# GENETIC ALGORITHMS: THE BINARY GA 

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Lecturer: Robin T. Bye

## Components of binary GA

## Algorithm flow

1. Define cost function, cost, variables. Select GA parameters.
2. Generate initial population.
3. Decode chromosomes.
4. Find cost for each chromosome.
5. Select mates for reproduction.
6. Mating.

## Algorithm flow

7. Mutation.
8. Check stopping criteria

- IF (reached max number of iterations OR converged) THEN stop
- ELSE go to Step 3.


Figure 2.2 Flowchart of a binary GA.
Adapted from [1].

## Variables and cost function

Nvar-dimensional problem $\rightarrow$ chromosome has Nvar variables
(genes), $i=1, .$. , Nvar

- chrom = [p1, p2,...,pNvar]
- Cost $=f($ chrom $)=f(p 1, p 2, \ldots, p N v a r)$
- Example: 2D height map in xy-plane
- chrom = $[\mathrm{x}, \mathrm{y}]$
- cost $=$ height $=f($ chrom $)=f(x, y)$


## Variables and cost function

- If too many variables $\rightarrow$ slow GA
- Eg. $f=2 x+3 y+z / 10000+\sqrt{ } w / 9876$ with constraints $1 \leq x, y, z, w \leq 10$
- Due to constraints, $z$, w terms relatively small $\rightarrow$ ignore: $f=2 x+3 y$
- Variable interaction (epistasis)
- GA good for medium/high interaction
- Random search good for high interaction
- Minimum-seeking good for low interaction


## Encoding/decoding

- Encoding: Convert variable values to binary genes
- Decoding: Convert binary genes back to human-readable variable values
- Example:

| Bin | Dec | Numbers | Alt. Numbers | Colour | Speed |
| :--- | :--- | :--- | :--- | :--- | :--- |
| 00 | 0 | 10 | 13.75 | Red | Slow |
| 01 | 1 | 20 | 21.25 | Green | Medium |
| 10 | 2 | 30 | 28.75 | Blue | Fast |
| 11 | 3 | 40 | 36.25 | Yellow | Superfast |

## Encoding/decoding

- Example continued:
- gene1 $=01 \Leftrightarrow$ medium
- gene2 $=10 \Leftrightarrow$ fast
- gene3 $=11 \Leftrightarrow$ superfast
- gene4 $=00 \Leftrightarrow$ slow
- chrom $=$ [gene1, gene2, gene3, gene4]
$=[01101100]=[$ med,fast,supfast,slow]


## Encoding/decoding

- Goal: Sort categories in increasing order (slow,medium,fast,superfast)
- Cost: 0 for correct place, 1 for one place off, 2 for two places off, etc.
- [01101100] =[medium,fast,superfast,slow]

$$
\cdot \rightarrow \text { Cost }=1+1+1+3=6
$$

- [00100111] =[slow,fast,medium,superfast]
- $\rightarrow$ Cost $=0+1+1+0=2$


## Encoding/decoding

- Number of bits Nbits in chromosome:
- Ngene = number of bits in each gene/var
- Nvar = number of genes/variables
- Nbits $=$ Ngene $\times$ Nvar $=$ number of bits


## Population

- Set of Npop chromosomes
- Each chromosome has Nbits
- Represented as matrix of binary digits
- Dimensions are Npop $\times$ Nbits
- Initial population randomly assigned:
- pop=round(rand(Npop, Nbits));


## Natural selection

1. Rank chromosomes (low cost better)
2. Only keep best fraction (selection rate Xrate) of Npop chromosomes $\rightarrow$ Nkeep
$=$ Xrate $\times$ Npop chromosomes survives
3. Let kept chromosomes mate and replace discarded chromosomes

## Pairing methods

- From top to bottom (1+2, 3+4, etc.)
- Uniform random pairing
- Weighted random pairing
- rank weighting
- cost weighting
- Tournament selection
- Others


## Mating

- Randomly pick a crossover point
- Parent1 passes left-bits to offspring1 and right-bits to offspring2
- Parent2 passes left-bits to offspring 2 and right-bits to offspring1
- $\mathrm{p} 1=[\mathrm{L} 1 \mid \mathrm{R} 1], \mathrm{p} 2=[\mathrm{L} 2 \mid \mathrm{R} 2] \rightarrow$
- $\mathrm{o} 1=[\mathrm{L} 1 \mid \mathrm{R} 2], \mathrm{o} 2=[\mathrm{L} 2 \mid \mathrm{R} 1]$
- Other schemes exist


## Elitism

- Always keep best chromosome in population and never mutate it!
- Do not throw away a good solution!


## Next generation

- Insert offspring into population
- Recalculate costs and repeat process until
- convergence
- max number of iterations reached
- you are happy for some reason


## Example 2D problem

## Example 2D problem




Three-dimensional view of the cost surface with a view of Long's Peak.
Adapted from [1].

## Example 2D problem

## Encoding

TABLE 2.2 Binary Representations

| Variable | Binary | Decimal | Value |
| :--- | :---: | :---: | :---: |
| Latitude | 0000000 | 1 | $40^{\circ} 15^{\prime}$ |
| Latitude | 1111111 | 128 | $40^{\circ} 16^{\prime}$ |
| Longitude | 0000000 | 1 | $105^{\circ} 36^{\prime}$ |
| Longitude | 1111111 | 128 | $105^{\circ} 37^{\prime} 30^{\prime \prime}$ |

$$
\text { chromosome }=[\underbrace{1100011}_{x} \underbrace{0011001}_{y}]
$$

Adapted from [1].

## Example 2D problem

## Initial population

| TABLE 2.3 Example Initial Population of 8 <br> Random Chromosomes and Their Corresponding <br> Cost |  |
| :--- | ---: |
| Chromosome | Cost |
| $\mathbf{* 0 0 1 0 1 1 1 1 0 0 0 1 1 0}$ | -12359 |
| 11100101100100 | -11872 |
| $* 00110010001100$ | -13477 |
| $* 00101111001000$ | -12363 |
| 11001111111011 | -11631 |
| 01000101111011 | -12097 |
| $* 11101100000001$ | -12588 |
| 01001101110011 | -11860 |

[^0]

## Example 2D problem

## Natural selection

TABLE 2.4 Surviving Chromosomes after a 50\%
Selection Rate

## Best 50\% <br> Nkeep = 4

| Chromosome | Cost |
| :---: | :---: |
| $* 00110010001100$ | -13477 |
| $* 11101100000001$ | -12588 |
| $* 00101111001000$ | -12363 |
| $* 00101111000110$ | -12359 |

$$
N_{\text {keep }}=X_{\text {rate }} N_{\text {pop }}
$$

## Example 2D problem

## Crossover



Figure 2.11 Two parents mate to produce two offspring. The offspring are placed into the population.

## Example 2D problem

Create offspring and replace bad chromosomes
TABLE 2.7 Pairing and Mating Process of SinglePoint Crossover

| Chromosome | Family | Binary String |
| :--- | :--- | :---: |
| 3 | $\mathrm{ma}(1)$ | 00101111001000 |
| 2 | pa(1) | 11101100000001 |
| 5 | offspring $_{1}$ | 00101100000001 |
| 6 | offspring $_{2}$ | 11101111001000 |
| 3 | ma(2) | 00101111001000 |
| 4 | pa(2) | 00101111000110 |
| 7 | offspring $_{3}$ | 00101111000110 |
| 8 | offspring $_{4}$ | 00101111001000 |

## Example 2D problem

New population after mating

## TABLE 2.8 Mutating the Population

| Population after Mating | Population after Mutations | New Cost |
| :--- | :---: | :---: |
| 00110010001100 | 00110010001100 | -13477 |
| 11101100000001 | 11101100000001 | -12588 |
| 00101111001000 | 00101111010000 | -12415 |
| 00101111000110 | 00001011000111 | -13482 |
| 00101100000001 | 00101000000001 | -13171 |
| 11101111001000 | 11110111010010 | -12146 |
| 00101111000110 | 00100111001000 | -12716 |
| 00101111001000 | 00110111001000 | -12103 |

Adapted from [1].

## Example 2D problem

Members of population after first generation


Figure 2.12 A contour map of the cost surface with the 8 members at the end of the first generation.

## Example 2D problem

New ranked population at start of second generation

| New best $\rightarrow$ chromosome | TABLE 2.9 New Ranked Population at the Start of the Second Generation |  |
| :---: | :---: | :---: |
|  | Chromosome | Cost |
|  | 00001011000111 | -13482 |
|  | 00110010001100 | -13477 |
|  | 00101000000001 | -13171 |
|  | 00100111001000 | -12716 |
|  | 11101100000001 | -12588 |
|  | 00101111010000 | -12415 |
|  | 11110111010010 | -12146 |
|  | 00110111001000 | -12103 |

## Example 2D problem

Population after crossover/mutation in 2nd generation TABLE 2.10 Population after Crossover and Mutation in the Second Generation

|  | Chromosome | Cost |
| :--- | :--- | :---: |
| Note that 2nd | 00001011000111 | -13482 |
| best | 00110000001000 | -13332 |
| chromosome | 01101001000001 | -12923 |
| has been | 0110011101000 | -12128 |
| replaced by | 10100111000001 | -12961 |
| one with | 10100010001000 | -13237 |
| higher cost | 00110100001110 | -13564 |
|  | 00100010000001 | -13246 |
|  |  | AdapR8d from [1]. |

## Example 2D problem

Members of population after 2nd generation


Adap $\mathrm{E}^{\mathrm{Od}}$ from [1].

## Example 2D problem

- Example converged after only 3 gen's
- Height found: 14199 m


Figure 2.15 Graph of the mean cost and minimum cost for each generation. Ad3apted from [1].

## References

[1] Haupt \& Haupt, Practical Genetic Algorithms, 2nd Ed., Wiley, 2004.


[^0]:    * best chromosomes

