a piston “half-stroke,” the volume flow boundary condition overestimates the peak pressure (Figure 60). Despite this, the pressure wave noticeably smoothes out as it travels along the pipeline and forms a clear peak which is consistent with the experimental profile.

The pressure boundary condition performs roughly the same in the presence of small amounts of air. It should be noted that two 6$^\text{th}$ order polynomials were used to fit the P1
profile (polynomials up to $25^{th}$ order could not accurately fit the data).

As shown in Figure 61, the pressure peak narrows and rises in a manner consistent with the empirical data, however, does not coincide perfectly with the timing of the pulse. This is partly due to the inaccuracy of the polynomial used to fit the boundary condition.

For a pipeline with 0.654in$^3$ of air (a piston “full-stroke”), the volume flow boundary condition is very accurate at P4 (Figure 62). The average pressure also follows the experimental measurements closely at P1.

The pressure boundary condition, here poorly fit due to the large number of fluctuations in the P1 data, shows a dampening, rather than increase, of the pressure peak at P4.
Despite the inaccuracies presented above, the pressure boundary condition is clearly a good approximation of the experimental data when the pipeline contains little or no air. This is probably because, at high wave speeds, a relatively small addition of water (i.e. small volume flow rate) produces large pressure fluctuations. Since the experimental flow rates are on the order of $10^{-3}$ ft³/s, the assumption of a zero flow rate at the inlet is valid.

Furthermore, the flow rate boundary condition exhibits extraordinary success at modeling the pressure of the tee containing the air pocket. This suggests that the proposed volume flow boundary condition is most appropriate for large volumes of air. Furthermore, it suggests that the accompanying pressure boundary condition (Eq. 5.32c) becomes more accurate as the quantity of air increases. This is because the air dampens out the pressure waves in the system reducing the difference in pressure between pipe segments.

With these observations, it is possible to conclude that the MOC code developed for this thesis, combined with the proposed boundary conditions, is valid for this experimental set up.
5.4. Evaluating and Optimizing the APU

In light of the coupling between the piston pump’s movement, and the behavior of
the pipeline, it is clear that the evaluation of a pump’s performance is specific to the
pump and the pipeline. In addition to this, a plugged pipeline has the added complexity of
the solid-fluid interaction between the plug, the pipe walls, and the surrounding fluid. For
short pipelines with large wave speeds in which the flow cannot be reversed (i.e.
pipelines that resemble pressure vessels), the specification of an arbitrary pressure profile
at the inlet is not enough to properly evaluate the pipeline’s performance. In this case,
there is no guarantee than a given piston pump will have the ability to reproduce a given
pressure profile. Therefore, the evaluation of a pipeline will require some
experimentation to determine the maximum and minimum pressures achievable with a
given piston pump, and the frequencies at which those pressures can be generated. Given
the bounds of a pump’s performance, however, an optimization algorithm can be used to
determine the optimal performance of the pump.

During earlier experiments, it was determined that if the pipeline conditions are
right and the timing of the pulses is right, pressure waves reaching the blockage can cause
a buildup of pressure at the surface of the blockage that will ultimately dislodge the plug
[57]. To avoid damaging the pipeline, the DOE designated safety limit for pipeline
pressure is 300 psi [57]. Furthermore, to avoid cavitation the pressure in the pipeline must
remain above -14.447 psi [66]. With this information, it is now possible to formulate the
APU operation process as an optimization process.

5.4.1. Optimization Problem Set Up

The goal of the APU unplugging process is to maximize the difference in
pressures at each face of the plug over the period \((t_{\text{max}})\) of the operation of the APU. Therefore, the objective function can be written as:

\[
U = - \frac{\sum_{i=0}^{t_{\text{max}}} |\Delta P(t)|}{t_{\text{max}}}
\]  

(5.35)

where \(\Delta P\) represents the difference in pressure across the plug. The objective function is negative because the hybrid optimization algorithm is designed to locate the global minimum.

The pressure at one surface of the plug is a function of the frequency and peak pressures produced by the piston. Assuming a sequence of 5 consecutive half-pulses at the same frequency, the design vector representing the piston’s pulse schedule can be written as:

\[
\tilde{x} = (P_1, P_2, P_3, P_4, P_5, P_6, P_7, P_8, P_9, P_{10}, f_1, f_2)
\]  

(5.36)

Where \(P_i\) is the amplitude of the pressure of the \(i^{th}\) pulse, \(f_1\) and \(f_2\) are the frequency of the first and second five pulses, respectively. The first five amplitudes correspond to the piston acting upstream of the plug; while the second set of five correspond to a second piston acting downstream of the plug. Here, a “half-pulse” means the pressure generated during the first half of the period of a sinusoidal pulse.

Given the limitations on the pressure in the pipeline, this optimization problem contains two constraints:

\[
g - 300 < 0
\]

\[
-14.448 - g < 0
\]  

(5.37)

For each constraint violation at any point during the simulation, anywhere in the pipeline, a penalty of 300psi is added to the objective function value.
In order to better represent the physical limitations of the experimental piston pump, the frequencies will be between .25 Hz and 2 Hz, and the peak pressures will be between -13.5psi and 257psi. Note that the pressure peaks correspond to a reverse stroke of 0.5in and a forward stroke of 9.5in, respectively, for a fully flooded pipeline.

Each simulation was executed for 20 seconds of physical time, and the pipeline was assumed symmetrical. The mesh for each side of the pipeline contained 20 elements to reduce computational cost. The pipeline was assumed straight. The hybrid was run for 50 iterations. Optimization was performed on a fully flooded pipeline as well as a pipeline containing an air bubble on one side of the plug. Each of those two cases was executed for synchronized-piston \((f_1 = f_2)\) and asynchronous-pulsing \((f_1 \neq f_2)\) cases.

### 5.4.2. Optimization Results

The fully flooded, synchronized case resulted in the logical (arguably trivial) result, where the pistons apply positive pressure on one side of the pump, and a negative pressure on the other side of the pump. This result is the natural consequence of the high wave speed present in fully flooded pipelines. More importantly, it serves as a validation of the optimization algorithm. Despite the multi-modal nature of the objective function,
(the trivial result is not unique) the optimization algorithm converged to the logical answer given the constraints applied to the pipeline, and the limitations of the piston pump.

The second case involves a pipeline that is fully flooded on the downstream side of the plug, but, contains an air bubble (“full-stroke” of air) on the upstream side of the plug with synchronized pistons. This profile is very similar to the first case, except that the upstream side of the pipeline displays the slow response time typical of air bubbles. The solution obtained by the optimization algorithm is clearly the reasonable solution: the maximum positive pressure is applied on the flooded side of the plug, whereas the suction is applied on the side with the air. If the positive pressure was applied on the side with air, the bubble would dramatically reduce the peak pressure, and since the minimum pressure has a low magnitude compared to the maximum positive pressure, it would be impossible to maximize the pressure across the plug in this way.
Figure 66, above, shows the results for the fully flooded and multiphase cases in which the pistons are allowed to pulse asynchronously. Unlike the synchronized case, the new piston operation schedule for the fully flooded case favors operating at the upper limits of the frequency range. The downstream (negative pulse) pistons were operated at a lower frequency of 0.2698Hz, than the upstream (positive pulse) pistons, for which the frequency was 1.9209Hz. The asynchronous case with air arrives at nearly the same solution (0.25Hz and 0.2698 Hz), as the synchronized case. Table 15, below, shows the
results and average pressure across the plug for each case.

Table 15: Optimal Piston Schedules

<table>
<thead>
<tr>
<th>Figure</th>
<th>Upstream Piston Schedule</th>
<th>Downstream Piston Schedule</th>
<th>Avg. ΔP</th>
</tr>
</thead>
<tbody>
<tr>
<td></td>
<td>P₁, P₂, P₃, P₄, P₅, f₁</td>
<td>P₆, P₇, P₈, P₉, P₁₀, f₂</td>
<td></td>
</tr>
<tr>
<td>Figure 64</td>
<td>257, -13.5, -13.5, -13.5, -13.5, 0.2698</td>
<td>-13.5, 257, 257, 257, 257, 0.2698</td>
<td>173.69</td>
</tr>
<tr>
<td>Figure 65</td>
<td>257, 257, 257, -13.5, 257, 0.2698</td>
<td>-13.5, -13.5, -13.5, 257, -13.5, 0.2698</td>
<td>172.30</td>
</tr>
<tr>
<td>Figure 66(a)</td>
<td>-13.5, -13.49, -13.5, -13.49, -13.49, 1.9209</td>
<td>257, 257, 257, 257, 257, 0.2698</td>
<td>173.61</td>
</tr>
<tr>
<td>Figure 66(b)</td>
<td>257, 257, 257, 257, 257, 0.2698</td>
<td>-13.5, -13.5, -13.5, -13.5, -13.5, 0.25</td>
<td>173.61</td>
</tr>
</tbody>
</table>

At this stage the reader is cautioned again, that the true behavior of a piston can only be studied within the context of a coupled solver that simultaneously computes the pressure transients as well as the solid-fluid interaction with the piston and any other relevant physical considerations. Nevertheless, the optimization algorithm clearly provides reasonable results and demonstrates that it is possible to get higher average pressures across the plug by operating the pistons asynchronously, rather than synchronously, when air is present in the pipeline.
6. CONCLUSIONS

This thesis has achieved its primary goals. First, a novel, hybrid optimization algorithm switching logic was proposed. After being tested against other algorithms using standard benchmarking techniques, it was found to be superior to many algorithms over a broad set of test cases. Due to the added overhead, however, the BST-SPC algorithm was found to be superior to the hybrid in a variety of test cases as well. Nevertheless, the hybrid is a strong competitor outperforming OPTRAN in the majority of unconstrained test cases, and performing competitively against other hybrid algorithms in other test cases.

Second, a general framework for gaging the performance of hybrid optimization algorithms was proposed. This framework can be applied to any hybrid whose express aim is to outperform its constituent algorithms. Within this context, the hybrid is found to perform “typically” suggesting that its extra computational expense prevents it from maximizing the potential given the constituent algorithms at its disposal. Therefore, in the future it would be useful to incorporate surrogate models, or improve the search vector/constituent selection scheme so that the overhead costs of the hybrid are reduced.

Third, a Method of Characteristics code was developed to model the Asynchronous Pulse Unit. Using an updated version of the text book, and including new boundary conditions proposed for the purpose of this thesis, the code was found to accurately model a variety of experimental flows. With this, it became possible to connect the hybrid optimization algorithm to the MOC code so that the operation of the
piston pumps could be performed. Once done, it was found that higher average pressures across the plug could be achieved by pulsing the pistons asynchronously.

The research performed for this thesis also opened up new topics of study. For example, the question of how to decompose an objective function’s topology into effective topologies remains unsolved. For many problems it is clear that this decomposition is not unique. Therefore, if it is done carelessly, it is possible that an incorrect set of effective topologies will be selected, which will misguide the optimization algorithm. Furthermore, the question of how to best identify the iterative heuristic that solves a problem given this set of effective topologies remains unanswered. It is clear from the results in this thesis that matching the fittest search vector to the constituent algorithm centroid is a useful approach, but it is only part of the solution. Could there be rules, or a probabilistic selection scheme that would improve the hybrid’s robustness? What additional measurements should the selection criteria be based on (e.g. standard deviation of the range)?

What is clear from these questions and the effectiveness of the proposed hybrid, is that this is a fruitful area of research with ample room for continued investigation. Although it may not be possible to create a single algorithm that solves all problems, a well-constructed hybrid algorithm has clear advantages over any other kind of optimization algorithm.
REFERENCES


APPENDIX

A.1. Additional Global Optimization Algorithms

The information below was previously presented in [67] and [68]. The interested reader is referred to those publications for more in-depth analysis.

A.1.1. Quantum-Behaved Particle Swarm (QPS)

The fundamental difference between this algorithm and PSO is simply the update equation. QPS no longer utilizes an inertia term, and Eq. 2.4 is replaced with,

\[
\tilde{X}_{i}^{g+1} = \tilde{p}_{i} \pm \alpha |\tilde{C}^{g} - \tilde{X}_{i}^{g}| \ln \left( \frac{1}{R_{3}} \right)
\]  \hspace{1cm} (A.1)

Where \( \tilde{p}_{i} \) is the “local attractor” defined using an equation reminiscent of Eq. 2.5,

\[
\tilde{p}_{i} = \phi \tilde{X}_{best,i} + (1-\phi) \tilde{X}_{best,G}
\]  \hspace{1cm} (A.2a)

\[
\phi = \frac{\beta R_{3}}{\beta R_{1} + \gamma R_{2}}
\]  \hspace{1cm} (A.2b)

and \( \tilde{C}^{g} \) is the “mean best value” which is simply the arithmetic mean of the individual best vectors for each generation. The scalars \( R_{1}, R_{2} \) and \( R_{3} \) are random numbers \( \epsilon [0,1] \), and \( \alpha \) is a user-defined parameter, set here to decrease linearly during each generation from 1 to 0.5, per the authors’ recommendation in [69]. The scalars \( \beta \) and \( \gamma \) are also user-defined scalars (here \( \beta = \gamma = 2 \)). Note that the \( \pm \) operation in Eq. A.1 is resolved simply by assigning a 50% probability that the term will be either positive or negative.

A.1.2. Modified Quantum-Behaved Particle Swarm (MQP)

The modification to QPS presented here modifies the local attractor randomly according to the scheme recommended in [69]. The rest is identical to QPS. The authors suggest setting \( \alpha \) to decrease linearly each generation from 1 to 0.4 [69].
A.1.3. Firefly Algorithm (FFA)

Under certain circumstances, FFA reduces to a special case of PSO [70]. Unlike any other method presented here, it uses two loops over the population with one loop nested inside the other. The attractiveness equation used in this thesis is a replica of the equation provided in Dr. Yang’s MATLAB code, which is,

$$\beta = (\beta_0 - \beta_{\text{min}}) e^{-\gamma^2} + \beta_{\text{min}}$$  \hspace{1cm} (A.3)

where $\gamma$ is a user-defined parameter (here set to 1). The scalars $\beta_0$ and $\beta_{\text{min}}$ are also user-defined parameters (here set to 1 and 0.2, respectively). Finally, $r$ is the Euclidean distance. The small random perturbation vector is generated in the MATLAB code according to,

$$\bar{u} = \sum_{i=1}^{\text{dim}} (\alpha(R - 0.5)L) \hat{e}_i$$  \hspace{1cm} (A.4)

where $\alpha$ is a user-defined parameter that decreases linearly with the generation number from 0.25 to 0, $R$ is a uniformly distributed random number $\epsilon [0,1]$, and $L$ is the width of the search domain in the $\hat{e}_i$ direction. Finally, the update equation is given as,

$$\bar{X}_{i}^{g+1} = (1 - \beta) \bar{X}_{i}^{g} + \beta \bar{X}_{j}^{g} + \bar{u}$$  \hspace{1cm} (A.5)

The resemblance to the velocity equation of PSO is striking.

A.1.4. Bat-Inspired Algorithm (BAT)

This algorithm resembles DE in that it uses comparisons to update each new generation of candidate solutions. Reminiscent of best/2/bin, it partially incorporates perturbations of the best design vector into its formulation, and like sociability-only-PSO, it searches the function space based on a multiple of the difference between the global
best and a particular design vector. However, each of these steps are performed separately, and the DE-style comparison is used at the end to determine which candidate solutions are retained and which are discarded. Similar to PSO, the algorithm has a velocity equation is given by,

$$V_{i,k}^{g+1} = V_{i,k}^{g} + (X_{i,k}^{g} - X_{bestG,k}^{g}) \times f$$  \hspace{1cm} \text{(A.6)}$$

where a new frequency is generated for each velocity vector component (denoted by k).

As in PSO, The bat update equation is the same as Eq. 2.4.

**A.1.5. Cuckoo Search (CKO)**

Another method reviewed for this thesis is CKO [71]. Like BAT, it also uses DE style comparisons. The main drawback of this algorithm is that it requires two objective function evaluations per generation. The first (Lévy flight) update is executed according to the following equation,

$$X_{i,k}^{g+1} = X_{i,k}^{g} + \left(0.01 \frac{\sigma N_1}{N_2}^{1/\beta} (X_{i,k}^{g} - X_{best G,k}^{g}) \right) \times N_3$$  \hspace{1cm} \text{(A.7)}$$

where N1, N2 and N3 are normally distributed numbers centered at zero, $\beta$ is a user-defined parameter (here, $\beta = 1.5$), k is the dimension number, and $\sigma$ is a constant. The second (empty nest) population update is executed as follows,

$$\text{if } (R_1 > p_a)
\tilde{X}_{i}^{g+1} = \tilde{X}_{i}^{g} + R_2 (\tilde{X}_{j}^{g} - \tilde{X}_{k}^{g})$$  \hspace{1cm} \text{(A.8)}$$

where R1 and R2 are uniformly distributed random numbers $\epsilon [0,1]$, and $p_a$ is a user-defined parameter (here $p_a = 0.25$). The two population members denoted by j and k are randomly selected from the population.

**A.1.6. Biogeography-Based Optimization (BBO)**
BBO is modeled after the migration of species from one habitat to another [3]. Unlike the previous methods, BBO does not use an update equation, but rather, creates new design vectors by creating permutations of existing design vectors based on their fitness. BBO contains three user-defined parameters: $\mu$, $\lambda$ and the mutation probability. Researchers have found that varying the emigration and immigration profiles can improve the algorithm’s performance [72], but that is outside the scope of this thesis. The value of $\mu$ must be nonnegative, or the algorithm may never perform emigration (the update). The update condition is given by the following inequality,

$$\text{if } \sum_{i=1}^{k} \mu_i > R \times \sum_{i=1}^{\text{pop}} \mu_i, \text{then select individual } k \text{ for emigration}$$  \hspace{1cm} (A.9)

where $R$ is a random number between zero and one, and $k$ is the design vector whose $\mu$ value makes the condition statement true.

A.1.7. Particle Swarm With Random Differences (RD)

Inspired by the concept of random walks and the modified local attractor of MQP, this proposed modification adds versatility to PSO without dramatically changing its basic procedure. Traditionally, each design vector in PSO is subtracted from its own individual best vector. This method, proposed in [67], introduces a probability that a design vector will be subtracted from an individual best other than its own based on the value $R_p$, which is the “probability of random differencing,” a user-defined parameter $\epsilon$ [0,1]. The above algorithm is roughly equivalent to rewriting Eq. 2.4 as follows,

$$\tilde{v}_i = \alpha \tilde{v}_{i-1} + \beta R_1 (\tilde{x}_{\text{best},i} - \tilde{x}_i) + \beta R_2 (\tilde{x}_{\text{best,G}} - \tilde{x}_i)$$

where $i, j \in [1, \text{pop}]$ and $i \neq j$  \hspace{1cm} (A.10)
where pop is the number of candidate design vectors in the population. The requirement that \( i \) not equal \( j \) is not strictly enforced when the vectors are shuffled.

**A.1.8. Modifications to DE**

Over the years many modifications to DE have been proposed. Some of these modifications relate to the update equation [4], while others involve merging DE with some other technique [73] [74]. Recently, three types of modifications were proposed in [68] that draw more performance out of DE while minimizing the changes made to the implementation of DE.

1. **Randomly Varying Parameters**

   Randomly varying both \( F \) and \( CR \) dramatically improves the convergence speed and robustness of the DE methods in many test functions.

2. **Special Vectors**

   Since DE involves the weighted average of a collection of vectors, including special vectors in the population (such as the weighted average) after each iteration has been shown to improve performance. An average vector, and a weighted average vector are used for this purpose. This was the precursor to the search vector concept.

3. **Sorted Comparisons**

   This method has been shown to greatly increase convergence speed but reduce robustness, causing the algorithm to converge to local minima more frequently.
A.2. Comparison of Standard Optimization Algorithms

Figure 67, below, compares the relative accuracy and speed of each algorithm. Although PSO obtained the single best answer in 58.4% of the test cases, STD and DN3 outperformed all other methods in speed and accuracy in nearly half of the test cases.

Second to STD and DN3 in both speed and accuracy were PSO and MQP. These four methods combined account for nearly 75% of the best-performing algorithms. Although BST, TDE and QPS have accuracies above 10%, they suffer in the combined speed-accuracy category due to their lack of speed relative to other methods. BBO rarely outperforms any of the other algorithms. This may be due to the fact that BBO relies on two processes in order to produce new designs:

1. Permutations: given an existing population, each new member generated by the “migration” operation is simply a permutation of existing members.

Each of these processes relies entirely on the quality of the RNG as well as shape of the topology. BBO will converge to the global minimum if and only if the population already contains components of that exact vector (neglecting the effects of mutation). Now let the definition of convergence to the global minimum be relaxed to mean convergence to the convex region containing the global minimum. This will enable BBO to converge to the global minimum if the population contains components from the set of vectors that occupy that convex region. If even one component of these vectors is missing from the population, and if this component is never randomly generated during the optimization process, it is impossible for BBO to converge to the global minimum. Therefore, BBO is at a disadvantage when searching a continuous domain because it can only search using permutations.

On the other extreme, there is a subset of the Schittkowski & Hock test cases for which some algorithms are perfectly suited. The test cases are given in Table 16,

**Table 16: Selection of Test Cases – Exceptionally Good Algorithm Performance**

<table>
<thead>
<tr>
<th>TC</th>
<th>Dim</th>
<th>Objective Function</th>
<th>Constraint Equation(s)</th>
</tr>
</thead>
<tbody>
<tr>
<td>4</td>
<td>2</td>
<td>$U(x) = \frac{(x_1 + 1)^3}{3} + x_2$</td>
<td>n/a</td>
</tr>
<tr>
<td>45</td>
<td>5</td>
<td>$U(x) = 2 - \frac{x_1x_2x_3x_4x_5}{120}$</td>
<td>n/a</td>
</tr>
<tr>
<td>220</td>
<td>2</td>
<td>$U(x) = x_1$</td>
<td>$(x_1 - 1)^2 - x_2 = 0$</td>
</tr>
<tr>
<td>234</td>
<td>2</td>
<td>$U(x) = (x_2 - x_1)^3 - (1 - x_1)$</td>
<td>$-x_1^2 - x_2^2 + 1 &lt; 0$</td>
</tr>
<tr>
<td>236</td>
<td>2</td>
<td>$U(x) = \left( b_1 + b_2x_1 + b_3x_1^3 + b_4x_1^4 + b_5x_2 + b_6x_1x_2 + b_7x_1^2 + b_8x_1x_2^2 + b_9x_2^2 + \ldots \right)$</td>
<td>$x_1x_2 - 700 &lt; 0$</td>
</tr>
<tr>
<td></td>
<td></td>
<td>$\ldots + b_{10}x_1^3x_2^3 + b_{11}x_1^4x_2^4 + b_{12}x_1^5x_2^5 + \ldots$</td>
<td>$x_2 - \frac{x_1^2}{125} &lt; 0$</td>
</tr>
<tr>
<td>238</td>
<td>2</td>
<td>$U(x) = \left( b_{20}e^{50-4x_1x_2\ldots} + b_{21}x_1^3x_2^3 + b_{22}x_1x_2^2 + b_{23}x_1^2x_2^2 + \ldots \right)$</td>
<td>$x_1x_2 - 700 &lt; 0$</td>
</tr>
<tr>
<td></td>
<td></td>
<td>$\ldots + b_{24}x_1^5x_2^5 + b_{25}x_1^6x_2^6 + b_{26}x_1^7x_2^7 + \ldots$</td>
<td>$x_2 - \frac{x_1^2}{125} &lt; 0$</td>
</tr>
<tr>
<td></td>
<td></td>
<td>$\ldots + b_{27}x_1^8x_2^8 + b_{28}x_1^9x_2^9 + \ldots$</td>
<td>$(x_2 - 50) - 5(x_1 - 55) &lt; 0$</td>
</tr>
</tbody>
</table>
In many cases (TC 4, 45, 220, 234, 236, 238, 239), the global minimum is located at the boundary of the search domain. This can make minima easier to find when the design domain is enforced by projecting the violating vector component back to the boundary. The global minimum can be found simply by overshooting the domain’s boundaries.
As shown in Figure 68-75, BBO converges very slowly. This is due, in part, to the fact that BBO cannot overshoot a domain boundary (assuming the domain is closed and simply connected). Therefore, it must approach the boundary at whatever speed the population of design vectors allows.

In several cases, such as Figure 69 above, an optimization algorithm (BAT in this case) locates the global minimum within two iterations. It is important to note that the convergence history plots show the error of the lowest population member each iteration, and not the average error of the population.
Cases 236, 238, and 239 share the same objective function but have different constraints as well as different search domain sizes. These differences do not appear to alter the performance of the optimization algorithms.
Figure 71: Convergence Histories, Test Case 234
Figure 72: Convergence Histories, Test Case 236

The wavy surface that appears in TC 236 and 239 is barely visible once the search domain is expanded, as in TC 238. Effectively flattening the topology in this way causes FFA to become trapped in local minima.
Figure 73: Convergence Histories, Test Case 238
FFA also performs poorly on TC 242 and 220. Recall that although TC 220 has a linear objective function, the application of penalties for constraint violations causes the topology to change.
This change in topology is what causes optimization algorithms to apparently become caught in local minima, despite the smooth objective function topology. TC 331 is a remarkable case in which BBO converges quickly to the global minima, relative to its performance in many other test cases.
Figure 76: Convergence Histories, Test Case 331
TC 378 has a higher dimensionality than the previous cases. Here, the so called “curse of dimensionality” is not yet observed, although it will become rapidly apparent in other test cases.

TC 111 is very similar to TC 378, and the optimization algorithm perform similarly well on it.

As one might expect, there is overlap in the performance of the various DE optimization algorithms. In some cases, such as those shown in Table 17, the DE optimization algorithms perform very well, showing only minor differences between algorithms.
Table 17: Selection of Test Cases – Similar, Good Performance: DE Algorithms

<table>
<thead>
<tr>
<th>TC</th>
<th>Dim</th>
<th>Objective Function</th>
<th>Constraint Equation(s)</th>
</tr>
</thead>
<tbody>
<tr>
<td>3</td>
<td>2</td>
<td>$U(\bar{x}) = 10^2 (x_2 - x_1)^2 + x_2$</td>
<td>n/a</td>
</tr>
<tr>
<td>21</td>
<td>2</td>
<td>$U(\bar{x}) = 0.01 x_1^2 + x_2^2 - 100$</td>
<td>$10x_1 - x_2 - 10 &lt; 0$</td>
</tr>
<tr>
<td>204</td>
<td>2</td>
<td>2nd order polynomial</td>
<td>n/a</td>
</tr>
<tr>
<td>245</td>
<td>2</td>
<td>$U(\bar{x}) = \sum_{i=1}^{10} (e^{-0.1x_i} - e^{-0.1i\pi} - x_i e^{-0.1i} - e^{-i})^2$</td>
<td>n/a</td>
</tr>
</tbody>
</table>

Note that the difference between TC 245 and TC 242 is the size of the search domain. In TC 242, the domain is a cube with sides [0, 10]. In [13], the domain for TC 245 is $(-\infty, \infty)$, but since this is impossible to program, the domain was set to [-300, 300]. This was the limit set for most test cases having an infinite domain.

![Figure 78: Convergence Histories - Similar, Good Performance: DE Algorithms](image_url)
In other cases the DE optimization algorithms all perform poorly, such as in the following table,

**Table 18: Selection of Test Cases – Similar, Bad Performance: DE Algorithms**

<table>
<thead>
<tr>
<th>TC</th>
<th>Dim</th>
<th>Objective Function</th>
<th>Constraint Equation(s)</th>
</tr>
</thead>
<tbody>
<tr>
<td>47</td>
<td>5</td>
<td>(U(\bar{x}) = (x_1 - x_2)^2 + (x_2 - x_3)^2 + (x_3 - x_4)^2 \ldots + (x_4 - x_5)^2)</td>
<td>(x_1 + x_2^2 + x_3^2 - 3 = 0)  \(x_2 - x_3^2 + x_4 - 1 = 0)  \(x_4 x_5 - 1 = 0)</td>
</tr>
<tr>
<td>48</td>
<td>5</td>
<td>(U(\bar{x}) = (x_1 - 1)^2 + (x_2 - x_3)^2 + (x_4 - x_5)^2)</td>
<td>(x_1 + x_2 + x_3 + x_5 + x_5 - 5 = 0)  \(x_3 - 2(x_4 + x_5) + 3 = 0)</td>
</tr>
<tr>
<td>293</td>
<td>50</td>
<td>(U(\bar{x}) = \left(\sum_{j=1}^{50} i x_j^2\right)^2)</td>
<td>n/a</td>
</tr>
</tbody>
</table>

The poor performance in TC 293 can be attributed to its high dimensionality. The functions for TC 290, 291, 292, and 293 vary only in their upper limits, which are 2, 10, 30, and 50 respectively.
Therefore, despite the fact that the DE family of optimization algorithms performs well on several low-dimension, even functions, as the number of equality constraints or problem dimensionality increases performance can rapidly decrease.
In some test cases, the donor formulation of the DE optimization algorithms becomes important. A few examples include,

<table>
<thead>
<tr>
<th>TC</th>
<th>Dim</th>
<th>Objective Function</th>
<th>Constraint Equation(s)</th>
</tr>
</thead>
<tbody>
<tr>
<td>79</td>
<td>5</td>
<td>( U(\vec{x}) = (x_1 - 1)^2 + (x_1 - x_2)^2 + (x_2 - x_3)^2 \ldots ) ( + (x_3 - x_4)^4 + (x_4 - x_5)^4 )</td>
<td>( x_1 + x_2^2 + x_3^2 - 2 - 3\sqrt{2} = 0 ) ( x_2 - x_3^2 + x_4 + 2 - 2\sqrt{2} = 0 ) ( x_4x_5 - 2 = 0 )</td>
</tr>
<tr>
<td>241</td>
<td>3</td>
<td>Sixth order polynomial</td>
<td>n/a</td>
</tr>
<tr>
<td>393</td>
<td>48</td>
<td>See [13]</td>
<td>1 Nonlinear Inequality, 2 Linear Equality</td>
</tr>
<tr>
<td>395</td>
<td>50</td>
<td>( U(\vec{x}) = \sum_{i=1}^{50} \left(x_i^2 + x_i^4\right) )</td>
<td>(-1 + \sum_{i=1}^{50} x_i^2 = 0)</td>
</tr>
</tbody>
</table>

Figure 15, below, demonstrates the variability in performance. Not only is the donor formulation for each DE algorithm different, but in the case of BST four vectors are used to perturb the donor rather than two.
Therefore, while they share a common structure, each variation of DE can have markedly different performance on a design problem. No single variation of DE is superior to the others, or representative of the others’ behavior.

The Particle Swarm family of optimization algorithms (PSO, QPS, and MQP) also suffers from the curse of dimensionality, but QPS and MQP appear to perform slightly better on TC 290-293 than the DE algorithms. As shown in Figure 16, QPS and MQP are still able to locate the global minimum within the allotted 200 iterations on the 10-dimensional TC 291. After that, the performance gradually worsens, but is still better than PSO and the DE optimization algorithms.
Similar to the DE optimization algorithms, the PSO family of algorithms sometimes performs the same, and other times it does not (see Figure 82). Each of the algorithms in this group represents a viable alternative to DE, although PSO generally outperforms QPS and MQP.
TC 5, 9, and 110 are given in Table 20, below. From this it is clear that PSO, QPS, and MQP are capable of handling highly nonlinear problems, although problem dimensionality is still an issue.

Table 20: Selection of Test Cases – Varying Performance Among DE OAs

<table>
<thead>
<tr>
<th>TC</th>
<th>Dim</th>
<th>Objective Function</th>
<th>Constraint Equation(s)</th>
</tr>
</thead>
<tbody>
<tr>
<td>5</td>
<td>2</td>
<td>$U(\vec{x}) = \sin(x_1 + x_2) + (x_1 - x_2)^2 - 1.5x_1 + 2.5x_1 + 1$</td>
<td>n/a</td>
</tr>
<tr>
<td>9</td>
<td>2</td>
<td>$U(\vec{x}) = \sin(\pi x_1 / 12)\cos(\pi x_2 / 16)$</td>
<td>$4x_1 - 3x_2 = 0$</td>
</tr>
<tr>
<td>110</td>
<td>10</td>
<td>$U(\vec{x}) = \sum_{i=1}^{10} \left[ (\ln(x_1 - 2))^2 + (\ln(10 - x_1))^2 \right] - \left( \prod_{i=1}^{10} x_i \right)^{0.2}$</td>
<td>n/a</td>
</tr>
</tbody>
</table>
The algorithms by Dr. Yang and Dr. Simon rarely outpace the PSO family of algorithms, although they can match their accuracy. In some cases, however, such as TC 395 below, the FFA greatly outperforms other methods.

![Convergence Histories – Varying Performance of Yang/Simon Algorithms](image)

Figure 83: Convergence Histories – Varying Performance of Yang/Simon Algorithms

The success of algorithms like FFA on high dimensional problems (using the nonlinear population function) shows that the curse of dimensionality is the result of the relationship between an algorithm and the objective function topology, not just the population size. In other words, while conventional wisdom might call for increasing the population size as dimensionality increases, it is also possible to overcome the curse of dimensionality by modifying an algorithm to better suit a given topology, changing from one algorithm to
another, or creating an algorithm capable of adapting on its own. Simply increasing the population size in hopes of obtaining better results is a merely a brute-force approach.

A.3. Comparison of Modified Optimization Algorithms

The modifications to DE and PSO discussed in [67] and [68] greatly enhance each algorithm’s performance. Figure 87, below, compares the relative performance of all modified optimization algorithms, and their standard counterparts. The single greatest improvement in performance can be seen in the difference between BST and BST-SPC, as well as BST and BST-R to a lesser degree. The randomization of parameters dramatically improves the performance of BST because it allows for a wider range of perturbations to the global best vector. It also improves the performance of STD but appears to decrease the performance of DN3 as well as TDE.

![Figure 84: Best Performance of Unmodified vs Modified Optimization Algorithms](image)

This apparent decrease, however, is misleading. In fact, it is more appropriate to say that randomizing the parameters of DN3 as done here shifts the class of functions for which it
is well suited from one type to another. Consider, for example, the 50-dimensional problems, TC 293 and 395, as well as the 100-dimensional problems, TC 299, and 302.

In each of the above cases, the algorithm’s accuracy (and speed) is improved by several orders of magnitude. Therefore, there is a tradeoff between improvement in some problems and worsening in others. It should be noted that modified DN3’s performance also improves in the 100-dimensional TC 305, but by less than one order of magnitude. Therefore, randomization may not lead to performance improvements in all higher dimension problems.

Figure 85: Convergence Histories – Modified DN3
Table 21: Selection of Test Cases – Modifications to DN3

<table>
<thead>
<tr>
<th>TC</th>
<th>Dim</th>
<th>Objective Function</th>
<th>Constraint Equation(s)</th>
</tr>
</thead>
<tbody>
<tr>
<td>299</td>
<td>100</td>
<td>$U(\bar{x}) = 10^{-4} \sum_{i=1}^{99} \left[ 100(x_{i+1} - x_i^2)^2 + (1-x_i)^2 \right]$</td>
<td>n/a</td>
</tr>
<tr>
<td>302</td>
<td>100</td>
<td>$U(\bar{x}) = x_1^2 - 2x_1 + \sum_{i=2}^{100} \left[ 2x_i^2 - 2x_{i+1}x_i \right]$</td>
<td>n/a</td>
</tr>
<tr>
<td>305</td>
<td>100</td>
<td>$U(\bar{x}) = \left( 0.5 \sum_{i=1}^{100} i^2 x_i \right)^4 + \left( 0.5 \sum_{i=1}^{100} i x_i \right)^2 + \sum_{i=1}^{100} x_i^2$</td>
<td>n/a</td>
</tr>
</tbody>
</table>

Similar to DN3, the addition of random differencing (PRD) to PSO tends to improve its performance in higher dimensional problems. Although PRD decreases PSO’s convergence speed (PSO appears to converge rapidly to local minima), it can enable PSO to escape local minima and continue moving toward the global minimum.

Figure 86: Convergence Histories – PSO and PRD

Another notable feature of PRD, is that PRD obtains the single-worst minimum in a
problem far less frequently than PSO, but also obtains the single-best minimum less frequently than PSO. PRD is very sensitive to the selection of $R_p$. While a linearly decreasing $R_p$ value was selected for this thesis, linearly increasing and constant values of $R_p$ have all been tested. Like randomization for DN3, each $R_p$ value shifts the suitability of PRD from one class of problems to another.

![Figure 87: Convergence Histories – Various Improvements Due to Modifications](image)

Some notable convergence history improvements are shown in Figure 87. As indicated with DN3-R and PRD, modifications to these algorithms should be thought of as shifting an algorithm from being well suited for one class of problems to another. Therefore, even though STD and BST become more robust as a result of their improvements, these
improvements still come at the expense of performance in certain test cases that STD and BST were previously well suited for. TC 282 is given in Table 22 below.

Table 22: Selected Test Case – Improvement to STD Due to Modifications

<table>
<thead>
<tr>
<th>TC</th>
<th>Dim</th>
<th>Objective Function</th>
<th>Constraint Equation(s)</th>
</tr>
</thead>
<tbody>
<tr>
<td>282</td>
<td>10</td>
<td>( U(\bar{x}) = (x_1 - 1)^2 + (x_{10} - 1)^2 + 10\sum_{i=1}^{9} (10 - i)(x_i^2 - x_{i+1})^2 )</td>
<td>n/a</td>
</tr>
</tbody>
</table>

Direct comparisons between standard optimization algorithms and their modified counterparts are given in Table 23 – Table 26.

Table 23: Relative Performance of Standard and Modified STD

<table>
<thead>
<tr>
<th></th>
<th>STD</th>
<th>STD-R</th>
<th>STD-R,S</th>
</tr>
</thead>
<tbody>
<tr>
<td>Accuracy</td>
<td>20.6%</td>
<td>32.1%</td>
<td>65.9%</td>
</tr>
<tr>
<td>Accuracy &amp; Speed</td>
<td>9.5%</td>
<td>26.0%</td>
<td>64.5%</td>
</tr>
<tr>
<td>Standard Dev.</td>
<td>50.7%</td>
<td>38.2%</td>
<td>29.7%</td>
</tr>
<tr>
<td>Single Best Result</td>
<td>31.4%</td>
<td>41.6%</td>
<td>90.9%</td>
</tr>
<tr>
<td>Single Worst Result</td>
<td>64.9%</td>
<td>23.6%</td>
<td>43.6%</td>
</tr>
</tbody>
</table>

Table 24: Relative Performance of Standard and Modified BST

<table>
<thead>
<tr>
<th></th>
<th>BST</th>
<th>BST-R</th>
<th>BST-R,S</th>
<th>BST-SPC</th>
</tr>
</thead>
<tbody>
<tr>
<td>Accuracy</td>
<td>15.9%</td>
<td>46.6%</td>
<td>24.0%</td>
<td>47.3%</td>
</tr>
<tr>
<td>Accuracy &amp; Speed</td>
<td>6.8%</td>
<td>32.4%</td>
<td>15.5%</td>
<td>45.3%</td>
</tr>
<tr>
<td>Standard Dev.</td>
<td>62.5%</td>
<td>25.7%</td>
<td>16.6%</td>
<td>29.1%</td>
</tr>
<tr>
<td>Single Best Result</td>
<td>21.3%</td>
<td>62.5%</td>
<td>75.0%</td>
<td>77.7%</td>
</tr>
<tr>
<td>Single Worst Result</td>
<td>61.8%</td>
<td>24.0%</td>
<td>45.9%</td>
<td>15.5%</td>
</tr>
</tbody>
</table>

Tables Table 23 and Table 24 clearly demonstrate that the modifications significantly improve the performance of the optimization algorithms.

Table 25: Relative Performance of Standard and Modified DN3

<table>
<thead>
<tr>
<th></th>
<th>DN3</th>
<th>DN3-R</th>
<th>DN3-R,S</th>
</tr>
</thead>
<tbody>
<tr>
<td>Accuracy</td>
<td>63.5%</td>
<td>18.2%</td>
<td>21.3%</td>
</tr>
<tr>
<td>Accuracy &amp; Speed</td>
<td>61.5%</td>
<td>18.2%</td>
<td>20.3%</td>
</tr>
<tr>
<td>Standard Dev.</td>
<td>56.4%</td>
<td>32.8%</td>
<td>13.9%</td>
</tr>
<tr>
<td>Single Best Result</td>
<td>50.0%</td>
<td>35.1%</td>
<td>62.5%</td>
</tr>
<tr>
<td>Single Worst Result</td>
<td>27.4%</td>
<td>12.2%</td>
<td>65.9%</td>
</tr>
</tbody>
</table>

Table 26: Relative Performance of Standard and Modified TDE

<table>
<thead>
<tr>
<th></th>
<th>TDE</th>
<th>TDE-R</th>
</tr>
</thead>
<tbody>
<tr>
<td>Accuracy</td>
<td>45.6%</td>
<td>63.9%</td>
</tr>
<tr>
<td>Accuracy &amp; Speed</td>
<td>36.1%</td>
<td>63.9%</td>
</tr>
<tr>
<td>Standard Dev.</td>
<td>56.4%</td>
<td>53.0%</td>
</tr>
<tr>
<td>Single Best Result</td>
<td>54.4%</td>
<td>74.3%</td>
</tr>
<tr>
<td>Single Worst Result</td>
<td>70.6%</td>
<td>51.4%</td>
</tr>
</tbody>
</table>
Although randomization improves TDE, the improvements are less than those in other methods. The standard form of DN3 algorithm, however, is apparently more robust than its modified form.

<table>
<thead>
<tr>
<th>Table 27: Relative Performance of PSO and PRD</th>
</tr>
</thead>
<tbody>
<tr>
<td>PSO</td>
</tr>
<tr>
<td>---------------------------------------------</td>
</tr>
<tr>
<td>Accuracy</td>
</tr>
<tr>
<td>Accuracy &amp; Speed</td>
</tr>
<tr>
<td>Standard Deviation</td>
</tr>
<tr>
<td>Single Best Result</td>
</tr>
<tr>
<td>Single Worst Result</td>
</tr>
</tbody>
</table>

The improvements to PSO are similar to those of TDE in that it does not make PSO more robust relative to other methods (for the Rp value selected in this thesis). However, it makes the method perform much more consistently as shown by the increased standard deviation. Furthermore, the improvements (shift) in PRD accuracy is generally accompanied by superior speed as evidenced by the equal Mean and M&C values.