NAVAL POSTGRADUATE SCHOOL
MONTEREY, CALIFORNIA

THESIS

OPTIMIZING COVERAGE AND REVISIT TIME IN SPARSE MILITARY SATELLITE CONSTELLATIONS: A COMPARISON OF TRADITIONAL APPROACHES AND GENETIC ALGORITHMS

by

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September 2004

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Sparse military satellite constellations were designed using two methods: a traditional approach and a genetic algorithm. One of the traditional constellation designs was the Discoverer II space based radar. Discoverer II was an 8 plane, 24 satellite, Low Earth Orbit (LEO), Walker constellation designed to provide high-range resolution ground moving target indication (HRR-GMTI), synthetic aperture radar (SAR) imaging and high resolution digital terrain mapping. The traditional method designed 9-ball, 12-ball, 18-ball, and 24-ball Walker constellations. The genetic algorithm created constellations by deriving a phenotype from a triploid genotype encoding of orbital elements. The performance of both design methods were compared using a computer simulation. The fitness of each constellation was calculated using maximum gap time, maximum revisit time, and percent coverage. The goal was to determine if one design method would consistently outperform the other. The genetic algorithm offered a fitness improvement over traditional constellation design methods in all cases except the 24-ball constellation where it demonstrated comparable results. The genetic algorithm improvement over the traditional constellations increased as the number of satellites per constellation decreased. A derived equation related revisit time to the number of ship tracks maintained.
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I. TRADITIONAL CONSTELLATION DESIGN

A. HISTORY

1. Origin

Scientists have been studying astrodynamics (the motion of natural and human-made bodies in outer space) since the ancient era. Some definitions of astrodynamics leave out the "natural" portion, establishing the origin of astrodynamics on 10 October 1946 with the launch of the V2 rocket into space or 04 October 1957 with the launch and subsequent orbit of Sputnik 1. However, regardless of the definition, astronomers contributed to the study of astrodynamics beginning about 200 A.D.

The Chaldeans were probably the first to develop astrology. They developed the Saros cycle by measuring the time between eclipses. Focusing on the moon rather than the sun, the Babylonians developed the lunar month by observing the phases of the moon.

The first recognized astrologer was probably Thales of Miletus (c. 640-546 B.C.). Thales determined the length of the year, predicted eclipses, founded the Ionian school of astronomy and philosophy and taught that the world was spherical. Pythagorous (569-470 B.C.) taught astronomy and philosophy. He believed that comets revolved around the sun and the earth rotated about its own axis. Aristarchus (310-250 B.C.) was possibly the first person to suggest the earth rotated around the sun. Eratosthenes (275-194 B.C.) was the first to calculate the radius of the earth accurately.

Hipparchus (c. 161-126 B.C.) developed spherical geometry and also declared the sun to be the center of the universe. In addition, he began cataloging stars based on brightness. Without the aid of instruments, Hipparchus categorized over 1000 stars based on magnitude, separating each star into one of six categories of brightness separated by about 2.5 times the brightness of the previous star. Hipparchus also developed theories of orbital motion.
Claudius Ptolemaceus (100-170 A.D.) continued Hipparchus’ work. Unfortunately he was not exposed to earlier work declaring the sun as the center of the universe. Instead, he published a 13-volume work explaining the motion of celestial bodies around the earth.

There was a large gap of time between Ptolemaceus’ work and the next significant scientific contribution to astrodynamics provided by Nicholas Copernicus (1473-1543).

2. Copernicus

Copernicus proposed a Sun-centered solar system. He also disagreed with Ptolemaceus in some numbers and data and the motion of the planets. Unfortunately, Copernicus' theories were controversial at the time and, consequently, his work was not published until he was near death. It is thought that Copernicus theorized elliptical motion but the sections were not included in his work.

3. Kepler

Johann Kepler (1571-1630) determined how to relate mean and true anomalies in the orbit to time in order to predict future occurrences for planets. Using Copernicus' Sun-centered solar system as a starting point, Kepler developed three laws to describe the kinematics of motion of celestial bodies:

1. The orbit of each planet is an ellipse with the Sun at one focus.
2. The line joining the planet to the Sun sweeps out equal areas in equal times.
3. The square of the period of a planet is proportional to the cube of its mean distance to the Sun.

Kepler printed three books; Astronomia Nova (1609) containing his first two laws, De Cometis (1618) about comets, and Harmonices Mundi Libri V (1619) describing physical motion.

4. Newton

Isaac Newton (1642-1727) discovered the mathematical solution to the dynamics of motion. Newton explained the law of gravitational attraction that
accounts for the elliptical motion of planets using an inverse square law. At Edmond Halley's urging (and funding), Newton published *Philosophiae Naturalis Principia Mathematica* in 1687. In his book, referred to as the Principia, Newton introduced his three laws of motion:

1. Every body continues in its state of rest, or of uniform motion in a right [straight] line, unless it is compelled to change that state by forces impressed upon it.

2. The change of motion is proportional to the motive force impressed and is made in the direction of the right line in which that force is impressed.

3. To every action there is always opposed an equal reaction: or, the mutual actions of two bodies upon each other are always equal and directed to contrary parts.

Newton's Universal Law of Gravitation - in combination with his three laws of motion - allowed scientists to model planetary and satellite motion.

Joseph Louis Lagrange (1736-1813), Leonhard Euler (1707-1783), Carl Gustav Jacob Jacobi (1804-1851) and Henri Poincaré (1854-1912) also made significant contributions to the study of astrodynamics, particularly in modeling three body object interaction. [1]

B. ORBIT DETERMINATION

1. **Classic Orbital Elements (COE)**

   Six orbital elements are used to define a satellite's orbit for positioning and tracking. Once all of the COE are known, the orbit size, shape, orientation and location of the spacecraft in that orbit can be determined.

   a. **Semimajor Axis (a)**

   The semimajor axis is used to determine the size of the orbit. The semimajor axis is half the distance along the long axis of the ellipse around which the spacecraft travels. For circular orbits, the semimajor axis is simply the radius of the circle made by the orbiting satellite.
b. **Eccentricity (e)**

The eccentricity is used to determine the shape of the orbit. The eccentricity is the ratio of the distance between the two foci and the semimajor axis. The eccentricity is 0 for a circle and 0 < e < 1 for an ellipse.

c. **Inclination (i)**

The inclination is used to determine the tilt of the orbit. The inclination is the angle between the fundamental plane of the coordinate system (the equatorial plane in an earth centered system) and the orbital plane. An equatorial orbit has an inclination of 0° or 180°, a polar orbit has an inclination of 90°, a direct or prograde orbit has an inclination of 0° ≤ i < 90°, and an indirect or retrograde orbit has an inclination of 90° < i ≤ 180°.

d. **Right Ascension of the Ascending Node (Ω)**

The right ascension of the ascending node is used to determine the angular orientation of the orbit relative to some principal direction. The right ascension of the ascending node is the angle from the vernal equinox (a line drawn from earth through the sun on the first day of Spring) to the ascending node (the intersection of the orbital plane and the fundamental plane as the spacecraft travels from the Southern Hemisphere to the Northern Hemisphere).

e. **Argument of Perigee (ω)**

The argument of perigee is used to determine the orbital ellipse’s orientation within the orbital plane. The argument of perigee is the angle, along the orbital path, between the ascending node and perigee (the point of the orbit closest to earth).

f. **True Anomaly (ν)**

The true anomaly is used to determine the spacecraft’s location within the orbit. The true anomaly is the angle, along the orbital path, between perigee and the spacecraft’s position vector (from earth’s center to the satellite) measured in the direction of the spacecraft’s motion. [2]

2. **Orbit Selection**

Engineers spend many hours in the design process determining spacecraft orbital parameters. Orbit altitudes are often of primary interest –
based on mission requirements – and are generally grouped into one of four
categories based on the distance from earth the spacecraft will travel while on
orbit.

a. **Low Earth Orbit (LEO)**

The Low Earth Orbit includes all orbits from an altitude of a few hundred kilometers to the Van Allen Radiation Belts (≤ 1500 km). They generally have a period of approximately 90 minutes and have more than ten revolutions per day. Some advantages of LEO orbits are: better resolution for remote sensing, less expensive launch costs, and less power needed to transmit signals. Some disadvantages are: limited viewing area, many satellites needed to achieve continuous global coverage, and shorter spacecraft lifetimes due to drag and orbit decay.

b. **Geosynchronous Orbit (GEO)**

The Geosynchronous Orbit is at an altitude of nearly 36,000 km. GEO orbits have a 24 hour period and revolve with the earth. Some advantages of GEO are: the ability to "stare" at one area of the earth, approximately 1/3 earth coverage with one satellite and global coverage with as few as five satellites (lower satellite construction cost than LEO to achieve the same coverage). Some disadvantages of GEO are: very poor remote sensing resolution, high power needed to transmit to earth, large antennas needed for sensing, and expensive to get to orbit.

c. **Medium Earth Orbit (MEO)**

The Medium Earth Orbit includes all orbits from the Van Allen Belts to an altitude around 30,000 km. However, typically, MEO satellites are at an altitude of just over 20,000 km. They have a 12 hour period with two revolutions per day. MEO orbits have all of the advantages and disadvantages of both LEO and GEO. As compared to LEO, MEO satellites are able to see a greater percentage of the earth at a time and therefore require fewer satellites to cover the globe (The Global Positioning System uses 24 satellites for four-fold global coverage). However, MEO satellites require more power to transmit signals to earth and cannot achieve the same remote sensing resolution as LEO sensors.
d. **Highly Elliptical Orbit (HEO)**

A Highly Elliptical Orbit is one that does not fit well into one of the previous classifications. They are characterized by large values for eccentricity or large differences between the perigee and apogee altitudes. Typically HEO pass through all of the other orbital regimes, for example the Molniya orbit has a perigee of 500 km and an apogee of 39,850 km. Molniya orbits have a 12 hour period with two revolutions per day and an inclination of 63.4°. Satellites in HEO orbits are typically at an inclination of 63.4°; at this inclination the perigee remains fixed, at other inclinations the perigee rotates. The advantage of HEO is a long dwell time at apogee. The disadvantage of HEO is very limited coverage near perigee.

3. **Design Process**

   a. **Establish Orbit Types**

   Step 1 involves examining the four types of orbits. The Earth-referenced and space-referenced orbits are operational orbits (the satellite/s will remain in the orbit for the majority of their lifetime). The transfer and parking orbits are used to get the satellites to their operational orbits

   (1) Earth-referenced orbits are used to cover the earth (The Global Positioning System).

   (2) Space-referenced orbits are used to cover space (the Hubble Space Telescope).

   (3) Transfer orbits are used to transition a satellite from one orbit to another (Hohmann Transfer).

   (4) Parking orbits are used as a transition orbit between the initial and final orbit.

   b. **Determine Orbit-related Mission Requirements**

   Step 2 involves examining the mission requirements. Earth-referenced mission requirements drive the orbit to one of the orbital regimes described above (LEO, MEO, HEO or GEO). For example, a need for high resolution pictures would drive the orbit to LEO but a need for long dwell times would drive the orbit to HEO or GEO.
Three other orbits that must be considered are transfer, parking and reference orbits. Transfer orbits are used to get the spacecraft where it is needed when it is needed. Design of the transfer orbit is generally uncomplicated with cost ($\Delta V$ or propellant) as the driver.

Parking orbits – or storage orbits - are used to provide a spacecraft a place to linger while waiting for transfer into the ultimate orbit destination. Often parking orbits are sparsely populated orbits at an altitude high enough to minimize drag but low enough to retrieve the spacecraft easily. Some typical spacecraft in parking orbits are on-orbit spares, spacecraft being tested following launch and spacecraft waiting for the proper conditions to be met before transferring to the mission orbit.

Mission orbits are either earth referenced or space referenced. Earth-referenced orbits allow the spacecraft sensors to provide coverage of the earth or the space near earth. Space-referenced orbits allow the spacecraft sensors to point toward space. Often, for space-referenced orbits, specific orbit parameters are not crucial.

c. Assess Applicability of Specialized Orbits

Step 3 involves examining whether the unique characteristics of a specialized orbit offers an advantage over traditional LEO, MEO and GEO. Usually the advantage must be significant to offset the added cost usually associated with specialized orbits.

Some specialized orbits include:

- Geostationary. A geosynchronous orbit with inclination and eccentricity approximately zero. Geostationary satellites maintain their relative position fixed over one geographic area.
- Sun-synchronous. The orbit rotates as the earth rotates about the sun. Consequently, the orbit maintains a constant orientation with the sun and allows the satellite to cross the equator at the same local time each pass.
- Polar. A 90° inclination orbit used to provide polar coverage.
- Molniya. A highly elliptical, highly inclined orbit which provides coverage to northern latitudes where geostationary satellites do not.
• Repeating ground track. An orbit for which the ground track of the satellite will repeat itself after one or more days.
• Frozen. Stable orbits with low eccentricity and an argument of perigee of 90° or 270°.
• Super Synchronous. An Earth orbit with a semi-major axis greater than a geosynchronous orbit.

d. **Evaluate Whether a Single Satellite or a Constellation is Needed**

Step 4 involves determining whether a single satellite is sufficient or an entire constellation is required. As a rule, single satellites are less expensive and therefore desired if they can accomplish the mission. However, single satellites are often complex and provide no redundancy (if you lose the satellite you cannot accomplish the mission). A constellation of small inexpensive satellites may be a better solution.

e. **Perform Mission Orbit Design Trades**

Step 5 involves choosing an orbit based on the mission the satellite will be asked to perform. Some considerations when selecting orbits are; coverage, sensitivity or performance, environment and survivability, launch capability, ground communications, orbit lifetime, and legal or political constraints.

Some missions can be completed regardless of altitude. In this case, mission requirements must be weighted based on preference. For example, communications can be done from all altitudes. An adequately populated LEO constellation, using several satellites and relay, could enable communications between two users anywhere on the globe. A GEO satellite constellation of 5 inclined satellites could do the same. The trade may be the cost of populating a LEO constellation versus populating a constellation in GEO.

f. **Evaluate Constellation Growth and Replenishment or Single-satellite Replacement Strategy**

Step 6 involves determining how to keep the constellation useful. Satellite constellations are generally populated a few satellites at a time. Ideally, the constellation will be at least partially serviceable while waiting for the remaining satellites in the constellation to be launched and operational.
Another consideration is how to replace satellites when they fail. Traditionally, either replacement satellites are launched and put into orbit after the satellite fails or an on orbit spare is launched ahead of time and transferred into position to replace the failed satellite.

Finally, at the end of the constellations life (when failed satellites will not be replaced), ideally the system will not be useless with the loss of one satellite. As each satellite fails the quality of service depletes but the customers do not experience a total loss of service.

g. Assess Retrieval or Disposal Options
Step 7 involves determining how to deal with a failed satellite. If the satellite is in LEO, it is deorbited and either breaks up in the atmosphere or is landed in the ocean. In GEO, satellites compete for a finite amount of space and therefore disposal is also important. Typically, GEO satellites are boosted into super synchronous orbits at end of life.

Although rare, retrieval is another option. LEO satellites can be retrieved and refurbished or recovered and brought back down to earth using the space shuttle.

h. Create $\Delta V$ Budget
Step 8 involves calculating the cost for each mission orbit scenario.

i. Determine Launch Options and Cost
Step 9 involves calculating how much it will cost to get the satellites on orbit. More mass equates to more cost. A large satellite requires a large launch vehicle which requires more fuel and is consequently more expensive. Engineers try to design the satellite to be as light as possible to save cost. Often multiple smaller satellites can ride on the same launch vehicle.

Mass being the cost driver, higher altitude constellations are more expensive to populate than lower altitude constellations. To get to GEO, the satellite must first be placed in LEO with enough fuel to transfer to GEO. More fuel means more mass which means higher cost.
C. CONSTELLATION DESIGN CONSIDERATIONS

1. Parameters
   a. Swath Width

   The swath width is the area that can be covered by each individual satellite. The swath width (maximum earth central angle) is a function of altitude and minimum working elevation angle (grazing angle). Assuming a constant minimum working elevation angle, as the altitude increases the swath width increases. Conversely, assuming a constant altitude, as minimum working elevation angle increases the swath width decreases.

   b. Altitude

   Most constellations are designed with all satellites at the same altitude. In this scenario, a uniform relationship between the satellites over time can be maintained. Also, with all satellites at the same altitude (and inclination), the orbit planes maintain their relative orientation.

   c. Inclination

   Considering inclination is important because the inclination impacts how coverage patterns are formed and coverage as a function of latitude.

   d. Node Spacing

   As long as all of the nodes rotate at the same rate, actual location of the ascending node is irrelevant. Some common node spacings include:

   - Equal node spacing over the complete equator
   - Equal node spacing over half the equator
   - Equal node spacing except for a seam between satellites going up and coming down
   - Node spacing adjusted in pairs or triplets

2. Coverage

   Coverage is the principal performance parameter. The amount of coverage is dependent on mission needs. If the mission calls for periodic images of an installation, intermittent coverage might be sufficient. If the mission calls for
uninterrupted communications, continuous coverage by at least one satellite might be required. If the mission calls for triangulating a position, continuous coverage by multiple satellites may be required (GPS).

3. **Number of Satellites**

   The number of satellites is likely the principal cost driver. As illustrated above, more satellites mean more expense. The exception might be a complex, heavy satellite replaced by several simple, light satellites.

4. **Launch Options**

   Launch represents the largest risk. A launch failure may cost hundreds of millions of dollars. Not only is the expensive launch vehicle destroyed, but also the very expensive satellite. Often, several satellites are launched on a single vehicle, bringing the launch failure cost even higher.

5. **Environment**

   Environment has the greatest impact on constellation life. All orbits experience the harsh effects of the space environment. LEO satellites are affected by the Earth's atmosphere. MEO constellations reside inside or just outside the Van Allen Radiation belts and are subjected to harsh radiation effects. GEO constellations are more affected by the solar atmosphere than the earth's atmosphere, but also suffer some affects of the Van Allen Radiation belts.

6. **Stationkeeping**

   The purpose of stationkeeping is to maintain a relative position between satellites or inertial space. The dominant orbit perturbations are atmospheric drag – a function of altitude – and the oblateness of the earth – a function of altitude, inclination and eccentricity. Consequently, LEO constellations pose the greatest stationkeeping challenges as LEO constellations are most affected by atmospheric drag and the Earth's oblateness. If left unchecked, the decay caused by atmospheric drag will ultimately lead to spacecraft reentry.

   To prevent disassociation, satellites are often designed with the same altitude, eccentricity and inclination within the constellation. Also, constellations in eccentric orbits will usually be at the critical inclinations of 63.4° or 116.6° so apogee and perigee do not rotate.
7. **Collision Avoidance**

Collisions represent the largest long term threat to satellites. Debris caused by a collision between two satellites could destroy the entire constellation as each satellite (at the same altitude) eventually passes through the particles scattered by the crash. Consequently the entire system is designed for collision avoidance.

8. **Constellation Build-up, Replenishment, and End-of-Life**

Constellation build-up, replenishment, and end-of-life represent the plan for the constellation health. Constellation build-up is the plan to populate the constellation and concerns such issues as: one satellite at a time, several satellites per launch vehicle, how much of the mission will the constellation perform before fully populated? Replenishment is the method of replacing failed satellites and addresses: on orbit spares, launch on demand, or no replacement. Dead satellites must be removed from orbit to avoid collision. The end-of-life choices are deorbit or raise the satellite to a higher orbit.

9. **Number of Orbit Planes**

The number of orbit planes is important for satellite repositioning. Repositioning within a plane is much more efficient than repositioning to a different plane. Therefore, the fewer the planes the better (assuming the mission can still be accomplished). Usually, higher altitudes require fewer orbital planes.

D. **PATTERNS**

1. **Geosynchronous Constellations**

Geosynchronous constellations are the simplest constellation pattern. Three satellites provide worldwide coverage; only five are needed for continuous global coverage. GEO constellations are used for communications, LEO satellite tracking and gathering data for weather predictions. Geostationary constellations provide continuous coverage over a fixed area of the Earth.

2. **Streets of Coverage Constellations**

A streets of coverage constellation consists of satellites in polar or nearly polar orbits. The right ascensions of the ascending node of the orbit planes are spread evenly around one hemisphere of the earth. In this hemisphere all of the
satellites move northward. In the other hemisphere all of the satellites move southward. There are two seams where the hemispheres meet. At the seams the adjacent satellites are moving in opposite directions. To achieve constant coverage statistics the space between the orbital planes at the seams must be less than the spacing of the planes within the hemisphere. The sensor swath width determines the number of orbital planes and satellites required for global coverage.

Figure 1. Streets of Coverage Constellation [From 3]

3. **Walker Constellations**

Walker constellations are the most symmetric of the satellite patterns. The *Walker Delta Pattern* contains a set number of satellites distributed evenly within a set number of orbit planes at the same inclination. The ascending nodes of the orbital planes are uniformly distributed around the equator and the satellites are uniformly distributed within the orbital planes. Walker constellations are completely symmetrical in longitude, but perhaps their greatest advantage is
that there are a finite number of them and they can all be identified and investigated.

Figure 2. Walker Constellation [From 4]

4. **Elliptical Orbit Patterns**

Elliptical orbit patterns allow designers to optimize coverage to fit specific needs. Specifically, elliptical constellations provide additional free parameters to optimize the constellation. The disadvantage is the satellites must be designed with greater complexity to work at varying altitudes (variations in range, angular size of the Earth's disk, in-track velocity, and relative position for satellites in the same orbit) and components will need extensive radiation hardening.

5. **Other Constellation Patterns**

Some examples of other constellation patterns include:

- The Ellipso Constellation. Elliptical orbits are used to optimize coverage as a function of longitude, latitude, and time of day.
- 2–plane Polar
- 3 Mutually Perpendicular Planes
• 2 Perpendicular Non-polar Planes
• 5-plane Polar "Streets of Coverage"

E. SUMMARY

1. Process
   a. Establish Constellation-related mission requirements
   b. Do All Single Satellite Orbit Trades Except Coverage
   c. Do Trades Between Swath Width, Coverage, and Number of Satellites
   d. Evaluate Ground Track Plots
   e. Adjust Inclination and In-plane Phasing
   f. Review the Rules of Constellation Design
   g. Document Reasons for Choices and Iterate

2. Design Factors
   a. Principal
      (a) Number of Satellites
      (b) Constellation Pattern
      (c) Minimum Elevation Angle
      (d) Altitude
      (e) Number of Orbit Planes
      (f) Collision Avoidance
   b. Secondary
      (a) Inclination
      (b) Between Plane Phasing
      (c) Eccentricity
      (d) Size of Stationkeeping Box [5]
II. GENETIC ALGORITHMS

A. THEORY

Genetic algorithms (GA) use the concepts of natural selection and natural genetics to solve optimization problems. Genetic algorithms use a population of solutions to solve practical engineering optimization problems by estimating a series of unknown parameters within a model of a physical system. [6]

B. HISTORY

John Holland developed the concept of genetic algorithms at the University of Michigan in the 1960s and 1970s. Holland's original intention was to study the mechanisms of adaptation found in nature and to incorporate those mechanisms into computer-simulated systems. Holland, with his students and colleagues, developed a detailed approach to modeling natural evolution in the form of computer algorithms. In 1975, Holland published his book, *Adaptation in Natural and Artificial Systems*, in which he describes the basic approach to population-based search characteristics. [7]

The research in genetic algorithms was mainly theoretical until the early 1980s. During this time, a large amount of work was done with fixed length binary representation in function optimization. Specifically, Kenneth De Jong attempted to capture the features of the adaptive mechanisms in the family of genetic algorithms that constitute a robust search procedure. R. B. Hollstein analyzed the effect that different selection and mating strategies have on the performance of a genetic algorithm.

Throughout the 1980s, genetic algorithm applications were plentiful and diverse. The genetic algorithm community routinely added insight into generality, robustness and applicability of genetic algorithms. Each new insight contributed to improving performance through tuning and specializing the genetic algorithm operators. By the late 1980s, genetic algorithms were successfully applied to optimization problems, scheduling, data fitting and clustering, trend spotting and path finding. [8]
In 1989, David E. Goldberg published his book, *Genetic Algorithms in Search, Optimization, and Machine Learning*. This book marked the second major milestone in the history of genetic algorithms, accelerating the application of genetic algorithms. Goldberg wrote about how genetic algorithms could be used to solve a myriad of problems. He gives examples of researchers applying genetic algorithms to solve various problems. Goldberg also presented the theory of genetic algorithms, giving an unambiguous, succinct definition. Finally, the Pascal source code Goldberg included allowed researchers to experiment with genetic algorithms.

In 1991, Dave Davis further advanced the study of genetic algorithms through his published book, *The Handbook of Genetic Algorithms*. Davis used his book to teach the reader how to implement a genetic algorithm. Davis kept the literature fundamental and did not include theoretical details. Davis also included chapters outlining genetic algorithm applications written by researchers who had successfully used genetic algorithms in their field. In 1991, *The Handbook of Genetic Algorithms* contained the most current state of genetic algorithm application and effectiveness. Davis successfully showed the utility of a properly conceived genetic algorithm in advanced problem solving. Unfortunately, it also showed the lack of industry participation as most of the chapter authors were genetic algorithm specialists applying genetic algorithms to a field of study and not necessarily field specialists using a genetic algorithm to advance their research.

Interest and use of genetic algorithms has grown substantially since Davis' book was published in 1991. Not only have there been multiple genetic algorithm texts published in the last few years, but a well attended biannual international conference met to discuss genetic algorithms. Additionally, genetic algorithm usage increased dramatically as word of the many advantages and applications spread as the volume of publications increased. Karr and Freeman wrote, "The number of publications related to genetic algorithms is not growing, it has virtually exploded over the last decade." [7]
C. PROCESS

A genetic algorithm has five basic components:

1. A genetic representation of solutions to the problem
2. A way to create an initial population of solutions
3. An evaluation function rating solutions in terms of their fitness
4. Genetic operators that alter the genetic composition of children during reproduction
5. Values for the parameters of genetic algorithms [9]

Typically, a genetic algorithm begins with an initial population of individuals. The selection of the initial population is generally random and spread throughout the search space. The individuals, representing a possible solution to the problem, are evaluated based on defined wellness parameters and given a score based on some measure of fitness. Generally, the individuals are represented by binary encoding (strings) where bits are manipulated to create new individuals.

Next, the genetic algorithm transforms using selection, crossover and mutation. Selection alters the genetic algorithm through "picking" the best individual to move on in the process. In selection, the poor performing samples are discarded in favor of the better (fitter) performers. Thereby, the population improves through natural selection similar to biological evolution.

Crossover allows two individuals to reproduce, creating offspring with characteristics of each parent, similar to biological sexual reproduction. Crossover hopes to capture the desired traits of each parent in the offspring creating a fitter individual. One popular crossover method is to choose a breaking point in the binary string and swap genetic information between each individual before or after that point.

Mutation changes the characteristics of the individual without reproduction with another. During mutation, bits are altered at random to produce an individual with different characteristics. Through this process, some mutated
individuals will be fitter and some will be weaker. The fitter individuals are kept in
the population and the weaker individuals are discarded.

When selection, crossover and mutation are complete, a new population is
formed from a fresh generation of individuals. The genetic algorithm continues to
transform generation after generation, using selection, crossover and mutation,
until a set number of generations are met or a convergence point is reached.

D. APPLICATIONS

Genetic algorithms have been used to solve a wide variety of problems.
Genetic algorithms are used primarily in the scientific and engineering industries
to solve complex mathematical problems. However, as exposure to the
advantages genetic algorithms present propagates, uses will become more
creative. Some example applications:

• Optimization. Genetic algorithms have been used in numerical
  optimization and combinatorial optimization problems.

• Automatic programming. Genetic algorithms have been used to
evolve computer programs for specific tasks, and to design other
  computational methods.

• Machine learning. Genetic algorithms have been used to classify
  and predict tasks and to evolve aspects of particular machine
  learning systems.

• Economics. Genetic algorithms have been used to model
  processes of innovation. They have also been used in the
development of bidding strategies and the emergence of economic
  markets.

• Immune systems. Genetic algorithms have been used to model
  somatic mutation during an individual's lifetime. Genetic algorithms
  have also been used in the discovery of multi-gene families during
evolutionary time.

• Ecology. Genetic algorithms have been used to model biological
  arms races, host-parasite co-evolution, symbiosis, and resource
  flow.

• Population genetics. Genetic algorithms have been used in the
  study of population genetics.
• Evolution and learning. Genetic algorithms have been used to determine how individual learning and species evolution affect one another.

• Social systems. Genetic algorithms have been used to study the evolution of social behavior in insects and the evolution of cooperation and communication in multi-agent systems. [11]

• Image registration. Genetic algorithms have been used to generate three-dimensional visualizations of the human body, greatly increasing the scale of the search space.

• Recursive prediction of natural light levels. Genetic algorithms have been used to control artificial lights within buildings to act solely as a supplement to available daylight.

• Water network design. Genetic algorithms have been used as a design tool for water distribution network planning and management.

• Ground-state energy of the ±J spin glass. Genetic algorithms have been used to study spin glasses. Computational methods developed have been applied to questions in computer science, neurology and the theory of evolution.

• Liquid crystals. Genetic algorithms have been used in the estimation of the optical parameters of liquid crystals.

• Energy efficiency. Genetic algorithms have been used in the design of energy-efficient buildings.

• Human judgment. Genetic algorithms have been used as the fitness function of human judgment. The genetic algorithm converges at a useful rate and in the direction of improving subject preference. [1]

This list of applications is long but does not converge on the total number of applications being considered. However, this list does illustrate the diversity of genetic algorithms and gives a glimpse of future applications. The continued success of genetic algorithms in the scientific and engineering communities will pave the way for further experimentation across many disciplines.

E. VOCABULARY

Some terms commonly used when Genetic Algorithms are referenced include:

• Population – The set of individuals, items, or data from which a statistical sample is taken.

• Individuals – Separate and distinct from others of the same kind
• Search Space – The boundaries of the genetic algorithm
• Strings – Lines of binary code, analogous to chromosomes
• Selection – A natural or artificial process that favors or induces survival and perpetuation of one kind of organism over others that die or fail to produce offspring.
• Crossover – An exchange of genetic material between chromosomes.
• Mutation – A change of the DNA sequence within a gene or chromosome of an organism resulting in the creation of a new character or trait not found in the parental type.
• Fitness – The extent to which an organism is adapted to or able to produce offspring in a particular environment.
• Parent – An entity that produces or generates offspring
• Offspring – Something that comes into existence as a result
III. THE TRADITIONAL MODEL

A. BACKGROUND
Most satellite constellations are designed using the traditional methods (Walker, Streets of Coverage, etc.) described in Chapter I. This paper investigates the performance achieved using GA techniques relative to these traditional methods.

B. SPACE BASED RADAR (SBR)
SBR offers sufficient complexity to strenuously test the two constellation design methods for comparison.

1. Imaging radar
The two main advantages of radar imaging over visual imaging sensors are 24 hour capability (radar can "see" equally well in daylight and darkness) and all weather capability (radar can "see" through clouds). Another advantage radar has over other sensors is that radar can penetrate slightly beneath the surface of the earth (mine detection capability).

   a. Basics
   An SBR satellite moving through space in orbit sends microwave radiation pulses through its antenna at the speed of light. The pulses are directed in the range, look or across-track direction. Figure 1 illustrates the following definitions:

   • Slant Range – the line-of-site distance measured from the antenna to the target
   • Ground Range – the horizontal distance measured along the surface from the ground track to the target
   • Near Range – the area closest to the ground track at which a radar pulse intercepts the terrain
   • Far Range – the area of pulse termination farthest from the ground track
   • Depression Angle ($\beta$) – the angle measured from a horizontal plane downward to a specific part of the radar beam
   • Look Angle ($\theta$) – the angle measured from a vertical plane upward to a specific part of the radar beam
When measured to the same part of the beam, the depression angle and the look angle are complementary angles ($\beta + \theta = 90^\circ$).

- **Incidence Angle ($\phi$)** – the angle measured between the axis of the radar beam and a line perpendicular to the local ground surface that the beam strikes
- **Grazing Angle ($\gamma$)** – the complement of the incidence angle

Consequently, the incidence angle and the grazing angle are a function of both the illumination angle ($\beta$ or $\theta$) and the slope of the terrain. When the terrain is horizontal, the depression and grazing angles are equal ($\beta = \gamma$) and the look and incidence angles are equal ($\theta = \phi$).

- **Resolution** – the minimum separation between two objects of equal reflectivity that will enable them to appear individually in a processed radar image
- **Pulse Rectangle** – the surface area covered by the energy radiated from the sensor

When two or more objects fall within the same pulse rectangle they cannot be resolved as separate entities. Rather, they are presented as one echo to the radar system. If objects are separated by a distance exceeding the corresponding dimension of the pulse rectangle, they will be imaged separately.

- **Range Resolution** – determines resolution cell size perpendicular to the ground track
- **Azimuth Resolution** – establishes the cell size parallel to the ground track
b. **Detection**

Radar detection is defined as any object that reflects enough energy to be distinguished from the background noise by the receiver (a blip on the scope). Objects are categorized based on their ability to reflect microwave radiation. Highly reflective objects create large radar signatures. Flat metal surfaces produce large signatures; a significant portion of the microwave radiation is reflected back to sensor. Objects with multiple surface angles
produce small signatures; most of the microwave radiation is reflected away from the sensor.

c. **Range Resolution**

Range resolution is determined by the length of the emitted microwave pulse (pulse length). Pulse length is determined by multiplying the pulse duration (τ) by the speed of light.

Two objects will appear as one unless all parts of their reflected signals reach the radar sensor at different times. Consequently, objects must be separated by a slant-range distance greater than one half of a pulse length to be seen as separate entities.

Ground range resolution is half the pulse length divided by the cosine of the depression angle. Therefore, ground range resolution can be improved by increasing the distance from the ground track and by shortening the pulse length.

d. **Signal Shape**

Target resolution is determined based on the pulse length (“t” in the figure below). Related to pulse length, pulse repetition interval (PRI, “T” in the figure below) is the interval between pulses. As illustrated, the PRI duration is generally much longer than the pulse length. The pulse repetition frequency (PRF) is the inverse of the PRI.

![Figure 4. Radar Pulse](From 13)
Pulse length is also related to spectrum. Range resolution is proportional to the length of the pulse. Essentially, a short pulse length contains a wide spectrum and a long pulse length is restricted to a narrow spectrum.

One solution to the pulse vs. spectrum conundrum is using frequency differential. By modulating the frequency of the pulses and monitoring the frequencies of the returns, the two objects can be discerned even if they overlap in time.

e. Azimuth Resolution

Beam width is determined by antenna size and wavelength. Azimuth, or along-track, resolution is a function of the beam width. The beam width increases with range, therefore the greater the range the poorer the azimuth resolution. Two objects at the same range within the beam will appear as one because their returns will be received at the same time. Therefore, to distinguish between two objects, their ground separation distance in azimuth must be greater than the width of the radar beam.

f. Resolution

Azimuth resolution is the slant range multiplied by the wavelength divided by the length of the antenna. Therefore, azimuth resolution improves as range decreases, antenna length increases and wavelength decreases. To improve azimuth resolution, use a long antenna, a short operating wavelength, a close-in interval, or a combination of these factors. The problems are antenna size is limited and the all weather capability of radar is reduced when wavelength is less than 3 cm. The solution is Synthetic Aperture Radar (SAR).

2. Synthetic Aperture Radar (SAR)

SAR uses the motion of the emitting vehicle (the satellite) to increase the length of the effective antenna. The SAR vehicle carries a relatively short antenna and intercepts the emitted signal at different paths along the flight as shown in Figure 5.
The longer resulting antenna is simulated by using the coherence of radar signals. The main beam footprint is then twice the altitude of the sensor multiplied by the wavelength and divided by the antenna length. The array footprint on the ground is half the length of the antenna.

SAR sensors generally have two modes of operation: scan and spotlight. In scan mode, the SAR antenna is pointed in a fixed direction; the only motion is the motion of its platform (aircraft or satellite). Scan mode allows imaging of a large area with fixed resolution. In spotlight mode, the SAR antenna is articulated to continuously point at a specific location or target. Spotlight mode restricts the area of an image, but provides greater resolution.

The resolution of an object is proportional to the time it is in the radar beam. That period of time increases with range and therefore azimuthal resolution is range independent for scan mode. The along-track, linear resolution increases (gets worse) with range since the beam has a fixed angular extent. However, this is compensated for by the fact that as the beam diverges with increasing range, any target at a more distant slant range spends more time in the beam. The net effect is a fixed, "range independent" resolution.

In spotlight mode, antenna length is arbitrarily large because the interval at which radar energy is returned from the target is determined by the operator. As R. C. Olsen points out in reference 10:
In processing, the azimuth details are determined by establishing the position-dependent frequency changes or shifts in the echoes that are caused by the relative motion between terrain objects and the platform. To do this, a SAR system must unravel the complex echo history for a ground feature from each of a multitude of antenna positions.

For example, if we isolate a single ground feature, the following frequency modulations occur as a consequence of the forward motion of the platform:

Positive Doppler - the feature enters the beam ahead of the platform and its echoes are shifted to higher frequencies

Zero Doppler – the platform is perpendicular to the features position and there is no shift in frequency

Negative Doppler - the platform moves away from the feature, the echoes have lower frequencies than the transmitted signal.

The Doppler shift information is then obtained by electronically comparing the reflected signals from a given feature with a reference signal that incorporates the same frequency of the transmitted pulse. The output is known as a phase history, and it contains a record of the Doppler frequency changes plus the amplitude of the returns from each ground feature as it passed through the beam of the moving antenna. [12]

3. Ground Moving Target Indicator (GMTI)

In contrast to SAR imaging, GMTI detects the Doppler shift in frequency caused when the radar pulse is reflected by a moving object. This Doppler shift enables a GMTI sensor to rapidly distinguish between a moving object and the stationary background. GMTI is able to distinguish moving targets from water and surface clutter over large areas in all weather and in darkness.
The figure above is a fusion of a SAR image and GMTI data. The SAR image (shades of green) enables the terrain to be seen; this figure is of a beach area. The GMTI data displays as the white tracks, in this figure they are LCACs. The GMTI picture is presented in real time allowing simple track correlation. The GMTI picture can also be correlated with other intelligence sources and fed into the Common Operating Picture (COP) of the dynamic battlefield.

GMTI requires low data rates relative to a SAR image. This enables the GMTI picture to be transmitted to ground stations over a wide area.

There are two types of GMTI radars. Static radars provide a moment in time picture of all objects within its field of view that are moving. Static radars are used to give a picture, with infrequent updates, depicting moving target density. Static radars do not have the capability to develop and maintain a track on a single target. Dynamic radars provide continuous or nearly continuous coverage. This frequent coverage allows dynamic radars to provide precise tracking of a moving target.

A GMTI system that can detect, locate and track slow moving ground targets will be able to provide detailed, near-real time information on vehicular movements such as the vehicle length and the order of vehicles within a convoy.
To do this, a GMTI system must be able to generate and maintain numerous tracks automatically using the following metrics as listed in reference [15]:

- **Probability of Detection** – the probability of detecting a given target at a given range any time the radar beam scans across it
- **Target Location Accuracy** – a function of platform self-location performance, radar pointing accuracy, azimuth resolution, and range resolution
- **Minimum Detectable Velocity** – the rate of movement determining whether the majority of military traffic will be detected
- **Target Range Resolution** – the fidelity determining whether two or more targets moving in close proximity will be detected as individual targets
- **Stand-off Distance** – the distance separating a radar system from the area it is covering
- **Coverage Area Size** – the area the system can keep under continuous surveillance from a specific orbit. [See Figure 3 below]. The earth's curvature and screening from terrain, foliage, and buildings cause system altitude to be a key factor determining coverage area—the higher the altitude, the greater the coverage.
- **Coverage Area Revisit Rate** — the frequency with which the radar beam passes over a given area

![Joint STARS Coverage In Operation Desert Storm](From 15)
The accuracy of the GMTI picture is dependent on the system's performance in these matrices. Poor performance can lead to inaccuracies in both location and timeframe of the targets position. Precise positioning of slow moving vehicles is crucial for targeting. The advantage of using radar is the system can often counter the effects of enemy camouflage, concealment and deception.

Most GMTI systems can collect SAR and GMTI information (but usually not both at the same time). Using SAR still images with GMTI can be valuable in ascertaining whether a target has stopped moving or has been lost. Using GMTI and high-resolution SAR pictures allow target characterization by Automatic Target Recognition (ATR) algorithms. [15]

4. Future

The Air Force is the executive agent for space. As such they are tasked with providing an effective cross service SBR system.

Ideally, the new arrangement will amalgamate all SBR requirements into one SAR/GMTI capable system with the next decade. The SBR system goal will be to provide integrated data without the user knowing the source. The new program was to provide initial SBR capability by 2010.

a. The Requirements Dilemma

The challenge with any system is to integrate all service's requirements into one efficient design. The primary challenge is determining how the system's information will get to the battlefield commanders (an arguing point that has led to the demise of systems in the past). Another challenge is determining whether the SBR constellation will be flown in LEO (better operational access for GMTI but requires more hand-offs) or MEO (simultaneous target access by two satellites but long SAR integration times).

Several technologies have advanced since initial plans for a SBR constellation were scrapped, including active electronically scanned arrays and synthetic aperture radar.
b. From Scratch

The SBR constellation will be developed from the ground up. This means the constellation will be developed without using a past or existing constellation as a model. The SBR system was designated by Martin R. Sambur, Air Force acquisition chief, a "pathfinder" program; employing new, innovative and streamlined acquisition strategies. The SBR system is required to use the spiral development acquisition strategy, making incremental improvements to the system during development. The SBR will also be the first program developed under the pan-agency space hardware acquisition system.

SBR most likely will consist of 20 to 25 light, inexpensive satellites in LEO, MEO or a LEO/MEO orbit. The physical characteristics are still in the early stages of development. [16]

b. Reality

As 18 June 2004, the 108th Congress met and again all but cancelled the SBR program. The DoD Appropriations Committee directed the Air Force to fundamentally restructure the program. Some Committee findings are below:

The Air Force requested $327,732,000 for the Space Based Radar program. The Committee recommends $75,000,000, a reduction of $252,732,000, and directs that the Air Force fundamentally restructure the program to meet the concerns addressed below.

The Space Based Radar (SBR) program is intended to provide near continuous, global radar imagery and surface moving target indication (SMTI) as well as high resolution terrain information. Advocates describe the program as a key contributor to achieving "global persistent surveillance". Though the pursuit of persistent surveillance is a noble goal, the Committee believes the Space Based Radar program as currently structured:

--Is neither affordable nor likely to produce the results claimed by its advocates, within any reasonable definition of cost, technical challenge, or risk.

--Would consume a disproportionate share of resources from within an already highly stressed DoD space and surveillance budget;
Committee Views and Recommendations- In summary, in and of itself the SBR development program is fraught with enough uncertainties to call into question its viability. Indeed, even under the Administration's own plans the SBR program of record is underfunded in the current Future Year Defense Program by $2 billion, a shortfall resulting from the Department's unwillingness to fully fund this program. The Committee sees little prospect of this changing in light of the other fiscal challenges confronting the Department. These include the well-documented ‘procurement bow-wave’; this Administration's emphasis on missile defense and other transformational programs; and now, and most importantly, the as-yet-unbudgeted future manpower, operational, and equipment recapitalization requirements stemming from operations in Iraq and the Global War on Terrorism. The Committee concludes that against these demands, SBR simply cannot be afforded budget priority.

Without a new approach, the Committee sees little future for the Space Based Radar program. Accordingly, the Committee recommends $75,000,000, a reduction to the request of $252,732,000. These funds are provided to redirect the Air Force's development efforts towards technologies and concepts that would lead to program costs far lower than currently conceived. The focus should be on seeking breakthroughs that fundamentally change the cost-benefit equation for a space based radar system. [17]

C. DISCOVERER II

Discoverer II was chosen as a model for comparison because the Discoverer II system was the most complete SBR design.

The Discoverer II Joint Program was a joint venture between the National Reconnaissance Office (NRO), Defense Research Projects Agency (DARPA) and the U.S. Air Force. Discoverer II was designed to provide high-range resolution ground moving target indication (HRR-GMTI), synthetic aperture radar (SAR) imaging and high resolution digital terrain mapping data collection capabilities. The system was designed to allow joint task force commanders the ability to task the assets and receive data via downlink directly to theater ground stations.
1. **Background**

Discoverer II was to be a pioneer in the ability to detect and track moving targets on the earth's surface from space. Additionally, Discoverer II was designed to produce high-resolution imagery and collect precision Digital Terrain Elevation Data (DTED).

A successful Discoverer II program would have revolutionized the coverage and timeliness of ISR under the direct control of commanders. Additionally, a goal of the Discoverer II program was to demonstrate the ability to populate a constellation with satellites for under $100 million, and to reduce the 20 year life-cycle cost to less than $10 billion.

2. **Capabilities**

The Discoverer II program strived to improve the reconnaissance and surveillance support provided to the theater commanders. To this end, the Discoverer II engineers sought SBR capabilities that would complement, enhance and improve on the existing capabilities airborne assets, such as Joint Surveillance Target Attack Radar System (JSTARS), U-2s and unmanned aerial vehicles (UAVs), provide. With Discoverer II assistance, leaders envisioned providing U.S theater commanders the "information superiority" necessary to employ the smaller, lighter, highly mobile and often widely dispersed joint forces of the future.

The Discoverer II would have been capable of the following:

- Assured, on-demand reconnaissance of areas of interest worldwide during peacetime, periods of heightened tension, hostilities and operations other than war.
- Near-continuous surveillance of selected ground and maritime objects.
- Coverage of blind spots, or "shadowed" areas, obscured from view of standoff airborne platforms.
- Rapid acquisition and tracking of mobile, time-critical targets throughout the depth of the theater.
- Precise geolocation of objects to support munitions targeting.
- High-quality terrain mapping.
To accomplish these goals, the Discoverer II program would have required a large constellation with a large number of inexpensive satellites with the following capabilities:

- Day/night, all-weather, near-continuous, global GMTI search/track, and high resolution imagery,
- Theater dynamic tasking of space-based, or "overhead," GMTI and imagery collection,
- Near-real-time, direct downlink to theater of overhead GMTI and imagery collection; and,
- Collection of precision Digital Terrain Elevation Data (DTED).

Additional requirements, and costs, would have been in-theater infrastructure and additional manning as well as the ability to use the same tactical ground stations used by airborne collectors.

3. General Characteristics

- Function: HRR-GMTI, HR-SAR, HR digital mapping data collection
- Constellation Size: 24 satellites
- Power: Solar panels
- Launch Vehicle: MLV or HLV (2+ per launch vehicle)
- Mass: 1500 kg
- Frequency: X band (10 GHz)
- Antenna Type: 2D, electronically scanned array (ESA)
- Antenna Dimension: 40 m²
- Altitude: 770 km
- Design Life: 10 years
- In-Theater C²: Dynamic, on-demand collection tasking; near real-time, direct downlink of mission data
- Theater Ground Station: Common Imagery Ground/Surface Station (CIG/SS) compliant
- In-Theater Communications: Modified CDL (548 Mbps)
- Reach-Back CONUS Communications: Relay and/or satellite cross-links [18]
4. Reality

From House Rpt.106-644 - DEPARTMENT OF DEFENSE
APPROPRIATIONS BILL, 2001:

The Air Force, Defense Advanced Research Projects Agency, and the National Reconnaissance Office collectively requested $130,000,000 for the Discoverer II satellite technology demonstration program. The Committee recommends no funding, a decrease of $130,000,000. The fiscal year 2000 Defense Appropriations Act provided sufficient funding for the Discoverer II program to conclude the phase I studies and analysis portion of the program along with related risk reduction efforts. With phase I now funded to completion, the Committee recommends that the Discoverer II program be terminated.

The Committee makes this recommendation for the following reasons: (1) Discoverer II has no documented requirement or concept of operations; (2) the cost of the engineering and manufacturing development phase of the program, which the program office estimates at $702 million and which will in all likelihood exceed $1 billion, is of a magnitude ordinarily associated with the development of fully operational satellites and therefore unaffordable given the limited operational benefits of a technology demonstration program; (3) the Department has conducted no trade-off analysis between Discoverer II and other systems and processes that could deliver ground moving target indication data to warfighters; and, (4) the Department has failed to analyze the impact a Discoverer II constellation would have on an already overtaxed imagery processing, exploitation and dissemination system.

Even if successful, there is no guarantee the Air Force could ever build, launch, operate and maintain a Discoverer II constellation without a substantial top line increase to its budget. By some estimates the cost of a fully functional Discoverer II constellation could reach $25 billion. In the face of other severe shortfalls in space and aircraft modernization the Committee concludes that Discoverer II is of low priority and recommends its termination.

The Committee discusses its recommendation more fully in the classified annex to this report. [19]
D. THE MODEL

The Discoverer II constellation was modeled using Satellite Tool Kit for demonstration and comparison with the Genetic Algorithm constellation. The following proposed Discoverer II system parameters were used:

- Total Satellites: 24
- Number of Planes: 8
- Satellites/Plane: 3
- Type of Orbit: Walker
- Inclination: 53°
- Altitude: 770 km
- Latitude Coverage: 65°N to 65°S
- Grazing Angle: 12°
- Slope Angle: 70°
- Cone Angle: 45°
- Response Time: 90%/15 min*

*90% of imaging requests result in collection within 15 minutes

- Modes: National Image Interpretability Rating Scale (NIIRS)-3 (3 meter/360,000 km²/hr)
  NIIRS-5 (1 meter/27,000 km²/hr)
  (NIIRS-7) Point/Spot (.3 meter/4 km², 60/hr)
  MTI (3 meter/360,000 km²/hr)

Collection Rates:

- HR-GMTI: Minimum Detectable Velocity (MDV) = 4 kph
  Maximum Velocity = 100 kph
  2,000,000 km²/hr (600 km²/sec)

- STRIP SAR/GMTI: 3 meter impulse response (IPR) imagery
  700,000 km²/hr

- SCAN SAR: 1 meter IPR
  100,000 km²/hr

- SPOT SAR: 0.3 meter IPR
IV. THE GENETIC ALGORITHM

Chapter II examined the fundamentals of genetic algorithms. This chapter will focus on the specific implementation used for this thesis. First, some of the concepts presented earlier will be reviewed and expanded in light of the developed application. Following that, the major modules of the program will be examined with an example to illustrate how the genetic algorithm operates. A discussion of the interface with Satellite Tool Kit (STK) will follow. Finally, the chapter will conclude by suggesting follow-on research in implementing a genetic algorithm for constellation design.

A. BACKGROUND

A genetic algorithm is a search tool that leverages biologic, genetic, and evolutionary processes. Although each implementation of a genetic algorithm models biologic processes, the granularity of the model varies widely. Most implementations use binary encoding, in which the variables optimized are represented as binary strings. Other encoding methods are available, including Gray encoding and whole number encoding. Our main objective was to test the usefulness of genetic algorithms as a constellation design tool, binary encoding was chosen for the simplicity of implementation.

Most biological organisms, including all animals, are diploid, having paired chromosomes. In the process of reproduction, the offspring receives one chromosome of the pair from the mother and the other from the father. Typically, genetic algorithms are haploid, having only single chromosomes. Reproduction is the result of crossover between the mother’s and father’s chromosomes. Genetic algorithms are rarely constructed to be diploid, because it requires the user to handle the genetic concept of dominance. In biologic genetics, a dominant gene is expressed in the heterozygous or homozygous instance; in other words, if it is present in either or both of the paired chromosomes. The recessive gene is only expressed in the homozygous instance. Dominance enables greater genetic diversity to be maintained in a population, because it is possible for the recessive gene to be passed on to future generations without
having it expressed. This leads us into a quick explanation of genotype and phenotype. Genotype is the sum total of the genetic material present in an organism; all of the possible genes that could be passed on to offspring. Phenotype is the expressed or visible manifestation of an individual’s genotype. A non-haploid genetic algorithm would more closely model biologic genetic processes and provide greater genetic diversity. However, how can you model a diploid organism in a binary encoded genetic algorithm and resolve the dominance problem? Which instance, zero (0) or one (1), should be dominant in the calculation of the phenotype? The solution arrived at elegantly achieves increased genetic diversity without creating a dominance problem: triploid chromosomes. By using a three-chromosome genotype, the individuals have greater genetic diversity and the phenotype is calculated by the instance, zero (0) or one (1), in the majority in the genotype chromosomes. The problem with triploid individuals is: how do you get three chromosomes from two parents? The simple answer is you don’t; you have three parents. Therefore, the genetic algorithm would use binary encoding and a phenotype derived from a triploid chromosome genotype.

The final decision was how many chromosome triplets should be used to represent a constellation. It was a choice between using long chromosomes to represent an entire constellation, medium length chromosomes to represent each satellite in a constellation, or short chromosomes representing each orbital element of each satellite in a constellation. Also implicit in this step was deciding what dynamic range was required for each orbital element; this would affect the length of the chromosomes. Here, the programming language, Visual Basic, helped narrow the decision. Visual Basic supports bit-wise comparisons of numeric data types. The bit-wise comparison operators are used to calculating the phenotype from the triploid genotype and to effect crossover and mutation. Three integer data types in Visual Basic supported bit-wise comparisons. The long data type is a 32-bit number and can represent integer values from -2,147,483,648 to 2,147,483,648. The second data type is integer. It is a 16-bit number, representing values from -32,768 to 32,767. The final data type is byte,
8-bits long and represents values from 0 to 255. The first two data types were deemed too granular in this application. For example, with the integer data type, if the possible range of inclinations is 0 to 180 degrees, each step of the data represents a change in inclination of 0.00274658203125 degrees. Such a small difference cannot significantly affect a difference in the performance of a constellation and most certainly cannot be achieved or maintained by a satellite in orbit. Therefore, all of the orbital elements were represented by the byte data type, which equates to an 8-bit binary string.

Of the six classical orbital elements, only four needed to be represented. Synthetic Aperture Radar requires the altitude to remain relatively constant. For this reason, all of the orbits evaluated had an eccentricity of zero. With an eccentricity of zero, the argument of perigee no longer has meaning. Of the four remaining orbital elements, the altitude and inclination of all satellites in a constellation would be equal, but could vary between constellations. This limitation was placed on the search space to guarantee that the constellation would remain constant over time with no differential node rotation, to enable data fusion, and to ensure the constellation could be populated with identical satellites.

B. THE GENETIC ALGORITHM

Now that we understand what orbital elements are represented and how they are encoded, an 8-bit phenotype for each orbital element of each satellite derived from an 8-bit triploid genotype, let us examine the genetic algorithm itself. First, an overview of the basic flow of the genetic algorithm will be presented. A more detailed explanation of the major sub routines with a simplified example will be presented. The simplified example will help to illustrate the major concepts. The entire genetic algorithm code is included as Appendix A.

1. Overview

As presented before, every genetic algorithm has certain characteristics: an data encoding scheme; an initial population of individuals; a method for evaluating the fitness of the individuals; use of genetic operators (crossover, mutation, selection); a means of creating offspring; an iterative loop that
continues until the termination criteria is met. The application of these basic characteristics is what differentiates one genetic algorithm from another. The figure below illustrates the main execution loop of the genetic algorithm used in this study.

2. Create Initial Population

The genetic algorithm first must create a starting population. In all trials conducted as part of this study the initial population size was twelve constellations. The initial population was created by assigning a randomly generated 8-bit value to each genotype chromosome. This was repeated for each orbital element of each satellite in the constellation. In the example, a three individual population will be created. Each individual in the example will consist of 4-bit triploid chromosomes. The following three individuals were created by using a standard 25-cent random binary number generator.

<table>
<thead>
<tr>
<th>Individual</th>
<th>A</th>
<th>B</th>
<th>C</th>
</tr>
</thead>
<tbody>
<tr>
<td>Genotype 1</td>
<td>0100</td>
<td>1001</td>
<td>0111</td>
</tr>
<tr>
<td>Genotype 2</td>
<td>0011</td>
<td>0111</td>
<td>1110</td>
</tr>
<tr>
<td>Genotype 3</td>
<td>1011</td>
<td>1101</td>
<td>1101</td>
</tr>
</tbody>
</table>

3. Calculate Phenotype
Once the genetic algorithm assigned values to the genotype chromosomes, it derived the phenotype. The phenotype is simply whichever value is in the majority for each bit. This was accomplished through a combination of the bit-wise AND and OR operators. See the truth table below for a summary of the bit-wise operators.

<table>
<thead>
<tr>
<th>Bit 1</th>
<th>Bit 2</th>
<th>AND</th>
<th>OR</th>
<th>XOR</th>
</tr>
</thead>
<tbody>
<tr>
<td>0</td>
<td>0</td>
<td>0</td>
<td>0</td>
<td>0</td>
</tr>
<tr>
<td>0</td>
<td>1</td>
<td>0</td>
<td>1</td>
<td>1</td>
</tr>
<tr>
<td>1</td>
<td>0</td>
<td>0</td>
<td>1</td>
<td>1</td>
</tr>
<tr>
<td>1</td>
<td>1</td>
<td>1</td>
<td>1</td>
<td>0</td>
</tr>
</tbody>
</table>

Table 1. Bitwise Comparison

Here are the original triploid genotypes for our example. We will walk through the bit-wise comparison for individual A to derive the phenotype as an example of the genetic algorithm code.

Phenotype = \{[(Genotype 1 AND Genotype 2) OR (Genotype 2 AND Genotype 3)] OR (Genotype 3 AND Genotype 1)\}

Phenotype = \{[(0100 AND 0011) OR (0011 AND 1011)] OR (1011 AND 0100)\}

Phenotype = \{[0000 OR 0011] OR 0000\}

Phenotype = 0011

<table>
<thead>
<tr>
<th>Individual</th>
<th>A</th>
<th>B</th>
<th>C</th>
</tr>
</thead>
<tbody>
<tr>
<td>Genotype 1</td>
<td>0100</td>
<td>1001</td>
<td>0111</td>
</tr>
<tr>
<td>Genotype 2</td>
<td>0011</td>
<td>0111</td>
<td>1110</td>
</tr>
<tr>
<td>Genotype 3</td>
<td>1011</td>
<td>1101</td>
<td>1101</td>
</tr>
<tr>
<td>Phenotype</td>
<td>0011</td>
<td>1101</td>
<td>1111</td>
</tr>
</tbody>
</table>

4. **Evaluate Fitness**

Once the phenotype is derived from the genotype, the genetic algorithm evaluates the fitness of each constellation. This evaluation occurs in three parts. First, the genetic algorithm sends commands to STK to construct a constellation based on the derived phenotypes. STK evaluates the percent coverage, maximum revisit, and average gap figures of merit of the constellation and returns these values to the application. From these performance statistics, the genetic algorithm calculates a fitness score for the constellation. To optimize
the performance of the constellation, we desire to minimize the maximum revisit and average gap time and maximize the percent coverage. To drive the fitness score to a meaningful value, it was necessary to subtract the coverage percentage from one. This enabled the genetic algorithm to converge toward an optimal solution by minimizing the fitness score. In our trivial example to parallel the genetic algorithm, the fitness will be the number of ones in the phenotype, optimizing to a minimum.

<table>
<thead>
<tr>
<th>Individual</th>
<th>A</th>
<th>B</th>
<th>C</th>
</tr>
</thead>
<tbody>
<tr>
<td>Genotype 1</td>
<td>0100</td>
<td>1001</td>
<td>0111</td>
</tr>
<tr>
<td>Genotype 2</td>
<td>0011</td>
<td>0111</td>
<td>1110</td>
</tr>
<tr>
<td>Genotype 3</td>
<td>1011</td>
<td>1101</td>
<td>1101</td>
</tr>
<tr>
<td>Phenotype</td>
<td>0011</td>
<td>1101</td>
<td>1111</td>
</tr>
<tr>
<td>Fitness</td>
<td>2</td>
<td>3</td>
<td>4</td>
</tr>
</tbody>
</table>

5. Scale Fitness

In the first generation, the fitness scores are stored in an array from lowest score to highest, best fitness to worst. In every other generation, the fitness is compared to the stored scores and, if lower, is stored. All other scores cascade down one position and the worst score drops off. Once the entire generation has been evaluated against the best fitness scores, the array is scaled in preparation for the roulette wheel selection of parents. The process requires each fitness score to be weighted, normalized and finally scaled. Let us walk through this process with our example.

<table>
<thead>
<tr>
<th>Individual</th>
<th>A</th>
<th>B</th>
<th>C</th>
</tr>
</thead>
<tbody>
<tr>
<td>Genotype 1</td>
<td>0100</td>
<td>1001</td>
<td>0111</td>
</tr>
<tr>
<td>Genotype 2</td>
<td>0011</td>
<td>0111</td>
<td>1110</td>
</tr>
<tr>
<td>Genotype 3</td>
<td>1011</td>
<td>1101</td>
<td>1101</td>
</tr>
<tr>
<td>Phenotype</td>
<td>0011</td>
<td>1101</td>
<td>1111</td>
</tr>
<tr>
<td>Fitness</td>
<td>2</td>
<td>3</td>
<td>4</td>
</tr>
<tr>
<td>Weighted Fitness</td>
<td>4.5</td>
<td>3</td>
<td>2.25</td>
</tr>
<tr>
<td>Normalized Fitness</td>
<td>0.46</td>
<td>0.31</td>
<td>0.23</td>
</tr>
<tr>
<td>Scaled Fitness</td>
<td>0.46</td>
<td>0.77</td>
<td>1</td>
</tr>
</tbody>
</table>

Weighted Fitness = Fitness Summation / Individual Fitness
Normalized Fitness = Weighted Fitness / Weighted Fitness Summation
Scaled Fitness = Normalized Fitness + Previous Scaled Fitness
6. **Crossover Engine and Crossover**

The final step prior to selecting parents and generating the next generation of offspring is crossover. In haploid genetic algorithms, the process of crossover is used to create offspring. However, in this triploid genetic algorithm, crossover is an operation that occurs between genotype chromosomes of an individual. Each 8-bit chromosome has seven possible crossover loci. The algorithm generates a random number and compares this to the crossover probability for each crossover locus. If the random number is less than the crossover probability, crossover occurs at that locus between that chromosome and one of the other two. To ensure that crossover can take place between any and all of the genotype chromosomes, the genetic algorithm cycles through the genotypes. A crossover indicated in the first genotype occurs between chromosomes one and two; in the second genotype, the crossover occurs between the chromosomes two and three; and in the third genotype, the crossover occurs between chromosomes three and one. Because the orbital parameters of the constellations evaluated by STK are based on the phenotype, crossover has no effect on the fitness of a constellation. It does not change the derived phenotype because it does not change the overall majority of bits; rather it only changes the genotype chromosomes. Although crossover has no effect on the constellations fitness, it does affect the offspring it creates. By changing the genotype chromosomes, crossover affects the genotype string passed from parent to child. This enables a parent that displays one trait to pass the genetic material for another trait onto its child. The constellations that have the highest fitness are subject to crossover every generation. This enables a greater genetic diversity in the possible offspring. Genetic diversity is the single greatest protection against early convergence to a local optimum before thoroughly exploring the search space. The process of crossover is accomplished through bit-wise comparisons. A crossover mask is generated which effectively splits the two genotypes at the crossover locus. The tails are then switched between the genotypes and are then rejoined to the head to form two new genotype chromosomes. In our trivial example, we will set the crossover probability at 0.5.
We will walk through one example of using the crossover masks in the bit-wise comparison.

<table>
<thead>
<tr>
<th>Individual</th>
<th>A</th>
<th>B</th>
<th>C</th>
</tr>
</thead>
<tbody>
<tr>
<td>Genotype 1</td>
<td>0100</td>
<td>100/1</td>
<td>011/1</td>
</tr>
<tr>
<td>Genotype 2</td>
<td>0011</td>
<td>011/1</td>
<td>1/11/0</td>
</tr>
<tr>
<td>Genotype 3</td>
<td>10/11</td>
<td>1/101</td>
<td>1/10/1</td>
</tr>
<tr>
<td>Phenotype</td>
<td>0011</td>
<td>1101</td>
<td>1111</td>
</tr>
<tr>
<td>Fitness</td>
<td>2</td>
<td>3</td>
<td>4</td>
</tr>
<tr>
<td>Weighted Fitness</td>
<td>4.5</td>
<td>3</td>
<td>2.25</td>
</tr>
<tr>
<td>Normalized Fitness</td>
<td>0.46</td>
<td>0.31</td>
<td>0.23</td>
</tr>
<tr>
<td>Scaled Fitness</td>
<td>0.46</td>
<td>0.77</td>
<td>1</td>
</tr>
</tbody>
</table>

The first locus where crossover occurs is between the second and third bit in the third genotype chromosome of individual A. The third genotype chromosome crosses over with the first genotype chromosome.

- Genotype 1 0100
- Genotype 3 1011
- Low Crossover Mask = \((2 \^ \text{Crossover Locus}) - 1\)
- Low Crossover Mask = \((2 ^ 2) - 1\)
- Low Crossover Mask = 3
- High Crossover Mask = 15 XOR Low Crossover Mask
- High Crossover Mask = 1111 XOR 0011
- High Crossover Mask = 1100
- New Genotype 1 = \([(\text{Genotype 1 AND High Crossover Mask}) OR (\text{Genotype 3 AND Low Crossover Mask})]\)
- New Genotype 1 = \([(0100 AND 1100) OR (1011 AND 0011)]\)
- New Genotype 1 = \([(0100) OR (0011)]\)
- New Genotype 1 = 0111
- New Genotype 3 = \([(\text{Genotype 3 AND High Crossover Mask}) OR (\text{Genotype 1 AND Low Crossover Mask})]\)
- New Genotype 3 = \([(1011 AND 1100) OR (0100 AND 0011)]\)
- New Genotype 3 = \([(1000) OR (0000)]\)
- New Genotype 3 = 1000

After Crossover

<table>
<thead>
<tr>
<th>Individual</th>
<th>A</th>
<th>B</th>
<th>C</th>
</tr>
</thead>
<tbody>
<tr>
<td>Genotype 1</td>
<td>0111</td>
<td>1101</td>
<td>0110</td>
</tr>
<tr>
<td>Genotype 2</td>
<td>0011</td>
<td>0111</td>
<td>1101</td>
</tr>
<tr>
<td>Genotype 3</td>
<td>1000</td>
<td>1001</td>
<td>1111</td>
</tr>
<tr>
<td>Phenotype</td>
<td>0011</td>
<td>1101</td>
<td>1111</td>
</tr>
</tbody>
</table>
7. Mating Coupler and Create Offspring

After the fittest individuals are subject to the crossover operator, the genetic algorithm creates the next generation. A roulette wheel method is used to select three parents. A random number is chosen and compared to the scaled fitness probability. The parent selected is the first individual whose scaled fitness probability is greater than the random number. This roulette wheel selection method is repeated twice more to select three parents. Once all three parents are identified, the genetic algorithm creates three offspring. The creation of offspring is done by passing one genotype chromosome from each parent onto the offspring. The entire selection process is repeated, three parents chosen to create three offspring, until the number of offspring created equals the initial population. In the example, population size three, we will create three offspring by first randomly selecting the three parents.

<table>
<thead>
<tr>
<th>Individual</th>
<th>A</th>
<th>B</th>
<th>C</th>
</tr>
</thead>
<tbody>
<tr>
<td>Genotype 1</td>
<td>0111</td>
<td>1101</td>
<td>0110</td>
</tr>
<tr>
<td>Genotype 2</td>
<td>0011</td>
<td>0111</td>
<td>1101</td>
</tr>
<tr>
<td>Genotype 3</td>
<td>1000</td>
<td>1001</td>
<td>1111</td>
</tr>
<tr>
<td>Phenotype</td>
<td>0011</td>
<td>1101</td>
<td>1111</td>
</tr>
<tr>
<td>Fitness</td>
<td>2</td>
<td>3</td>
<td>4</td>
</tr>
<tr>
<td>Weighted Fitness</td>
<td>4.5</td>
<td>3</td>
<td>2.25</td>
</tr>
<tr>
<td>Normalized Fitness</td>
<td>0.46</td>
<td>0.31</td>
<td>0.23</td>
</tr>
<tr>
<td>Scaled Fitness</td>
<td>0.46</td>
<td>0.77</td>
<td>1</td>
</tr>
</tbody>
</table>

Random Number 1 = 0.26, 0 < 0.26 < 0.46, Parent 1 ~ Individual A
Random Number 2 = 0.92, 0.77 < 0.92 < 1, Parent 2 ~ Individual C
Random Number 3 = 0.01, 0 < 0.01 < 0.46, Parent 3 ~ Individual A

With the three parents identified, creating offspring is accomplished by transferring one genotype chromosome from each parent to each offspring.

<table>
<thead>
<tr>
<th>Parent</th>
<th>A</th>
<th>C</th>
<th>A</th>
</tr>
</thead>
<tbody>
<tr>
<td>Genotype 1</td>
<td>0111</td>
<td>0110</td>
<td>0111</td>
</tr>
<tr>
<td>Genotype 2</td>
<td>0011</td>
<td>1101</td>
<td>0011</td>
</tr>
</tbody>
</table>
8. Mutation Engine and Mutation

From the example one can see there is no mechanism in the genetic algorithm that requires all or any of the parents to be different. It is possible for all three parents to be the same individual. This does not mean that the offspring will be identical to the parent or each other; the mutation operation allows the offspring to vary from the parent. Again, because crossover occurs only within the genotype of an individual, it has no effect on the individual’s fitness, and it is unnecessary to re-evaluate the fitness of an individual. The only processes that require an individual’s fitness to be evaluated are creation and mutation. For this reason, mutation directly follows the creation of offspring. In order to make the genetic algorithm efficient, once an individual’s fitness is calculated, the individual is not permitted to mutate. Mutation is simply a bit flip.

The mutation operation is similar to the crossover operation in its implementation. A random number is compared to the mutation probability to determine if mutation occurs. If mutation occurs the program generates a mutation mask which is then bit-wise compared to the genotype chromosome to cause the mutation. The major difference is that crossover occurs at the loci between the bits whereas mutation affects the bits themselves. A reasonable starting value for the mutation probability is one over the chromosome length. With that in mind, the mutation probability for our example will be 0.25.

Before Mutation (bits to be mutated indicated with [ ])

<table>
<thead>
<tr>
<th>Offspring</th>
<th>D</th>
<th>E</th>
<th>F</th>
</tr>
</thead>
<tbody>
<tr>
<td>Genotype 1</td>
<td>0111</td>
<td>01[1]0</td>
<td>0111</td>
</tr>
<tr>
<td>Genotype 2</td>
<td>1101</td>
<td>0011</td>
<td>[0][0][1]1</td>
</tr>
<tr>
<td>Genotype 3</td>
<td>1[0][0]0</td>
<td>1000</td>
<td>1111</td>
</tr>
</tbody>
</table>
The first instance of mutation occurs in the third genotype chromosome of Offspring D at the third bit from the right. The following illustrates the bitwise comparison used to cause the mutation.

Mutation Mask = $2^{(\text{Mutation Bit} - 1)}$
Mutation Mask = $2^{(3 - 1)}$
Mutation Mask = $2^2$
Mutation Mask = 4
Mutation Mask = 0100
Genotype 3(after mutation) = Genotype 3(before mutation) XOR Mutation Mask
Genotype 3(after mutation) = 1000 XOR 0100
Genotype 3(after mutation) = 1100

After Mutation
Offspring D E F
Genotype 1 0111 0100 0111
Genotype 2 1101 0011 1001
Genotype 3 1100 1000 1111

9. Finishing the Example
The genetic algorithm is now at a point where it increments the generation counter by one and loops back to the phenotype calculator. To finish the simple example presented let us calculate the phenotype and fitness of the offspring.

<table>
<thead>
<tr>
<th>Individual</th>
<th>A</th>
<th>B</th>
<th>C</th>
</tr>
</thead>
<tbody>
<tr>
<td>Genotype 1</td>
<td>0111</td>
<td>1101</td>
<td>0110</td>
</tr>
<tr>
<td>Genotype 2</td>
<td>0011</td>
<td>0111</td>
<td>1101</td>
</tr>
<tr>
<td>Genotype 3</td>
<td>1000</td>
<td>1001</td>
<td>1111</td>
</tr>
<tr>
<td>Phenotype</td>
<td>0011</td>
<td>1101</td>
<td>1111</td>
</tr>
<tr>
<td>Fitness</td>
<td>2</td>
<td>3</td>
<td>4</td>
</tr>
<tr>
<td>Weighted Fitness</td>
<td>4.5</td>
<td>3</td>
<td>2.25</td>
</tr>
<tr>
<td>Normalized Fitness</td>
<td>0.46</td>
<td>0.31</td>
<td>0.23</td>
</tr>
<tr>
<td>Scaled Fitness</td>
<td>0.46</td>
<td>0.77</td>
<td>1</td>
</tr>
</tbody>
</table>

<table>
<thead>
<tr>
<th>Offspring</th>
<th>D</th>
<th>E</th>
<th>F</th>
</tr>
</thead>
<tbody>
<tr>
<td>Genotype 1</td>
<td>0111</td>
<td>0100</td>
<td>0111</td>
</tr>
<tr>
<td>Genotype 2</td>
<td>1101</td>
<td>0011</td>
<td>1001</td>
</tr>
<tr>
<td>Genotype 3</td>
<td>1100</td>
<td>1000</td>
<td>1111</td>
</tr>
<tr>
<td>Phenotype</td>
<td>1101</td>
<td>0000</td>
<td>1111</td>
</tr>
<tr>
<td>Fitness</td>
<td>3</td>
<td>0</td>
<td>4</td>
</tr>
</tbody>
</table>
In this simplified example, the genetic algorithm was able to find the optimum solution in just two generations. Typically, the optimum solution is unknown, and the genetic algorithm must be given some other termination criteria.

C. PARAMETERS

A genetic algorithm has certain parameters which must be set by the experimenter. In this application the parameters were: crossover probability, mutation probability, population size, and number of generations.

1. Population Size and Number of Generations

One problem that remains unresolved directly impacts the population size and the number of generations. The genetic algorithm establishes a socket TCP/IP connection with STK which it uses to send constellations to be evaluated and receives data in return. The problem is that the STK application overruns the virtual memory of the computer after evaluating about 30 generations. This causes the genetic algorithm to lock up. In order to generate the data for this thesis a workaround was developed. The genetic algorithm would have the option of either generating a random initial population or reading an initial population in from a data file. The algorithm ran for 20 generations and the fittest individuals at generation 20 were written into the data file. After 20 generations, the applications could be exited and the computer would empty the virtual memory page file. The genetic algorithm and STK could be re-launched and the data from the 20th generation read in as the 1st generation. To avoid running into the possibility of overrunning the virtual memory, the maximum generations was set to twenty and the population set at twelve.

The termination criterion for the genetic algorithm is number of generations. However, the need to restart the genetic algorithm after 20 generations enabled a different termination criterion to be used: if the fitness of the best constellation did not improve over 100 generations, the genetic algorithm was terminated.
2. Crossover and Mutation Probability

Crossover probability and mutation probability are typically varied over several runs of the genetic algorithm around an initial value that previous research in a similar field has indicated. Although previous research using genetic algorithms to optimize constellation design has been conducted, three differences between previous efforts and the current study exist. The existing research [21, 22, 23] has used haploid chromosomes instead of a phenotype derived from triploid genotype chromosomes. Previous research has used single chromosomes to represent constellations, whereas this study uses one derived phenotype to represent a single orbital element on a single satellite in the constellation. In addition, these studies have used crossover as the mechanism for creating offspring. Because each individual is represented by one binary chromosome, the offspring are created through uniform crossover. In uniform crossover the offspring has a 50% chance of inheriting each bit from one parent or the other. Also because these studies used single chromosomes to represent a constellation, the chromosome length varied dependent upon the number of satellites in the constellation. The mutation probabilities varied and were derived based on the length of the chromosome string and the size of the population. Finally, previous studies have used much larger populations than the current study.

To determine appropriate values for the crossover and mutation probabilities, this study employed a meta-genetic algorithm (meta-GA). A meta-GA is a genetic algorithm that optimizes another genetic algorithm. Along the way to the final iteration of the genetic algorithm used for constellation design, several previous versions were created. One of these previous versions was designed to try to achieve target values for the orbital elements. This version was used to debug the derivation of the phenotype from the triploid genotype and other issues without having to interface with STK. Over this test version, a simple genetic algorithm (two binary encoded, 8-bit, haploid chromosomes) was wrapped. The two chromosomes of this meta-GA determined the crossover and mutation probabilities of the test genetic algorithm. Each iteration of the test
A genetic algorithm consisted of 12 individuals over 100 generations. The meta-GA consisted of 32 individuals over 100 generations. The fitness function of the test genetic algorithm was the sum of the differences between the target value for each orbital element and the genetic algorithm result. The fitness function for the meta-GA was equal to the fitness of the best individual of the test genetic algorithm after 100 generations. Figure 9 illustrates the best 32 results from the meta-GA. The best crossover probabilities fall into two fairly distinct values: between 0.5 and 0.6, and approximately 0.1. The best mutation probabilities tend to fall between 0.2 and 0.3. Using this data the crossover probability for the genetic algorithm was set at approximately 0.51 (exactly 130/255) and the mutation probability was set at approximately 0.25 (exactly 63/255).

![Meta-GA Results](image)

**Figure 9. Meta-GA Data**

**C. STK INTERFACE**

The class module that implements the STK connection functionality, Visual Basic Connect Download, was downloaded directly from the STK website. It was used without modification. A documentation file is included in the download as well.
V. CONstellation Comparison

A. Background

To measure the effectiveness of the constellation generated using a genetic algorithm, the constellation designed using a genetic algorithm was tested against a constellation generated using parameters from the design of the Discoverer II Space Based Radar Constellation.

B. The Process

1. Constraints

Constraints were placed on the genetic algorithm constellation to allow a fair comparison against the Discoverer II constellation. Specifically, minimum altitude was limited to 500 km and the maximum altitude to 1,000 km. This was to force the genetic algorithm to produce a low earth orbit at an altitude similar to the 770 km altitude planned for Discoverer II.

Additionally, the genetic algorithm was constrained to produce a 24 satellite constellation – similar to the notional Discoverer II constellation. However, no constraints were placed on the number of satellites within a plane, or the spacing of the satellites. Every satellite in a constellation designed by the genetic algorithm had the same semi-major axis and inclination. The right ascension of the ascending node and the true anomaly were allowed to vary from satellite to satellite.

Finally, the genetic algorithm constellation was designed with the same sensor pattern as Discoverer II. The footprint created by the sensors will vary based on the altitude at which the satellites orbit. However, by keeping the sensor patterns equivalent, a constellation was created without factoring in differences in the physical satellites (a Discoverer II satellite could orbit in the genetic algorithm constellation).

2. Data Points

Fourteen data points were chosen to evaluate the fitness of the constellations. The data points were selected based on the global distribution of land and water by latitude as illustrated in Figure 10. The lower and middle
latitudes are dominated by water and the upper latitudes tend towards a more equitable distribution.

Figure 10. Distribution of Land and Water by Latitude [From 25]

A big advantage of SAR GMTI is the ability to track multiple targets in congested areas such as coastal harbors or high density vehicle traffic areas. Consequently, we focused our evaluation on land and coastal areas. In addition, the other products a space based radar system generates: SAR imagery and digital terrain elevation mapping, require the constellation to be optimized for coverage of land masses.

The data points were placed at 10° intervals between 55°S and 75°N latitude because the majority of the global land mass falls within those constraints. The only significant land mass below 55°S is Antarctica around which there is sparse traffic. Above 75°N lays only the Arctic Ocean and the northern parts of Canada and Greenland.

The longitudinal positions were determined based on an equitable distribution throughout the globe. With 14 latitudes 14 longitudes were needed. The earth was divided into 14 even segments in longitude. The speed of the genetic algorithm was highly dependent upon the speed of STK’s evaluation of
the coverage. For this reason coverage was evaluated for only 24 hours, and only at 14 data points. The 14 data points were selected from the 196 possible (14 latitudes by 14 longitudes) by applying the principles of the Eight Queens problem. The Eight Queens problem is a chess diversion that challenges a player to place eight queens on an eight by eight chessboard so that no two queens lie on the same row, column, or diagonal. The 14 points selected represent one possible solution to an extension of the Eight Queens problem.

The weights of the data points were based on the land area present between the latitudes in which each data point resided. The weight of each data point is listed in Table 2.

<table>
<thead>
<tr>
<th>Latitude</th>
<th>Longitude</th>
<th>Weight</th>
</tr>
</thead>
<tbody>
<tr>
<td>55 S</td>
<td>51.429</td>
<td>0.002</td>
</tr>
<tr>
<td>45 S</td>
<td>102.857</td>
<td>0.007</td>
</tr>
<tr>
<td>35 S</td>
<td>0.000</td>
<td>0.027</td>
</tr>
<tr>
<td>25 S</td>
<td>205.714</td>
<td>0.068</td>
</tr>
<tr>
<td>15 S</td>
<td>257.143</td>
<td>0.070</td>
</tr>
<tr>
<td>5 S</td>
<td>334.286</td>
<td>0.075</td>
</tr>
<tr>
<td>5 N</td>
<td>180.000</td>
<td>0.074</td>
</tr>
<tr>
<td>15 N</td>
<td>77.143</td>
<td>0.086</td>
</tr>
<tr>
<td>25 N</td>
<td>308.571</td>
<td>0.112</td>
</tr>
<tr>
<td>35 N</td>
<td>154.286</td>
<td>0.116</td>
</tr>
<tr>
<td>45 N</td>
<td>231.429</td>
<td>0.120</td>
</tr>
<tr>
<td>55 N</td>
<td>25.714</td>
<td>0.109</td>
</tr>
<tr>
<td>65 N</td>
<td>128.571</td>
<td>0.108</td>
</tr>
<tr>
<td>75 N</td>
<td>282.857</td>
<td>0.026</td>
</tr>
</tbody>
</table>

Table 2. Fitness Weighting by Latitude Land Area

3. Measurements of Fitness

An algorithm was generated for measuring fitness of the constellations using three figures of merit; average gap, maximum revisit and percent coverage. These three figures of merit were calculated for the 14 data points by STK for both the traditional Walker constellations and the genetic algorithm designed constellations. Toyon Research of Santa Barbara, California provided the data necessary to model the GMTI sensor in STK. These data are summarized in the table below.
<table>
<thead>
<tr>
<th>Sensor Mode</th>
<th>GMTI</th>
</tr>
</thead>
<tbody>
<tr>
<td>Minimum Grazing</td>
<td>Measured up from the tangential</td>
</tr>
<tr>
<td>Angle</td>
<td>6.0</td>
</tr>
<tr>
<td>Nadir Angle</td>
<td>Measured down from the horizontal</td>
</tr>
<tr>
<td></td>
<td>70.0</td>
</tr>
</tbody>
</table>

Table 3. GMTI Sensor Data

The values derived from the following three figures of merit contributed to the overall fitness. Each of the figures of merit contributed equally to the calculation of fitness in the genetic algorithm.

a. **Average Gap**
   The average gap is the total number of coverage gaps over the entire coverage interval divided by the time frame the measurement is taken, measured in seconds.

b. **Maximum Revisit**
   Revisit time is the amount of time a gap is present at a data point. Maximum revisit is the maximum duration of a gap in coverage over the entire coverage interval.

c. **Percent Coverage**
   Coverage time is the amount of time during which a data point is covered. Percent coverage is the total time (multiplied by 100) divided by the coverage interval, giving the percentage of time during which a point is covered.

d. **Weight**
   As listed in Table 2 above and illustrated in Figure 11, weights were assigned to each data point based on the percentage of land between the latitudes in which the data point is positioned.
e. Overall Fitness

Overall fitness was calculated using the three figures of merit together with the weighted values determined for each data point. The equation used for each data point is:

\[
\text{weight} \times \left\{ \left( \frac{\text{average gap}}{86400} \right) + \left( \frac{\text{maximum revisit}}{86400} \right) + \left[ 1 - \left( \frac{\text{percent coverage}}{100} \right) \right] \right\}
\]  

All data point sub totals were added together to generate the overall fitness number we used to compare the genetic algorithm constellations against the traditional Walker constellations.

C. RESULTS

The orbital elements for all of the constellations designed and evaluated are listed in Appendix B. The complete fitness calculations for all of the constellations evaluated are included in Appendix C.
1. 9-ball Constellations

The traditional 9-ball constellation designed had three orbital planes with three satellites per plane; the complete orbital parameters are listed in Appendix B. By applying Equation 5.1, the total fitness score for the 9-ball Walker constellation was 0.793. The genetic algorithm designed constellation orbital parameters are also listed in Appendix B. The total fitness score for the 9-ball genetic algorithm design was 0.673. Fitness score calculations for the 9-ball constellations are included in Appendix C. Figures 12-14 compare the average revisit, maximum gap, and percent coverage of the 9-ball Walker and the 9-ball GA design.

![Comparison of 9-ball Constellation Average Revisit Time](image.png)

Figure 12. Comparison of 9-ball Constellation Average Revisit Time

The 9-ball GA has a faster average revisit and lower maximum gap than the 9-ball Walker between the latitudes of 15° and 35° both north and south and at 75° north latitude.
Figure 13. Comparison of 9-ball Constellation Maximum Gap Time

Comparing percent coverage, the 9-ball GA design outperforms the 9-ball Walker at all latitudes.
2. **12-ball Constellations**

The traditional 12-ball constellation designed had four orbital planes with three satellites per plane; the complete orbital parameters are listed in Appendix B. By applying Equation 5.1, the total fitness score for the 12-ball Walker constellation was 0.650. The genetic algorithm designed 12-ball constellation orbital parameters are also listed in Appendix B. The total fitness score for the 12-ball genetic algorithm design was 0.576. Fitness score calculations for the 12-ball constellations are included in Appendix C. Figures 15-17 compare the average revisit, maximum gap, and percent coverage of the 12-ball Walker and the 12-ball GA design.
Figure 15. Comparison of 12-ball Constellation Average Revisit Time

The average revisit and maximum gap of the 12-ball GA design outperformed the 12-ball Walker design between 15° north and south latitudes and at the extreme northern latitude.
Figure 16. Comparison of 12-ball Constellation Maximum Gap Time

In comparing percent coverage, the 12-ball GA design provides more coverage of the data points than the 12-ball Walker design at all latitudes.
Figure 17. Comparison of 12-ball Constellation Percent Coverage

3. **18-ball Constellations**

The traditional 18-ball constellation designed had six orbital planes with three satellites per plane; the complete orbital parameters are listed in Appendix B. By applying Equation 5.1, the total fitness score for the 18-ball Walker constellation was 0.456. The genetic algorithm designed 18-ball constellation orbital parameters are also listed in Appendix B. The total fitness score for the 18-ball genetic algorithm design was 0.386. Fitness score calculations for the 18-ball constellations are included in Appendix C. Figures 18-20 compare the average revisit, maximum gap, and percent coverage of the 18-ball Walker and the 18-ball GA design.
The average revisit of the 12-ball GA design outperformed the 12-ball Walker design only at 5° north, 65° north, and 75° north latitude. The maximum gap time of the 18-ball GA design outperformed the 18-ball Walker constellation at 75° north latitude only.
Although the 18-ball GA design did not perform as well as the 18-ball Walker design in average revisit time or maximum gap time at most latitudes, it did provide greater coverage at all latitudes.
Figure 20. Comparison of 18-ball Constellation Percent Coverage

4. 24-ball Constellations

The traditional design method used the Discoverer II space based radar as its base model. The Discoverer II constellation would have been a 24-ball constellation with 3 satellites in each of 8 orbital planes. In addition to modeling the Discoverer II constellation, the traditional design method also evaluated other permutations of a 24-ball Walker satellite. Walker constellations of 4 orbital planes, 6 orbital planes, and 12 orbital planes were also modeled. Orbital elements for all 24-ball Walker constellations modeled are included in Appendix B. Using Equation 5.1 the following fitness scores were calculated for the 24-ball Walker constellations: 4 orbital planes, 0.321; 6 orbital planes, 0.221; 8 orbital planes, 0.275; 12 orbital planes, 0.320. The calculations of fitness scores for all 24-ball Walker constellations are included in Appendix C. Two 24-ball constellations were developed by the genetic algorithm. For the dependent case constellation, the right ascension and anomaly of each satellite was referenced to the position of the previous satellite, with the exception of the first satellite which was referenced from the first point of Aries and to the ascending node. In the
independent case, the right ascension and anomaly of all the satellites were referenced from the first point of Aries and the ascending node. The orbital elements for both of the GA designed 24-ball constellations are listed in Appendix B. The 24-ball, dependent case GA designed constellation and the 24-ball, independent case GA constellation had fitness scores of 0.276. Complete fitness score calculations can be found in Appendix C. In Figures 21-23 below, the performance of the 24-ball independent case GA design; 24-ball, 6 orbital plane Walker; and the 24-ball, 8 orbital plane Walker are compared.

![Comparison of 24-ball Constellation Average Revisit Time](image)

**Figure 21. Comparison of 24-ball Constellation Average Revisit Time**

The 24-ball GA designed constellation outperforms the two Walker constellations only at 75° north latitude. The GA design outperforms the 8-plane Walker at 5° north and south latitude as well.
The 24-ball GA design was unable to outperform either of the Walker constellations in maximum gap time at any latitude. The 24-ball GA design was able to provide better coverage than the 8 plane Walker between the latitudes of 35° south and 35° north. It provided better coverage than both of the Walker constellations at 75° north latitude. In addition to 75° north, the GA design did provide a higher coverage percentage than the 6 plane Walker at 15° south, 5° south, and 15° north latitudes. At all other latitudes, the Walker constellations were able to provide better coverage.
Figure 23. Comparison of 24-ball Constellation Percent Coverage

D. SUMMARY

As Table 4 indicates, the constellations of 9, 12 and 18 satellites created using a genetic algorithm did perform better than the constellations of 9, 12 and 18 satellites using traditional constellation design methods. With 9, 12, or 18 satellites per constellation, the best performance in terms of average revisit and maximum gap was a function of the latitude, with the genetic algorithm performing better near the equator and at the northernmost latitude and the Walker constellations performing better at the mid-latitudes. However, in 9-ball, 12-ball, and 18-ball constellations, the genetic algorithm outperformed the Walker constellation in percent coverage at every latitude.

However, the 24 satellite genetic algorithm constellations were unable to outperform all of the traditional 24 satellite Walker constellations. The 24-ball genetic algorithm constellations were able to achieve better fitness scores than the 24-ball Walker constellations of 4 planes and 12 planes. The GA constellations performed nearly as well as the 8 plane Walker constellation which
modeled the proposed Discoverer II space based radar system. Clearly the 24-ball GA constellations were dominated by the performance of the 24-ball, 6 plane Walker constellation.

<table>
<thead>
<tr>
<th>Constellation</th>
<th>Fitness</th>
</tr>
</thead>
<tbody>
<tr>
<td>9-ball</td>
<td></td>
</tr>
<tr>
<td>GA Design</td>
<td>0.673</td>
</tr>
<tr>
<td>3-plane Walker</td>
<td>0.793</td>
</tr>
<tr>
<td>12-ball</td>
<td></td>
</tr>
<tr>
<td>GA Design</td>
<td>0.576</td>
</tr>
<tr>
<td>4-plane Walker</td>
<td>0.650</td>
</tr>
<tr>
<td>18-ball</td>
<td></td>
</tr>
<tr>
<td>GA Design</td>
<td>0.386</td>
</tr>
<tr>
<td>6-plane Walker</td>
<td>0.456</td>
</tr>
<tr>
<td>24-ball</td>
<td></td>
</tr>
<tr>
<td>GA Design, Dependent Case</td>
<td>0.276</td>
</tr>
<tr>
<td>GA Design, Independent Case</td>
<td>0.276</td>
</tr>
<tr>
<td>4-plane Walker</td>
<td>0.321</td>
</tr>
<tr>
<td>6-plane Walker</td>
<td>0.221</td>
</tr>
<tr>
<td>8-plane Walker, Discoverer II</td>
<td>0.275</td>
</tr>
<tr>
<td>12-plane Walker</td>
<td>0.320</td>
</tr>
</tbody>
</table>

Table 4. Summary of Fitness Scores for Designed Constellations

There are two possible explanations why the genetic algorithm was not able to design a constellation that could outperform the best 24-ball Walker constellation. The first possibility is that the termination criteria selected for the genetic algorithm, 100 generations with no improvement in the fitness score of the fittest individual, did not permit the genetic algorithm to thoroughly search the complex search space of a 24 satellite constellation. The second explanation is that as the number of satellites increases and the constellation becomes less sparse, a Walker constellation provides nearly optimal performance. In either case, the answer was beyond the scope of this thesis. Further study modifying
the termination criteria will determine if the first hypothesis is true. If the first hypothesis is false, further study to investigate the number of satellites where a genetic algorithm design can no longer outperform a Walker constellation is indicated.
VI. THE BENEFITS OF SUPERIORITY

A. THE THREAT

The uncertainty among security officials is not if al-Qaeda will attack again, but when. Equally important questions are where and how. One theory on how the next attack will occur is via the sea. Specifically, many security officials believe al-Qaeda will acquire a merchant ship and use it as a floating bomb or blow up a merchant ship similar to the attack on the USS Cole.

An article entitled *Merchant Shipping 'At Risk of Attack'* by Michael Evans, was recently published in the *London Times*. In his article, Mr. Evans writes of Admiral Sir Alan West, the head of the Royal Navy, stating al-Qaeda is plotting to blow up merchant ships. According to Mr. Evans, Admiral West stated, "What we have noticed is that al-Qaeda and other organizations have an awareness about maritime trade. They've realized how important it is for world trade in general (and) they understand that significance." Admiral West revealed, "We have seen plans from intelligence of (proposed) attacks on merchant shipping. We are aware that they have plans and they have looked at this."

Attacks on the French supertanker, *Limburg*, and the USS COLE prove al-Qaeda is willing and able to attack ships. However, most susceptible militaries have increased security on and around their vessels leaving merchant vessels as the most likely target.

As Admiral West points out, it is very difficult to attack a ship on the open sea. Therefore, the most likely target will be in ports and maritime choke points. According to Michael Evans, "A suspected al-Qaeda cell was arrested in Morocco two years ago after a plot was exposed to attack shipping in the Strait of Gibraltar." As a result, following the thwarted attack, western merchant ships were assigned escorts through the Strait.

 Escorting merchant ships in and out of port and through straits is certainly effective, but not efficient. The escort ships are being pulled from their primary
mission and the number of merchant ship targets outnumbers the number of available escorts. An alternate solution should be explored. [26] 

B. A POSSIBLE SOLUTION

1. The Theory

A number of assumptions are implicit regarding the concept of operations of a space based radar system in the equations derived below. First, the satellite will be given the initial position of a track of interest. Second, the constellation will have a common track correlation database and tasking system so that information about a track generated by one satellite is available to all satellites. Third, the satellite system will be able to identify a specific track with SAR imagery. Finally, the SAR will be able to image all candidate tracks designated by the GMTI sensor and no time delay exists in switching between SAR and GMTI modes.

With these assumptions in mind, the number of tracks a space based radar system can maintain can be derived from either the maximum gap time or the average revisit time of the constellation. If the space based radar system operates autonomously, receiving no data from other sources to help maintain a track, the maximum gap is used in calculating the number of tracks maintained. If the space based radar has the ability to receive cues or supplemental data from other sources to assist in maintaining a track, average revisit time should be used in calculating the number of tracks maintained.

The number of tracks maintained is equal to the time an area is in view divided by the time it takes to locate a specific track.

\[
N = \frac{T_v}{T_l}
\]

The time required to locate a specific track is equal to the time required to search for a track added to the time to identify a track.
Time required to search for a track

Time required to identify a track

\[ Tl = Ts + Ti \]

The time required to search for a specific track is equal to the search area divided by the search rate.

\[ Sa = \text{Search area} \]
\[ Sr = \text{Search rate} \]
\[ Ts = \frac{Sa}{Sr} \]

The time required to identify a specific track is equal to the number of candidate tracks multiplied by the time to image one track. On average, the system will have to image half of the candidate tracks to locate a specific track.

\[ Nc = \text{Number of candidate tracks} \]
\[ Tsar = \text{Time to image a track} \]
\[ Ti = \frac{1}{2} * Nc * Tsar \]

The number of candidate targets equals the area of uncertainty multiplied by the target density.

\[ Au = \text{Area of uncertainty} \]
\[ Td = \text{Target density} \]
\[ Nc = Au * Td \]

The search area equals the area of uncertainty which is the area of the circle with radius equal to the revisit time multiplied by the target velocity.

\[ Tr = \text{Revisit time} \]
\[ Vt = \text{Target velocity} \]
\[ Sa = Au = \pi * (Tr * Vt)^2 \]

Substituting and simplifying.
\[
N = \frac{TV}{Tl}
\]
\[
N = \frac{TV}{Ts + Ti}
\]
\[
N = \frac{TV}{\left(\frac{Sa}{Sr}\right) + \left(\frac{1}{2} \cdot Nc \cdot Tsar\right)}
\]
\[
N = \frac{TV}{\left(\frac{\pi \cdot (Tr \cdot Vt)^2}{Sr}\right) + \left(\frac{1}{2} \cdot Au \cdot Td \cdot Tsar\right)}
\]
\[
N = \frac{TV}{\left(\frac{\pi \cdot (Tr \cdot Vt)^2}{Sr}\right) + \left(\frac{1}{2} \cdot \pi \cdot (Tr \cdot Vt)^2 \cdot Td \cdot Tsar\right)}
\]
\[
N = \frac{TV}{\pi \cdot (Tr \cdot Vt)^2 \cdot \frac{1}{Sr} + \frac{Td \cdot Tsar}{2}}
\]
\[
N = \frac{TV}{\pi \cdot (Tr \cdot Vt)^2 \cdot \frac{2 + Sr \cdot Td \cdot Tsar}{2 \cdot Sr}}
\]
\[
N = \frac{2 \cdot TV \cdot Sr}{\pi \cdot (Tr \cdot Vt)^2 \cdot (2 + Sr \cdot Td \cdot Tsar)}
\]

The average time in view for a point is approximately 0.63 times the maximum time in view for a point, based on a circular footprint. In addition, the search area (area of uncertainty) may or may not be completely within by the footprint. The space based radar system cannot reliably maintain track on a target if the area of uncertainty is not completely covered by sensor footprint. The calculation of tracks maintained discounts a track if any portion of the area of uncertainty falls outside the sensor footprint. This is accounted for by taking the inverse of the ratio of the radius of the search area to the radius of the footprint.

\[Rf = \text{Radius of the access footprint}\]
\[
N = \left(1 - \frac{Tr \cdot Vt}{Rf}\right) \cdot \frac{2 \cdot TV \cdot Sr}{\pi \cdot (Tr \cdot Vt)^2 \cdot (2 + Sr \cdot Td \cdot Tsar)}
\]
\[
N = \left(1 - \frac{Tr \cdot Vt}{Rf}\right) \cdot \frac{4 \cdot TV \cdot Sr}{\pi^2 \cdot Tr^2 \cdot Vt^2 \cdot (2 + Sr \cdot Td \cdot Tsar)}
\]
2. The Math

The equations derived above can next be applied with the figure of merit values calculated for the constellations designed. The example below is the calculation of the number of tracks maintained by the Discoverer II constellation at 35 N° based on average revisit time, the latitude of San Francisco, CA and Washington, D.C.

To calculate the radius of the sensor footprint, it is necessary to know the elevation or grazing angle and the altitude of the satellite. The figure below illustrates the geometry of the earth, satellite, and target which is used in deriving the footprint radius. The radius of the footprint is the distance from the subsatellite point to the target which equals the earth central angle times the radius of the earth.

Figure 24. Geometry between the Earth, Satellite, and Target [After 5]
$R_e = 6378.1363 km$

\[ \sin \rho = \frac{R_e}{R_e + H} \]

\[ \varepsilon = 6^\circ = \frac{\pi}{30} \]

\[ \sin \eta = \cos \varepsilon \sin \rho \]

\[ \eta = \sin^{-1} (\cos \varepsilon \sin \rho) \]

\[ \lambda + \eta + \varepsilon = 90^\circ = \frac{\pi}{2} \]

\[ \lambda = \frac{\pi}{2} - (\eta + \varepsilon) \]

\[ \lambda = \frac{\pi}{2} - \left( \sin^{-1} (\cos \varepsilon \sin \rho) + \frac{\pi}{30} \right) \]

\[ \lambda = \frac{\pi}{2} - \left[ \sin^{-1} \left( \cos \frac{\pi}{30} \sin \rho \right) + \frac{\pi}{30} \right] \]

\[ \lambda = \frac{\pi}{2} - \left[ \sin^{-1} \left( \cos \frac{\pi}{30} \cdot \frac{R_e}{R_e + H} \right) + \frac{\pi}{30} \right] \]

\[ Rf = R_e \cdot \lambda \]

\[ Rf = 6378.1363 km \left[ \frac{\pi}{2} - \left( \sin^{-1} \left( \cos \frac{\pi}{30} \left( \frac{6378.1363 km}{(6378.1363 + 770) km} \right) \right) + \frac{\pi}{30} \right) \right] \]

\[ Rf = 2388.105 km \]

\[ T_v = 2 \cdot \left( \frac{2388.105 km}{7.467 km/s} \right) = 639.642 s \]

\[ T_d = \frac{1}{km^2} \]

\[ Tsar = 20 s \]

In this example, the average revisit time will be used. This will result in an answer that would be applicable in the case where the space based radar received or integrated data from other sources to assist it in maintaining a target track.

\[ Tr = 49.22 s \]
The GMTI sensor search rate is dependent on altitude and target velocity. It was necessary to develop an equation for search rate that would enable the comparison of constellations at different altitudes. Toyon Research provided the sensor data from which the search rate equations were derived. The data were provided for a GMTI sensor at altitudes of 1000 km and 10000km and for target velocities of 4 kph, 8kph, and 16kph. The equations derived were linear extrapolations of the data provided. The data and the equations are summarized in the table below; search rate is in km²/s.

<table>
<thead>
<tr>
<th>Target Velocity</th>
<th>Altitude</th>
<th>4 kph</th>
<th>8 kph</th>
<th>16 kph</th>
</tr>
</thead>
<tbody>
<tr>
<td>1000 km</td>
<td>105</td>
<td>258</td>
<td>1074</td>
<td></td>
</tr>
<tr>
<td>10000 km</td>
<td>527</td>
<td>1044</td>
<td>1233</td>
<td></td>
</tr>
<tr>
<td>Derived Equation</td>
<td>y = 0.0469x + 58.111</td>
<td>y = 0.0873x + 170.67</td>
<td>y = 0.0177x + 1056.3</td>
<td></td>
</tr>
</tbody>
</table>

Table 5. Search Rate Data and Equations

In this example, the number of tracks maintained will be calculated for a target velocity of 4 kph.

\[
V_t = 4 \frac{km}{h} = 0.00111 \frac{km}{s}
\]

\[
Sr = 0.0469 (altitude) + 58.111 = 0.0469 (770) + 58.111 = 94.224 \frac{km^2}{s}
\]

Now that all of the known variables have been identified and defined, the number of 4 kph tracks maintained by the Discoverer II constellation at 35 N can be calculated.
\[ Sa = Au = \pi \left( 49.22 \times 0.00111 \right)^2 = 0.00940 km^2 \]

\[ Nc = 0.00940 km^2 \times \frac{1}{km^2} = 0.00940 \]

\[ Ti = \frac{1}{2} \times 0.00940 \times 20s = 0.0940s \]

\[ Ts = \frac{0.00940 km^2}{94.224 \frac{km^2}{s}} = 9.972 \times 10^{-5} s \]

\[ TI = 0.0940 s + 9.972 \times 10^{-5} s = 0.0941s \]

\[ N = \left( \frac{639.642s}{0.0941s} \right) \left( 1 - \frac{49.22s \times 0.00111 \frac{km}{s}}{2388.105 km} \right) = 6799.898 \approx 6800 \]

3. The Solution

By repeating the calculation above for the fittest constellations designed, the number of tracks reliably maintained by each constellation can be calculated. The number of tracks maintained varies by latitude, the target’s velocity, and track density. Figures 13, 14, 15, and 16 illustrate the number of tracks maintained by latitude and target velocity for the 24-ball Walker constellation, the 18-ball GA constellation, the 12-ball GA constellation, and the 9-ball GA constellation, respectively. These results were calculated using average revisit time and a track density of 1/km$^2$. Calculating the number of tracks maintained using average revisit time models the performance of a space based radar system that could integrate data or receive track information from other sources. In all of the graphs, a significant decrease in performance is evident at the lower latitudes.
Figure 25. 24-ball Walker Performance

18 ball GA Constellation Performance Derived from Average Revisit Time

Figure 26. 18-ball GA Performance
12 ball GA Constellation Performance Derived from Average Revisit Time

Figure 27. 12-ball GA Performance

9 ball GA Constellation Performance Derived from Average Revisit Time

Figure 28. 9-ball GA Performance
In the calculation of tracks maintained, only the average revisit time figure of merit was used. This may raise the question why not modify the fitness function so that the genetic algorithm would either only seek to minimize the average revisit time or at least give the average revisit time more weight in the calculation. That would be an option if the concept of operations of the space based radar system specified that it would never operate autonomously and that it only would be used for GMTI. However, the space based radar system will have other functions that will not depend as heavily on average revisit time.

C. CONCLUSION

If fielded, Discoverer II Space Based Radar Constellation would have been able to track 6800 targets traveling at 4kph at 35°N. Space based radar constellations comprised of less than 24 satellites suffer a drop in performance. However, if an SBR constellation is constrained to less than 24 satellites, genetic algorithms offer a fitness improvement over traditional constellation design methods. The critical step in implementing a genetic algorithm to design a constellation is defining the fitness function in a manner that will translate into performance gains.

D. FURTHER STUDY

1. Caveat

These satellite constellation comparisons are based on revisit and coverage performance only. Other considerations such as satellite dimensions, sensor complexity, design timeline, acquisition resources or launch vehicles were not included. Additionally, constellations were designed with the same sensor capability in varying orbital patterns.

The consequence of not considering the parameters listed above is that predictions cannot be made concerning the feasibility of the constellations designed. For instance, a constellation designed by the genetic algorithm may have performed better than traditional constellations. However, the cost of launching satellites in many different orbital planes and into retrograde orbits may be prohibitive.
2. Next Steps

Follow on research should be conducted using the genetic algorithm from this thesis as a constellation design tool. Additional runs of the genetic algorithm should be conducted using a coverage definition consisting of a different set of data points to validate the results. Rather than choosing either to have a satellite’s position referenced to the previous satellite, the dependent case, or referenced to a fixed point, the independent case, a combination of the two methods could be implemented. This combination may result in the construction of a parallel genetic algorithm that would build on the current genetic algorithm. This could be accomplished by having the current genetic algorithm generate satellites referenced to each other, the dependent case. Following the evaluation of fitness, the fittest members of the population would go through an additional step where the each satellite would be permitted to mutate independently and this mutation evaluated. Additional work should be done in deriving constellation performance (tracks maintained) from the figure of merit values. This may lead to developing a better fitness function in which the figures of merit are not weighed evenly. It may be desirable to have the number of tracks maintained by a space based radar system remain nearly constant across all latitudes. Further research should be conducted in the weights derived for the latitudes and the derivation of the fitness function to reward the genetic algorithm for designing constellations with similar revisit and gap statistics across all latitudes. The genetic algorithm has demonstrated the ability to design sparse constellations better than traditional Walker designs. Further study could be conducted to determine if the genetic algorithm can design constellations achieving the same level of performance as a Walker constellation with fewer satellites.
LIST OF REFERENCES


APPENDIX A GENETIC ALGORITHM CODE

'COMPGA
'Constellation Optimization via Multi-Ploid Genetic Algorithm
'Version 3.0
'Programmed by Douglas J. Pegher
'July 2004

Option Explicit

dimension module constants and variables
'Integer value of last generation
Const mLastGeneration As Integer = 20
'Population size; must be divisible by 3
Const mTotalConstellations As Integer = 9
'Number of satellites in constellation
Const mSatelliteNumber As Integer = 24
'Number of orbital element variables ~ no eccentricity, no arg. of perigee ~ alt. and inc. same for all sats
Const mOrbitalElement As Integer = 4
'Probability of crossover
Const mCrossoverProbability As Single = 0.6
'Probability of mutation
Const mMutationProbability As Single = 0.0625
'define semimajor axis range
Const MinSemiMajorAxis As Single = 500 'min value
Const MaxSemiMajorAxis As Single = 1000 'max value
'define inclination range
Const MinInclination As Single = 0 'min value
Const MaxInclination As Single = 180 'max value
'define rightascension range
Const MinRightAscension As Single = 0 'min value
Const MaxRightAscension As Single = 360 'max value
Dim mPreviousRightAscension As Single
'define anomaly range
Const MinAnomaly As Single = 0 'min value
Const MaxAnomaly As Single = 360 'max value
Dim mPreviousAnomaly As Single
'current generation counter
Dim mCurrentGeneration As Integer
'data array
Dim mGAArray(3, (mOrbitalElement - 1), (mSatelliteNumber - 1), (mTotalConstellations - 1), (mLastGeneration - 1)) As Byte
'working array
Dim mFitnessArray(5, mTotalConstellations - 1, (mLastGeneration - 1)) As Single
Dim Weights(14) As Single
Dim AvgGap(14) As Single
Dim MaxRevisit(14) As Single
Dim PercentCoverage(14) As Single
'STK connection
Dim mConnection As StkCon
Private Sub cmdConnect_Click()
If cmdConnect.Caption = "Connect to STK" Then
    Set mConnection = New StkCon
    Call mConnection.connect("localhost:5001")
    'Call mConnection.sendCommand("ConControl /AsyncOn")
    optConnected.Value = mConnection.connected
    cmdConnect.Caption = "Disconnect from STK"
Else
    mConnection.disconnect
    optConnected.Value = mConnection.connected
    cmdConnect.Caption = "Connect to STK"
End If
    cmdRun.Enabled = mConnection.connected
End Sub

Private Sub cmdRun_Click()
' main sequence in compga
' disable connection button
cmdConnect.Enabled = False
' initialize random number generator
Randomize
' loops through for each generation
For mCurrentGeneration = 0 To (mLastGeneration - 1)
    If mCurrentGeneration = 0 And chkReadData.Value = Unchecked Then
        ' the original random population is generated in this call
        Call CreateInitialPopulation
        Call Phenotype
        Call EvaluateFitness ' evaluate fitness of calculated phenotype
        Call ScaleFitness ' recalculate weighted, normalized and scaled fitnesses
        Call CrossoverEngine ' crossover engine
    End If
Next
End Sub
ElseIf mCurrentGeneration = 0 And chkReadData.Value = Checked Then
    Call ReadData
    Call Phenotype
    Call EvaluateFitness 'evaluate fitness of calculated phenotype
    Call ScaleFitness 'recalculate weighted, normalized and scaled fitnesses

Else
    Call MatingCoupler 'mating engine
    Call MutationEngine 'mutation engine
    Call Phenotype
    Call CopyFitnessArray 'copies previous fitness scores into current generation
    Call EvaluateFitness 'evaluate fitness of calculated phenotype
    Call ScaleFitness 'recalculate weighted, normalized and scaled fitnesses
    Call CrossoverEngine 'crossover engine
End If
Next mCurrentGeneration

cmdDisplay.Enabled = mConnection.connected
cmdConnect.Enabled = True
'when GOMPGA is finished running write all data to file.
Call WriteData
Call WriteFitness

End Sub
Private Sub CreateInitialPopulation()
    'called from Private Sub btnRun_Click
    'create initial population
    'dimension subprocedure variables
    Dim Constellation As Integer
    Dim Satellite As Integer
    Dim OrbitalElement As Integer
    Dim Genotype As Integer

    'loop through GAArray assigning random numbers to the data cells for the first generation
    'random numbers range from 32767 to -32768
    'random numbers of 0.9999847412109375 and greater yield 32767
'random numbers of less than 0.0000152587890625 yield -32768
'the short data type represents the values of -32768 to 32767 as 16 bit binary
'as binary strings the values range from 0 up to 32767 and then -32768 to -1
'-1 is represented as 16 ones
For Constellation = 0 To (mTotalConstellations - 1)
    For Satellite = 0 To (mSatelliteNumber - 1)
        For OrbitalElement = 0 To (mOrbitalElement - 1)
            For Genotype = 0 To 2
                mGAArray(Genotype, OrbitalElement, Satellite, Constellation, mCurrentGeneration) = CByte(255 * Rnd())
            Next Genotype
        Next OrbitalElement
    Next Satellite
Next Constellation
End Sub
Private Sub Phenotype()
    'called from Private Sub CreateInitialPopulation
    'called from Private Sub Mutation
    'calculate phenotypes for most recently created generation

    'dimension subprocedure variables
    Dim Constellation As Integer
    Dim Satellite As Integer
    Dim OrbitalElement As Integer

    'loop through current generation and calculate phenotypes
    'phenotype is the result of the bitwise comparison of the three genotype values
    'the bit is decided by a simple majority of the bits in the genotype
    For Constellation = 0 To (mTotalConstellations - 1)
        For Satellite = 0 To (mSatelliteNumber - 1)
            For OrbitalElement = 0 To (mOrbitalElement - 1)
mGAArray(3, OrbitalElement, Satellite, Constellation, mCurrentGeneration) =
((mGAArray(1, OrbitalElement, Satellite, Constellation, mCurrentGeneration) And mGAArray(2, OrbitalElement, Satellite, Constellation, mCurrentGeneration)) Or
(mGAArray(2, OrbitalElement, Satellite, Constellation, mCurrentGeneration) And mGAArray(3, OrbitalElement, Satellite, Constellation, mCurrentGeneration)) Or
(mGAArray(3, OrbitalElement, Satellite, Constellation, mCurrentGeneration) And mGAArray(1, OrbitalElement, Satellite, Constellation, mCurrentGeneration)))
Next OrbitalElement
Next Satellite
Next Constellation

Private Sub EvaluateFitness()
'called from Private Sub Phenotype
'evaluate fitness of current generation

'dimension variables
Dim Constellation As Integer
Dim Satellite As Integer
Dim Fitness As Single
Dim NewSatelliteString As String
Dim NewSensorString As String
Dim OrbitalElementString As String
Dim SetStateString As String
Dim SatellitePath As String
Dim SensorPath As String
Dim CoveragePath As String
Dim AvgGapReportPath As String
Dim MaxRevisitReportPath As String
Dim PercentCoverageReportPath As String
Dim Counter As Integer
Dim AvgGapFitness As Single
Dim MaxRevisitFitness As Single
Dim PercentcoverageFitness As Single

Weights(0) = 0 'unused
Weights(14) = 0.071379704 '75deg N
Weights(13) = 0.180843786 '65deg N
Weights(12) = 0.134321551 '55deg N
Weights(11) = 0.11972634 '45deg N
Weights(10) = 0.099885975 '35deg N
Weights(9) = 0.087571266 '25deg N
Weights(8) = 0.062941847 '15deg N

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Weights(7) = 0.052451539    '5deg N
Weights(6) = 0.05359179    '5deg S
Weights(5) = 0.051083238    '15deg S
Weights(4) = 0.05313569    '25deg S
Weights(3) = 0.023717218    '35deg S
Weights(2) = 0.007069555    '45deg S
Weights(1) = 0.002280502    '55deg S

CoveragePath = "*/CoverageDefinition/55s_75n_10deg"
AvgGapReportPath = CoveragePath & "/FigureOfMerit/Avg_Gap ""Avg Gap"
MaxRevisitReportPath = CoveragePath & "/FigureOfMerit/Max_Revisit ""Max Revisit"
PercentCoverageReportPath = CoveragePath & "/FigureOfMerit/Percent_Coverage ""Percent Coverage"
'convert the phenotype short into a positive integer ~ conversion in functions
'send positive integer to be scaled ~ scale range defined in functions
'determine fitness
'a lower fitness is better
'fitness is evaluated by generation, but the fitness array is updated for each individual constellation
For Constellation = 0 To (mTotalConstellations - 1)
  'create new constellation
  For Satellite = 0 To (mSatelliteNumber - 1)
    'create new satellite
    NewSatelliteString = "New / */Satellite 
    & Satellite
    mConnection.sendCommand
    (NewSatelliteString)
    'input satellite ephemeris
    OrbitalElementString = SemiMajorAxis(mGAArray(3, 0, 0, Constellation, mCurrentGeneration)) & " 0.0 " & Inclination(mGAArray(3, 1, 0, Constellation, mCurrentGeneration)) & " 0.0 " & RightAscension(mGAArray(3, 2, Satellite, Constellation, mCurrentGeneration), Satellite) & " " & Anomaly(mGAArray(3, 3, Satellite, Constellation, mCurrentGeneration), Satellite)
    SatellitePath = "*/Satellite/" & Satellite
    'SetState Scenario/stkDemo/Satellite/ERS1
    Classical J2Perturbation "1 Nov 2000 00:00:00.00" "1 Nov
SetStateString = "SetState " & SatellitePath & " Classical J2Perturbation " & OrbitalElementString
mConnection.sendCommand (SetStateString)

NewSensorString = "New / " & SatellitePath & "/Sensor SMTI"
mConnection.sendCommand (NewSensorString)

SensorPath = SatellitePath & "/Sensor/SMTI"

SensorPath & " Conical 20.0 " & OuterHalfAngle(SemiMajorAxis(mGAArray(3, 0, 0, Constellation, mCurrentGeneration))) & " 0.0 360.0")
mConnection.sendCommand ("Define " & SensorPath & " Cov " & CoveragePath & " Asset " & SensorPath & " Assign")

mConnection.sendCommand ("Cov " & CoveragePath & " Access Compute")
mConnection.sendCommand ("GetReport " & AvgGapReportPath)

mConnection.returnedLine(Counter) = AvgGap(Counter) =
mConnection.sendCommand ("GetReport " & MaxRevisitReportPath)

mConnection.returnedLine(Counter) = MaxRevisit(Counter) =
mConnection.sendCommand ("GetReport " & PercentCoverageReportPath)

mConnection.returnedLine(Counter) = PercentCoverage(Counter) =

'm calculate fitness
For Counter = 1 To 14
    PercentCoverage(Counter) =
    mConnection.returnedLine(Counter) =

Next Counter
For Counter = 1 To 14
    PercentCoverage(Counter) =

Next Counter
For Counter = 1 To 14
    PercentCoverage(Counter) =

Next Counter
For Counter = 1 To 14
    PercentCoverage(Counter) =

Next Counter
For Counter = 1 To 14
    PercentCoverage(Counter) =

Next Counter
Fitness = Fitness + (Weights(Counter) * ((AvgGap(Counter) / 86400) + (MaxRevisit(Counter) / 86400) + (1 - (PercentCoverage(Counter) / 100))))

Next Counter
  'update fitness array
  Call FitnessArray(Constellation, Fitness)
  Fitness = 0
  'clear accesses
  mConnection.sendCommand ("Cov " & CoveragePath & " Access Clear")
  'unload constellation
  mConnection.sendCommand ("UnloadMulti / */Satellite/*")

Next Constellation

End Sub

Private Sub FitnessArray(ByVal Constellation As Integer, ByVal Fitness As Single)
  'called from Private Sub EvaluateFitness
  'compare a constellation to current fitest constellations and replace the less fit
  'dimension subprocedure variables
  Dim FitnessColumn As Integer

  'loop through fitness array and compare candidate constellation fitness to previous constellation fitness
  'fitness in the test function is arranged from low to high, low fitnesses are better
  If mCurrentGeneration = 0 Then
    For FitnessColumn = 0 To Constellation
      If FitnessColumn < Constellation And Fitness < mFitnessArray(2, FitnessColumn, mCurrentGeneration) Then
        Call ChangeOrder(FitnessColumn)
        mFitnessArray(0, FitnessColumn, mCurrentGeneration) = Constellation
        mFitnessArray(1, FitnessColumn, mCurrentGeneration) = mCurrentGeneration
        mFitnessArray(2, FitnessColumn, mCurrentGeneration) = Fitness
        Exit For
      ElseIf FitnessColumn = Constellation
      Then
        mFitnessArray(0, FitnessColumn, mCurrentGeneration) = Constellation
  End If

Then
  mFitnessArray(0, FitnessColumn, mCurrentGeneration) = Constellation
mFitnessArray(1, FitnessColumn) = mCurrentGeneration
mFitnessArray(2, FitnessColumn) = Fitness

End If
Next FitnessColumn

ElseIf mCurrentGeneration <> 0 Then
For FitnessColumn = 0 To (mTotalConstellations - 1)
If Fitness < mFitnessArray(2, FitnessColumn, mCurrentGeneration) Then
Call ChangeOrder(FitnessColumn)
End If
mFitnessArray(0, FitnessColumn, mCurrentGeneration) = Constellation
mFitnessArray(1, FitnessColumn, mCurrentGeneration) = mFitnessArray(0, FitnessColumn, mCurrentGeneration)
mFitnessArray(2, FitnessColumn, mCurrentGeneration) = Fitness
Exit For
End If
Next FitnessColumn
End If

End Sub
Private Sub ChangeOrder(ByVal FitnessColumn As Integer)
'called from Private Sub FitnessArray
'update fitness array
'dimension subprocedure variables
Dim Column As Integer

'loop through fitness array in reverse order and shift constellation order
'this copies the next best fitness into the current cell
For Column = (mTotalConstellations - 1) To (FitnessColumn + 1) Step -1
mFitnessArray(0, Column, mCurrentGeneration) = mFitnessArray(0, Column - 1, mCurrentGeneration)
mFitnessArray(1, Column, mCurrentGeneration) = mFitnessArray(1, Column - 1, mCurrentGeneration)
mFitnessArray(2, Column, mCurrentGeneration) = mFitnessArray(2, Column - 1, mCurrentGeneration)
Private Sub ScaleFitness()
    'called from Private Sub FitnessArray
    'calculate normalized fitness probabilities
    for use in mating
    'dimension subprocedure variables
    Dim FitnessColumn As Integer
    Dim FitnessSum As Single
    Dim WeightedSum As Single

    'loop through FitnessArray to sum fitnesses
    For FitnessColumn = 0 To (mTotalConstellations - 1)
        FitnessSum = FitnessSum +
        (mFitnessArray(2, FitnessColumn, mCurrentGeneration))
        Next FitnessColumn

    'loop through FitnessArray to calculate weighted fitnesses.
    For FitnessColumn = 0 To (mTotalConstellations - 1)
        mFitnessArray(3, FitnessColumn, mCurrentGeneration) = (FitnessSum / (mFitnessArray(2, FitnessColumn, mCurrentGeneration)))
        Next FitnessColumn

    'loop through FitnessArray to sum weighted fitnesses
    For FitnessColumn = 0 To (mTotalConstellations - 1)
        WeightedSum = WeightedSum +
        (mFitnessArray(3, FitnessColumn, mCurrentGeneration))
        Next FitnessColumn

    'loop through FitnessArray to calculate normalized fitnesses.
    For FitnessColumn = 0 To (mTotalConstellations - 1)
        mFitnessArray(4, FitnessColumn, mCurrentGeneration) = (mFitnessArray(3, FitnessColumn, mCurrentGeneration) / WeightedSum)
        Next FitnessColumn
'scaled fitness is running total of individual fitness percentage of total fitness
    For FitnessColumn = 0 To (mTotalConstellations - 1)
        If FitnessColumn = 0 Then
            mFitnessArray(5, FitnessColumn, mCurrentGeneration) = mFitnessArray(4, FitnessColumn, mCurrentGeneration)
        ElseIf FitnessColumn <> 0 And FitnessColumn <> (mTotalConstellations - 1) Then
            mFitnessArray(5, FitnessColumn, mCurrentGeneration) = m_fitness_array(5, FitnessColumn - 1, mCurrentGeneration) + mFitnessArray(4, FitnessColumn, mCurrentGeneration)
        ElseIf FitnessColumn <> 0 And FitnessColumn = (mTotalConstellations - 1) Then
            mFitnessArray(5, FitnessColumn, mCurrentGeneration) = 1
        End If
    Next FitnessColumn
End Sub
Private Sub CrossoverEngine()
    'called from Private Sub btnRun_Click
    'crossover can occur in any of the constellations in the fitness array
    'crossover involves the exchange of bits between genotype strings
    'crossover does not affect fitness in the organism; it may affect the fitness in the offspring
    'determine if crossover occurs
    'dimension subprocedure variables
    Dim FitnessColumn As Integer
    Dim Satellite As Integer
    Dim OrbitalElement As Integer
    Dim Genotype As Integer
    Dim Bit As Integer
    Dim GenotypeA As Integer
    Dim GenotypeB As Integer

    'loop through possible crossover sites to determine where crossover occurs
    For FitnessColumn = 0 To (mTotalConstellations - 1)
        For Satellite = 0 To (mSatelliteNumber - 1)
            ...
For OrbitalElement = 0 To (mOrbitalElement - 1)
    For Genotype = 0 To 2
        For Bit = 1 To 7
            If Rnd() < mCrossoverProbability Then
                If Genotype = 0 Then
                    GenotypeA = 0
                    GenotypeB = 1
                ElseIf Genotype = 1 Then
                    GenotypeA = 1
                    GenotypeB = 2
                ElseIf Genotype = 2 Then
                    GenotypeA = 2
                    GenotypeB = 0
                End If
                Call Crossover(GenotypeA, GenotypeB, OrbitalElement, Satellite, CInt(mFitnessArray(0, FitnessColumn, mCurrentGeneration)), CInt(mFitnessArray(1, FitnessColumn, mCurrentGeneration)), Bit)
            End If
        Next Bit
    Next Genotype
Next OrbitalElement
Next Satellite
Next FitnessColumn
End Sub

Private Sub Crossover(ByVal GenotypeA As Integer, ByVal GenotypeB As Integer, ByVal OrbitalElement As Integer, ByVal Satellite As Integer, ByVal Constellation As Integer, ByVal Generation As Integer, ByVal CrossoverBit As Integer)
    'called from Private Sub CrossoverEngine
    'perform crossover of given values at given point
    'crossover bit values between 1 and 15
    'crossover takes place after the designated bit
    'there are 15 possible crossover loci
    'dimension subprocedure variables
    Dim NewGAArrayValueA As Byte
    Dim NewGAArrayValueB As Byte
Dim HiCrossoverMask As Byte
Dim LoCrossoverMask As Byte
Dim HiArrayValueA As Byte
Dim LoArrayValueA As Byte
Dim HiArrayValueB As Byte
Dim LoArrayValueB As Byte

' generate crossover masks
LoCrossoverMask = CByte((2 ^ CrossoverBit) - 1)
HiCrossoverMask = 255 Xor LoCrossoverMask

' generate partial crossovers
HiArrayValueA = mGAArray(GenotypeA, OrbitalElement, Satellite, Constellation, Generation) And HiCrossoverMask
LoArrayValueA = mGAArray(GenotypeA, OrbitalElement, Satellite, Constellation, Generation) And LoCrossoverMask
HiArrayValueB = mGAArray(GenotypeB, OrbitalElement, Satellite, Constellation, Generation) And HiCrossoverMask
LoArrayValueB = mGAArray(GenotypeB, OrbitalElement, Satellite, Constellation, Generation) And LoCrossoverMask

' recombine partial crossovers
NewGAArrayValueA = HiArrayValueA Or LoArrayValueB
NewGAArrayValueB = HiArrayValueB Or LoArrayValueA

' update GAArray
mGAArray(GenotypeA, OrbitalElement, Satellite, Constellation, Generation) = NewGAArrayValueA
mGAArray(GenotypeB, OrbitalElement, Satellite, Constellation, Generation) = NewGAArrayValueB

End Sub

Private Sub MatingCoupler()
' called from Private Sub btnRun_Click
' create mating triples
' dimension subprocedure variables
Dim Coupling As Integer
Dim Probability As Single
Dim FitnessColumn As Byte
Dim ParentA As Integer
Dim ParentB As Integer
Dim ParentC As Integer

'roulette wheel selection of parents
'requires the fitnesses to be scaled
For Coupling = 0 To CInt((mTotalConstellations / 3) - 1)
    Probability = Rnd()
    For FitnessColumn = 0 To (mTotalConstellations - 1)
        If Probability < mFitnessArray(5, FitnessColumn, mCurrentGeneration - 1) Then
            ParentA = FitnessColumn
        End If
    Next
    Probability = Rnd()
    For FitnessColumn = 0 To (mTotalConstellations - 1)
        If Probability < mFitnessArray(5, FitnessColumn, mCurrentGeneration - 1) Then
            ParentB = FitnessColumn
        End If
    Next
    Probability = Rnd()
    For FitnessColumn = 0 To (mTotalConstellations - 1)
        If Probability < mFitnessArray(5, FitnessColumn, mCurrentGeneration - 1) Then
            ParentC = FitnessColumn
        End If
    Next
    Call CreateOffspring(ParentA, ParentB, ParentC, Coupling)
Next Coupling

End Sub

Private Sub CreateOffspring(ByVal ParentA As Integer, ByVal ParentB As Integer, ByVal ParentC As Integer, ByVal Coupling As Integer)
    'called from Private Sub MatingCoupler
    'create next generation

    'dimension subprocedure variables
    Dim Satellite As Integer
    Dim OrbitalElement As Integer

    'create offspring 1

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For Satellite = 0 To (mSatelliteNumber - 1)
  For OrbitalElement = 0 To (mOrbitalElement - 1)
    mGAArray(0, OrbitalElement, Satellite, ((3 * (Coupling + 1)) - 3), (mCurrentGeneration)) =
    mGAArray(0, OrbitalElement, Satellite, CInt(mFitnessArray(0, ParentA, mCurrentGeneration - 1)),
               CInt(mFitnessArray(1, ParentA, mCurrentGeneration - 1)))
    mGAArray(1, OrbitalElement, Satellite, ((3 * (Coupling + 1)) - 3), (mCurrentGeneration)) =
    mGAArray(0, OrbitalElement, Satellite, CInt(mFitnessArray(0, ParentB, mCurrentGeneration - 1)),
               CInt(mFitnessArray(1, ParentB, mCurrentGeneration - 1)))
    mGAArray(2, OrbitalElement, Satellite, ((3 * (Coupling + 1)) - 3), (mCurrentGeneration)) =
    mGAArray(0, OrbitalElement, Satellite, CInt(mFitnessArray(0, ParentC, mCurrentGeneration - 1)),
               CInt(mFitnessArray(1, ParentC, mCurrentGeneration - 1)))
  Next OrbitalElement
Next Satellite

'create offspring 2
For Satellite = 0 To (mSatelliteNumber - 1)
  For OrbitalElement = 0 To (mOrbitalElement - 1)
    mGAArray(0, OrbitalElement, Satellite, ((3 * (Coupling + 1)) - 2), (mCurrentGeneration)) =
    mGAArray(0, OrbitalElement, Satellite, CInt(mFitnessArray(0, ParentB, mCurrentGeneration - 1)),
               CInt(mFitnessArray(1, ParentB, mCurrentGeneration - 1)))
    mGAArray(1, OrbitalElement, Satellite, ((3 * (Coupling + 1)) - 2), (mCurrentGeneration)) =
    mGAArray(0, OrbitalElement, Satellite, CInt(mFitnessArray(0, ParentC, mCurrentGeneration - 1)),
               CInt(mFitnessArray(1, ParentC, mCurrentGeneration - 1)))
    mGAArray(2, OrbitalElement, Satellite, ((3 * (Coupling + 1)) - 2), (mCurrentGeneration)) =
    mGAArray(0, OrbitalElement, Satellite, CInt(mFitnessArray(0, ParentA, mCurrentGeneration - 1)),
               CInt(mFitnessArray(1, ParentA, mCurrentGeneration - 1)))
  Next OrbitalElement
Next Satellite

'create offspring 3
For Satellite = 0 To (mSatelliteNumber - 1)
  For OrbitalElement = 0 To (mOrbitalElement - 1)
mGAArray(0, OrbitalElement, Satellite, ((3 * (Coupling + 1)) - 1), (mCurrentGeneration)) = mGAArray(0, OrbitalElement, Satellite, CInt(mFitnessArray(0, ParentC, mCurrentGeneration - 1)), CInt(mFitnessArray(1, ParentC, mCurrentGeneration - 1)))
mGAArray(1, OrbitalElement, Satellite, ((3 * (Coupling + 1)) - 1), (mCurrentGeneration)) = mGAArray(0, OrbitalElement, Satellite, CInt(mFitnessArray(0, ParentA, mCurrentGeneration - 1)), CInt(mFitnessArray(1, ParentA, mCurrentGeneration - 1)))
mGAArray(2, OrbitalElement, Satellite, ((3 * (Coupling + 1)) - 1), (mCurrentGeneration)) = mGAArray(0, OrbitalElement, Satellite, CInt(mFitnessArray(0, ParentB, mCurrentGeneration - 1)), CInt(mFitnessArray(1, ParentB, mCurrentGeneration - 1)))
Next OrbitalElement
Next Satellite

End Sub
Private Sub MutationEngine()
'called from Private Sub btnRun_Click
'determine if mutation occurs
'dimension subprocedure variables
Dim Constellation As Integer
Dim Satellite As Integer
Dim OrbitalElement As Integer
Dim Genotype As Integer
Dim Bit As Integer

'loop through possible mutation locations to
determine where mutation occurs
For Constellation = 0 To (mTotalConstellations - 1)
    For Satellite = 0 To (mSatelliteNumber - 1)
        For OrbitalElement = 0 To (mOrbitalElement - 1)
            For Genotype = 0 To 2
                For Bit = 1 To 8
                    If Rnd() < mMutationProbability Then
                        Call Mutation(Genotype, OrbitalElement, Satellite, Constellation, mCurrentGeneration, Bit)
                    End If
                Next Bit
            Next Genotype
        Next OrbitalElement
    Next Satellite
Next Constellation

End If
Next Bit
Next Genotype

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Private Sub Mutation(ByVal Genotype As Integer, ByVal OrbitalElement As Integer, ByVal Satellite As Integer, ByVal Constellation As Integer, ByVal Generation As Integer, ByVal MutationBit As Integer)
'called from Private Sub MutationEngine
'mutate given bit
'mutation is a bit flip
'dimension variables
Dim MutationMask As Byte

'create mutation mask
MutationMask = CInt(2 ^ (MutationBit - 1))

'update GAArray
mGAArray(Genotype, OrbitalElement, Satellite, Constellation, Generation) = mGAArray(Genotype, OrbitalElement, Satellite, Constellation, Generation) Xor MutationMask

End Sub

Private Sub CopyFitnessArray()
'copy previous generation fitness scores into current array for comparison.
Dim FitnessColumn As Integer
Dim FitnessRow As Integer

For FitnessColumn = 0 To (mTotalConstellations - 1)
    For FitnessRow = 0 To 5
        mFitnessArray(FitnessRow, FitnessColumn, mCurrentGeneration) = mFitnessArray(FitnessRow, FitnessColumn, mCurrentGeneration - 1)
    Next FitnessRow
Next FitnessColumn
End Sub

Private Function SemiMajorAxis(ByVal ArrayValue As Byte) As Single
'return a scaled altitude value to the fitness function
SemiMajorAxis = 1000 * (6378.1363 + (MinSemiMajorAxis + ((MaxSemiMajorAxis - MinSemiMajorAxis) * (ArrayValue / 255))))

End Function
Private Function Inclination(ByVal ArrayValue As Byte) As Single
    'return a scaled inclination to the fitness function
    Inclination = MinInclination + ((MaxInclination - MinInclination) * (ArrayValue / 255))
End Function
Private Function RightAscension(ByVal ArrayValue As Byte, ByVal Satellite As Integer) As Single
    'return a scaled right ascension to the fitness function
    If Satellite = 0 Then
        mPreviousRightAscension = 0
        RightAscension = MinRightAscension + ((MaxRightAscension - MinRightAscension) * (ArrayValue / 255))
    ElseIf mPreviousRightAscension + (MinRightAscension + ((MaxRightAscension - MinRightAscension) * (ArrayValue / 255))) > 360 Then
        RightAscension = (mPreviousRightAscension + ((MaxRightAscension - MinRightAscension) * (ArrayValue / 255))) - 360
    Else
        RightAscension = mPreviousRightAscension + (MinRightAscension + ((MaxRightAscension - MinRightAscension) * (ArrayValue / 255)))
    End If
    mPreviousRightAscension = RightAscension
End Function
Private Function Anomaly(ByVal ArrayValue As Byte, ByVal Satellite As Integer) As Single
    'return a scaled anomaly to the fitness function
    If Satellite = 0 Then
        mPreviousAnomaly = 0
        Anomaly = MinAnomaly + ((MaxAnomaly - MinAnomaly) * (ArrayValue / 255))
    ElseIf mPreviousAnomaly + (MinAnomaly + ((MaxAnomaly - MinAnomaly) * (ArrayValue / 255))) > 360 Then
        Anomaly = (mPreviousAnomaly + ((MaxAnomaly - MinAnomaly) * (ArrayValue / 255))) - 360
    Else
        Anomaly = mPreviousAnomaly + (MinAnomaly + ((MaxAnomaly - MinAnomaly) * (ArrayValue / 255)))
    End If
    mPreviousAnomaly = Anomaly
Anomaly = (mPreviousAnomaly + (MinAnomaly + ((MaxAnomaly - MinAnomaly) * (ArrayValue / 255)))) - 360
Else
Anomaly = mPreviousAnomaly + (MinAnomaly + ((MaxAnomaly - MinAnomaly) * (ArrayValue / 255)))
End If
mPreviousAnomaly = Anomaly

End Function

Private Sub cmdDisplay_Click()
' dimension variables
Dim Rank As Integer
Dim Generation As Integer
Dim GenerationCreated As Integer
Dim Constellation As Integer
Dim Satellite As Integer
Dim NewSatelliteString As String
Dim OrbitalElementString As String
Dim SetStateString As String
Dim SatellitePath As String
Dim NewSensorString As String
Dim SensorPath As String
Dim CoveragePath As String
CoveragePath = "*/CoverageDefinition/55s_75n_10deg"

'display the data values in the GAArray
If cmdDisplay.Caption = "Display" Then
Rank = CInt(txtRank.Text) - 1
Generation = CInt(txtGeneration.Text) - 1
GenerationCreated = CInt(mFitnessArray(1, Rank, Generation))
Constellation = CInt(mFitnessArray(0, Rank, Generation))
txtGenerationCreated.Text = GenerationCreated
txtConstellation.Text = Constellation
txtFitness.Text = mFitnessArray(2, Rank, Generation)
txtWeight.Text = mFitnessArray(3, Rank, Generation)
txtNormal.Text = mFitnessArray(4, Rank, Generation)
txtScale.Text = mFitnessArray(5, Rank, Generation)
'recreate constellation in STK
For Satellite = 0 To (mSatelliteNumber - 1)
    'create new satellite
    'assign satellite to constellation
    NewSatelliteString = "New / */Satellite " & Satellite
    mConnection.sendCommand (NewSatelliteString)

    NewSatelliteString = "New / */Satellite " & Satellite
    mConnection.sendCommand (NewSatelliteString)

    'input satellite ephemeris
    OrbitalElementString = SemiMajorAxis(mGAArray(3, 0, 0, Constellation, GenerationCreated)) & " 0.0 " & Inclination(mGAArray(3, 1, 0, Constellation, GenerationCreated)) & " 0.0 " & RightAscension(mGAArray(3, 2, Satellite, Constellation, GenerationCreated), Satellite) & " " & Anomaly(mGAArray(3, 3, Satellite, Constellation, GenerationCreated), Satellite)
    SatellitePath = "*/Satellite/" & Satellite

    SetStateString = "SetState " & SatellitePath & " Classical J2Perturbation ""1 Jun 2003 12:00:00.00"" ""2 Jun 2003 12:00:00.00"" 60 J2000 ""1 Jun 2003 12:00:00.00"" " & OrbitalElementString
    mConnection.sendCommand (SetStateString)

    NewSensorString = "New / */Sensor SMTI"
    mConnection.sendCommand (NewSensorString)

    SensorPath = SatellitePath & "/Sensor/SMTI"

    'input sensor parameters
    mConnection.sendCommand ("Define " & SensorPath & " Conical 20.0 " & OuterHalfAngle(SemiMajorAxis(mGAArray(3, 0, 0, Constellation, GenerationCreated))) & " 0.0 360.0")

    'assign sensor to coverage
    mConnection.sendCommand ("Cov " & CoveragePath & " Asset " & SensorPath & " Assign")
    Next Satellite

    'change caption
    cmdDisplay.Caption = "Unload"
Else
    mConnection.sendCommand ("UnloadMulti /*Satellite/*")
txtRank.Text = ""

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txtGeneration.Text = ""
cmdDisplay.Caption = "Display"
End If
End Sub
Private Function OuterHalfAngle(ByVal SemiMajorAxis As Single) As Single
'calculate outer half angle for given semi major axis
Const pi As Single = 3.14159265358979
Dim x As Single
Dim Radians As Single
x = Cos(pi / 30) * (6378.1363 / (SemiMajorAxis / 1000))
Radians = Atn(x / Sqr(-x * x + 1))
OuterHalfAngle = Radians * (180 / pi)
End Function
Private Sub ReadData()
Dim Constellation As Integer
Dim Satellite As Integer
Dim OrbitalElement As Integer
Dim Genotype As Integer
Dim Data As Integer
Dim Position As Integer
Position = 1
Open "C:\Documents and Settings\Douglas J. Pegher\My Documents\Thesis\VB6\COMPGAvbyte\data\gabytedata.txt" For Random Access Read As #1
' Read the contents of the file.
For Constellation = 0 To (mTotalConstellations - 1)
    For Satellite = 0 To (mSatelliteNumber - 1)
        For OrbitalElement = 0 To (mOrbitalElement - 1)
            For Genotype = 0 To 2
                Get #1, Position, Data
                mGAArray(Genotype, OrbitalElement, Satellite, Constellation, 0) = Data
                Position = Position + 1
            Next Genotype
        Next OrbitalElement
    Next Satellite
Next Constellation
Close #1
End Sub
Private Sub WriteData()
Dim Constellation As Integer
Dim Satellite As Integer
Dim OrbitalElement As Integer
Dim Genotype As Integer
Dim Data As Integer
Dim Position As Integer
Position = 1

Open "C:\Documents and Settings\Douglas J. Pegher\My Documents\Thesis\VB6\COMPGAvbyte\data\gabytedata.txt" For Random Access Write As #1

For Constellation = 0 To (mTotalConstellations - 1)
    For Satellite = 0 To (mSatelliteNumber - 1)
        For OrbitalElement = 0 To (mOrbitalElement - 1)
            For Genotype = 0 To 2
                Data = (mGAArray(Genotype, OrbitalElement, Satellite, mFitnessArray(0, Constellation, (mLastGeneration - 1)), mFitnessArray(1, Constellation, (mLastGeneration - 1))))
                Put #1, Position, Data
                Position = Position + 1
            Next Genotype
        Next OrbitalElement
    Next Satellite
Next Constellation
Close #1

End Sub
Private Sub WriteFitness()

Dim fso, txtfile
Set fso = CreateObject("Scripting.FileSystemObject")
Set txtfile = fso.OpenTextFile("C:\Documents and Settings\Douglas J. Pegher\My Documents\Thesis\VB6\COMPGAvbyte\data\fitnessbytedata.txt", 8)
    ' Write a line with a newline character.
    txtfile.WriteLine (mFitnessArray(2, 0, 19))
    txtfile.Close

End Sub
APPENDIX B  CONSTITUTION ORBITAL ELEMENTS

A.  9-BALL CONSTELLATIONS

1. Genetic Algorithm Design

<table>
<thead>
<tr>
<th>Satellite</th>
<th>Semi-major axis</th>
<th>Eccentricity</th>
<th>Inclination</th>
<th>RAAN</th>
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2. 3-plane Walker

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B.  12-BALL CONSTELLATIONS

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### C. 18-BALL CONSTELLATIONS

#### 1. Genetic Algorithm Design

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### 3. 24-BALL CONSTELLATIONS

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## APPENDIX C  CONSTELLATION FITNESS CALCULATIONS

### A. 9-BALL CONSTELLATIONS

#### 1. Genetic Algorithm Design

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Overall Fitness: 0.673

#### 2. 3-plane Walker

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Overall Fitness: 0.793
### B. 12-BALL CONSTELLATIONS

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**Overall Fitness**: 0.576

#### 2. 4-plane Walker

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**Overall Fitness**: 0.650
C. 18-BALL CONSTELLATIONS

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Overall Fitness: 0.386

2. 6-plane Walker

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Overall Fitness: 0.456
### D. 24-BALL CONSTELLATIONS

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Overall Fitness 0.276

2. **Genetic Algorithm Design, Dependent Case**

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<th>SubTotal</th>
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Overall Fitness 0.276
3. **4-plane Walker**

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Overall Fitness 0.321

4. **6-plane Walker**

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Overall Fitness 0.221
5. **8-plane Walker, Discoverer II**

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**Overall Fitness 0.275**

6. **12-plane Walker**

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**Overall Fitness 0.320**
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