

Overview

Thursday, November 3, 2022 1:09 PM

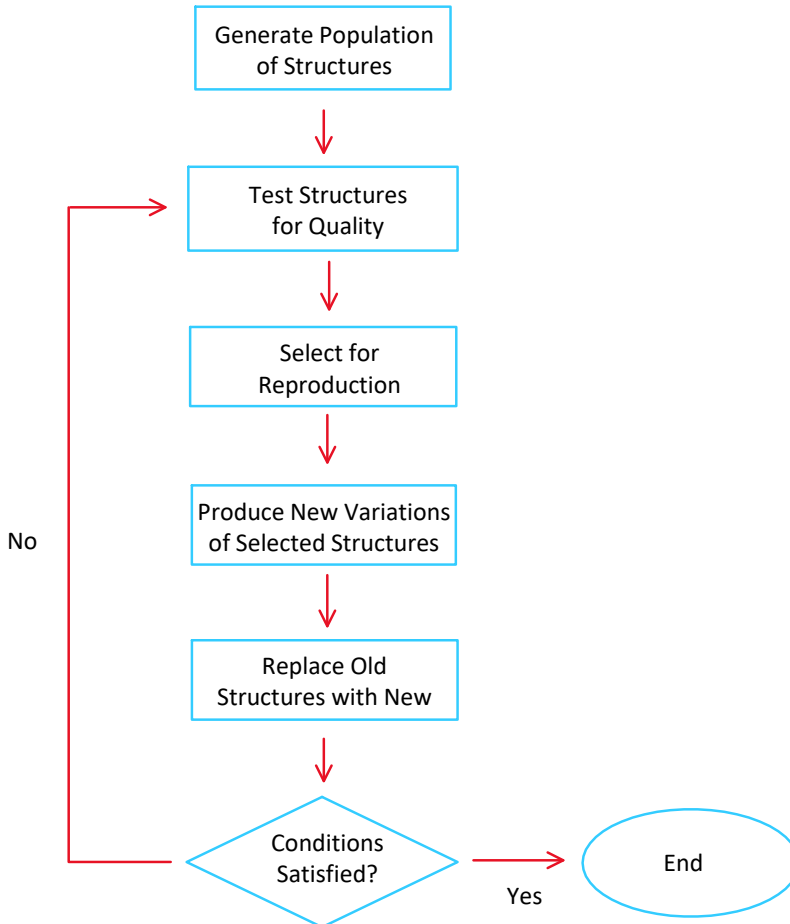
Advantages:

- Ignores discontinuities
- Since it uses random methods, it has good exploration vs exploitation
- Intuitive methods based on theories in biology
- Object-oriented creatures
- Embarrassingly parallel

Disadvantages:

- Computationally intensive

Founder: John Holland 1970's



Optimization: the search for the “best” solution for a problem

Fitness function: evaluates how close a given solution is to the optimum solution of the problem

Decision variables: independent variables to be manipulated when searching for the optimum solution

Constraint: restriction placed on decision variables or solution

Feasible solution: solution that does not violate any constraints

Search space: all values that the decision variables can assume

Mutation: (unary variation) changing a single structure

Crossover: (binary variation) exchanging material between structures

Penalty function: reduces the value of the fitness function when a constraint is violated

Niching: reducing fitness of population members in proportion to the number of other solutions that are the same. This makes solutions less good once they have been discovered by several other members of the population. This maintains population diversity, allowing to reach a global optima

Fitness landscape: graph of the fitness function of the entire space of data structures that are members of a population

Gene: a single encoded parameter

Chromosome: A set of genes making up an individual in a population

Simple Genetic Algorithm (SGA)

Steady State Genetic Algorithm (SSGA): No generations. Only the best fit individuals are added back into the population.

Selection Methods

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Bad selection can result in premature convergence at a local optima

n-tournament: the Population is shuffled randomly and divided into small groups. The most fit individuals in each small group are chosen to be parents.

Advantages:

- The most fit genes are guaranteed to survive
 - Efficient and simple
1. Choose some number of individuals randomly from a population (with or without replacement)
 2. Select the best individual from this group
 3. Repeat as often as desired (usually until the mating pool is filled)

Roulette wheel/Proportionate: chooses parents in proportion to their fitness.

Probability of selection = gene fitness / sum of population fitness

$$p(i) = \frac{f(i)}{\sum f}$$

- Slow

Roulette Wheel Parent Selection

1. Sum the fitnesses of all the population members; call the result total fitness.
2. Generate n , a random number between 0 and total fitness.
3. Return the first population member whose fitness, added to the fitnesses of the preceding population members, is greater than or equal to n .

Examples of Roulette Wheel Parent Selection

Chromosome	1	2	3	4	5	6	7	8	9	10
Fitness	8	2	17	7	2	12	11	7	3	7
Running Total	8	10	27	34	36	48	59	66	69	76

Random Number	23	49	76	13	1	27	57
Chromosome Chosen	3	7	10	3	1	3	7

Figure 1.5: Examples of roulette wheel parent selection. The first table shows the fitness of ten chromosomes and the running total of fitness. The second table shows the chromosome that would be chosen by the roulette wheel algorithm using these fitness values for each of seven randomly generated numbers.

From "Handbook of Genetic Algorithms" by Lawrence Davis 1991

Rank selection: ordered by fitness and selected by rank (1-least fit, n -most fit)

Probability of selection = gene rank/sum of population ranks

1. Rank population members
2. Generate random number between 0 and population size
3. Return the member whose rank equals the random number

Linear Ranking and binary tournaments have been shown to have identical performance

Absolute fitness replacement: replacing the least fit members of the population

Locally elite replacement: The two parents and the two children are examined in the two most fit are put into the population in the slots occupied by the parents

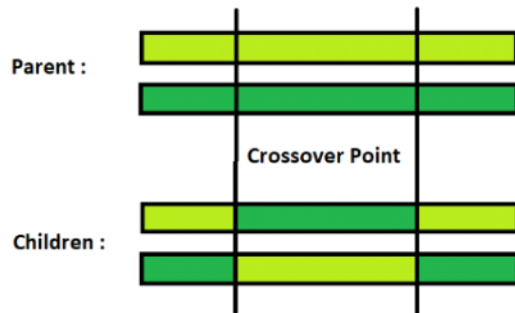
Goldberg 1991:

<https://www.cse.unr.edu/~sushil/class/gas/papers/Select.pdf>

Crossover Types

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n-point: n random points are chosen along a chromosome string and genetic material is exchanged between two parents



Uniform: Each gene is selected randomly from the genes of the parents (idea: have a probability based on fitness instead of random). Uses a binary mask, which is like a coin flip. Masks are selected every time crossover happens. Each mask value is random (value of .5, which makes it uniform)

Adaptive Methods:

"This can be achieved in a brute force manner by repeated calls to crossover until non-clones are produced, or in a more sophisticated fashion, such as Booker's reduced surrogate approach." - Dejong K, Spears WM. An analysis of the interacting roles of population size and crossover in genetic function optimization. 1990.

n-point having the point location as encoded material

Genetic material can determine what method to use! Pretty cool

Also:

Random elite: each child is compared with a random member of the population and is replaced only if it is least as fit

p_c is typically between .5 and 1.0

Mutation and Population Size

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Mutation rate can theoretically reduce population size by increasing the number of new cases, but this risks the possibility of losing a good solution

Mutation typically occurs during reproduction

Real number creep: If a function is likely continuous, then it is good to jump around the current solution in small, random increments

Too small of a population size can introduce [genetic drift](#)

50 seems to be a decent start (number of bit lengths?)

p_m (probability of mutation) is typically between .001 and .005

D. Thierens, "Adaptive mutation rate control schemes in genetic algorithms," Proceedings of the 2002 Congress on Evolutionary Computation. CEC'02 (Cat. No.02TH8600), 2002, pp. 980-985 vol.1, doi: 10.1109/CEC.2002.1007058.

Multi-Objective Optimization

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Subscripts

i	gene or decision variable index
j	chromosome index

Superscripts

n	GA generation or population index
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Each chromosome has a fitness vector $\mathbf{F}_j^n(x_1, x_2, \dots, x_n)$

Given:

If a lower x_1 and x_2 is better, then

$\mathbf{F}_1^n(x_1, x_2) = (2, 10) \rightarrow$ non-dominated

$\mathbf{F}_2^n(x_1, x_2) = (4, 6) \rightarrow$ non-dominated

$\mathbf{F}_3^n(x_1, x_2) = (8, 4) \rightarrow$ non-dominated

$\mathbf{F}_4^n(x_1, x_2) = (9, 5) \rightarrow$ dominated

$\mathbf{F}_5^n(x_1, x_2) = (7, 8) \rightarrow$ dominated

Goldberg Ranking:

1. Identify all non-dominated individuals in population
2. Assign them a rank of 1
3. Temporally remove from population
4. Repeat previous steps on the remaining set of non-dominated individuals until all have been assigned a rank (next round gets assigned a rank of 2)

This scheme assures equal reproductive potential.

Goldberg, D. E.: Genetic Algorithms in Search, Optimization and Machine Learning. AddisonWesley, Reading, MA, 1989,

Bin/Block Ranking:

1. Use greedy selection until the number of points reaches a certain threshold value
2. Bins are "constructed" by dividing the design space into $(M_{seg})^{N_o} = 5^2 = 25$ bins equally sized
3. If a bin is "full", or has a number of points belonging to it that is higher than some predetermined value, block all selections from that bin.
4. Randomly choose a nondominated chromosome from the accumulation file?

Pareto Optimal Set/Front

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"Includes optimal solutions for each of the individual objectives, as well as a range of tradeoff solutions in between, which are themselves optimal solutions. Providing a range of solutions to a multi-objective optimization problem is a powerful approach because it allows the designer to see the effect of decision variable variation on the design space in the form of optimal tradeoffs. Thus, the designer can choose individual objective weighting factors after their full influence is quantitatively known."

Genetic Algorithms Applied to Multi-Objective Aerodynamic Shape Optimization, Terry L. Holst



Pareto
Optimality

Preventing Premature Convergence

Saturday, December 3, 2022 10:55 AM

Goal: Maintain population diversity and selection pressure

Genetic Drift: The tendency to converge on a single solution as the diversity of a population decreases

Fitness sharing: a type of niching, where the fitness of each individual is scaled based on its proximity to others.

Deceptive: A class of challenging problems for conventional genetic algorithms which usually mislead the search to some local optima rather than the global optimum.

Homogeneous vs. Heterogenous

From <<https://ieeexplore.ieee.org/document/4630900>>

From <<https://stackoverflow.com/questions/37836751/what-are-fitness-sharing-and-niche-count-in-evolutionary-computation>>

Review of 24 different approaches:

Pandey, Hari Mohan et al. "A comparative review of approaches to prevent premature convergence in GA." *Appl. Soft Comput.* 24 (2014): 1047-1077.

From <<https://www.semanticscholar.org/paper/A-comparative-review-of-approaches-to-prevent-in-GA-Pandey-Chaudhary/590a7e80ad9fd564e79129acd978bd9d02fe710b>>

Determine convergence by subtracting avg fitness from max fitness. If it is low, then solution is likely well converged

Or by finding avg difference between genes

Metaselection algorithm: To share time among several evolving populations and delete older ones. Complex. Use if no progress has been made for several generations, or stopping conditions have not been met

Encoding

Saturday, December 3, 2022 11:23 AM

"The hardness of a problem is inherently related to the way we represent the problem. Therefore, the difficulty level of the problem can increase or decrease based on the representation" -

Pandey, Hari Mohan et al. "A comparative review of approaches to prevent premature convergence in GA." *Appl. Soft Comput.* 24 (2014): 1047-1077.

Also, check out

Malik, Ms. Shikha and Mr. Sumit Wadhwa. "Preventing Premature Convergence in Genetic Algorithm Using DGCA and Elitist Technique." (2014).

~~X~~ Talk about masking parameters

Penalty Functions

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1. Reject all unfeasible solutions (or a percentage of them) [not really a penalty function]
2. Repair unfeasible solutions by finding the closest feasible one
3. Superiority of feasible solutions over the unfeasible

Hyperparameter Control

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"The parameter control is a challenge in evolutionary computing for increasing the performance and obtaining a "universal" genetic algorithm, to be able to solve problems without manual parameter settings. The goal is to find not only the proper adjustments, but to do this in an efficient way." - R. Maniu and L. A. Dumitru, "Genetic algorithm - adaptive crossover based on solution distribution in search space," 2017

From <<https://ieeexplore.ieee.org/document/7975084>>

Deterministic: This rule modifies the strategy parameter in a fixed, predetermined (i.e., user-specified) way without using any feedback from the search.

Adaptive: Algorithm parameters are modified based on feedback about algorithm evolution

Static: Parameters are preset and do not change

Self-adaptive: Hyperparameters themselves are genetic material changed through evolution (Ex. genetically encoded mutation rates per individual)

Electrostatic Sieve Optimization Setup

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1. Need to figure parameters ✓
2. File I/O ✓
3. Fitness and param array ✓
4. Set up hyperparameters + end condition ✓
5. r.v. init pop. ✓
6. Fitness function ✓
7. Selection
8. Crossover
9. Mutation

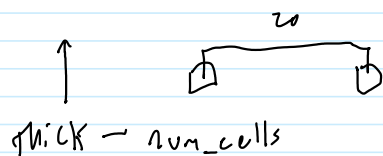
$$\text{fitness} = \% \text{ classified} \cdot \frac{1}{\text{abs}(\text{max_itvl_class} - \text{mean_class})}$$

Parameters:

- Number of electrodes (change thickness, pitch, to be constrained by a penalty or basic functions)
- Potential (float, not constrained [yet])
- Phases (2, 3, 4) (Integer)
- Frequency (makes sure time is consistent)
- ambient or continuous injection (binary)
- Also bias location as function to keep same place

$$\text{size} = \text{pitch} \cdot (\text{num} - 1) + \text{thickness}$$

$1 + \frac{\text{size} - \text{thick}}{\text{pitch}} \downarrow \text{rand 1st}$
 $1 - 10 \text{ integer}$

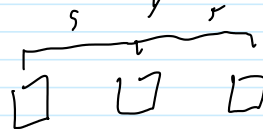


$$\text{pitch} = \frac{\text{size} - \text{thick}}{\text{num} - 1}$$

the check: size is over constrained

$$\frac{\text{size} \cdot \text{thick}}{\text{num} - 1}$$

$$n_{it} = \frac{\text{tot. time}}{\delta t}$$

$$t = \frac{1}{\text{freq}} \cdot \frac{1}{n_{\text{cell}}} \cdot \frac{1}{n_{\text{it_per_phase}}}$$


The diagram shows three square boxes arranged horizontally. Above the first box is the letter 's'. Above the second box is the letter 'r'. A horizontal line with a downward-pointing tick mark at its center spans across all three boxes, with short vertical lines connecting it to the top of each box.

pitch: size

Parallel GA

Wednesday, August 23, 2023 4:13 PM

Ridiculously parallel; a maximum speedup of the population size/number of processing elements (PEs) can be seen

Worker-master:

Master controls reproduction, selection, and genetic operators. Each worker process evaluates a single (or more) function evaluation.

This...

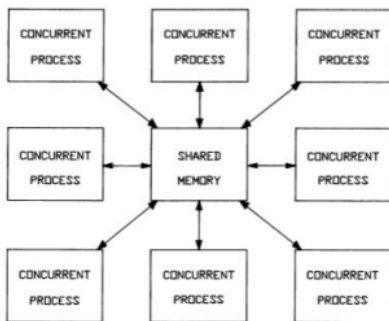
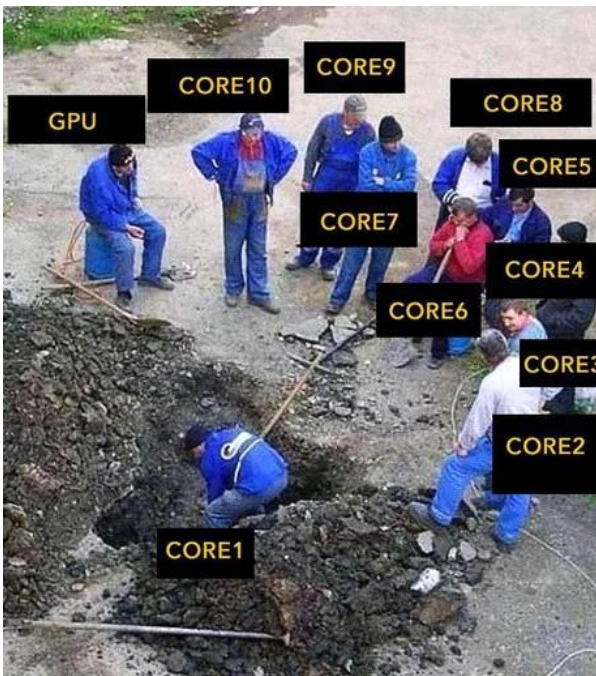


FIGURE 5.35 Schematic of an asynchronous concurrent genetic algorithm.

Not this...



Flow

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