

Assignment 44

Assuming that the rules for the 2-point-cross-over recombination-operator are analogous to the 1-point-cross-over recombination-operator there can only be two cases: The genome is split at two points. Either:

1. The part up to point 1 is taken from parent *A*, the part between point 1 and point 2 is taken from parent *B* and the final part from point 2 to the end is again taken from parent *A*.
2. or the part up to point 1 is taken from parent *B*, the part between point 1 and point 2 is taken from parent *A* and the final part from point 2 to the end is again taken from parent *B*.

So, regardless of the genome sequence length *L*, there can only be two different offsprings. Thus, the diversity of the population has to be provided by the size of the population or a mutation operator.

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Assignment 45

A binary genome of length *L* can be represented by a hypercube of dimension *L* of unit edge length. Every bit of the genome corresponds to an axis of the hypercube. Since the hypercube has unit edge length and the genome is binary, a genome corresponds to a specific corner of the hypercube. This is depicted for the three-bit-genomes (0,0,1), (1,1,0), (1,0,1) in figure (1).

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Assignment 46

Generating all possible sequences of *L* is equivalent to finding all permutations of *L*. Since every permutation can be expressed as a product of transpositions (the swapping of two elements), the algorithm can generate all sequences.

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Assignment 47

Done, needs further explanation $1 - (N \cdot (1 - p)^L)$

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Assignment 48

External selection selects the individuals to "survive", while the rest of the population is discarded. *Parent selection* selects those "survivors" which will generate new offspring to replenish the population.

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Assignment 49

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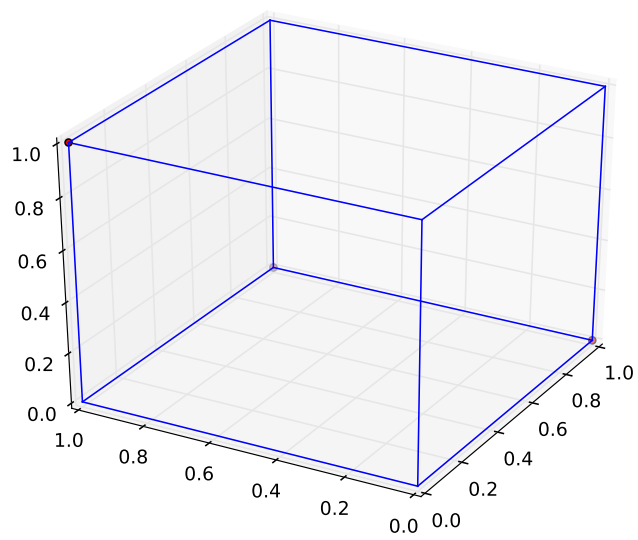


Figure 1: representation of three-bit-genomes by an three dimensional hypercube