Multimodal Optimization, Dynamic Optimization

- Particularities of multi-modal optimization
- Mechanisms for dealing with multiple optima
- Particularities of dynamic optimization
- Mechanisms for dealing with dynamic objective functions

Particularities of multi-modal optimization

- Aim: find all optima (global and/or local) of the objective function
- Motivation:
 - give to the decision maker not a single optimal solution but a set of good solutions
 - find all solutions with local optimal behavior
- Similar (in some sense) with: multiobjective optimization
- Applications:
 - Optimal design in engineering (e.g. induction motors)
 - Data analysis (e.g. clustering)
 - Protein structure prediction

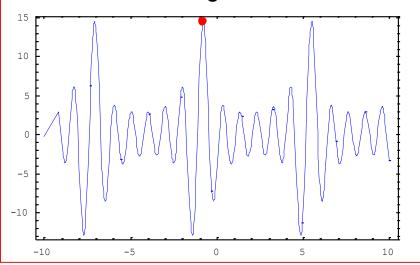
Global vs multi-modal optimization

Global optimization

Aim: find a global optimum

Evolutionary approach: population concentrates on the global optima (a single powerful species)

Premature convergence: bad

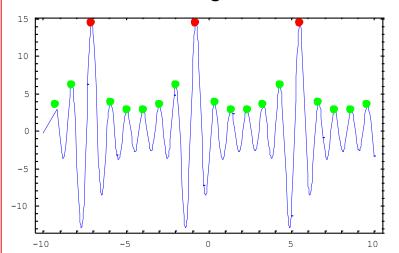


Multimodal optimization

Aim: find all (global/ local) optima

Evolutionary approach: multiple species are formed each one identifying an optimum

Premature convergence: not so bad



Approaches in multi-modal optimization

Implicit speciation (also called niching)

- species emerges in the population
- niching = finding and preserving multiple stables niches in the search space (around the potential optima)
 - Sequential the niches are identified in several stages characterized by different mechanisms or values of some control parameters (e.g. resolution of the search)
 - Parallel the niches are identified during one evolutionary process (based on specific selection mechanisms)
- Mechanisms which induce the formation of species:
 - fitness sharing
 - crowding
- Remark: it is usual combined with archiving (the "good" configurations are collected in the population or in an additional archive)

Approaches in multi-modal optimization

Explicit speciation

- The population is divided into communicating subpopulations which evolves in parallel; the communication should be rare in order to allow the preservation of the species (and avoid the full mixing of the population element)
- The division of a population in subpopulations can be based on a clustering process (based on a specific similarity measure)
- Each subpopulation corresponds to a species whose aim is to populate a niche in the fitness landscape and to identify an optimum

Implicit speciation: fitness sharing

Fitness sharing:

 Discourage the agglomeration of many elements in the same search region by penalizing the fitness function based on a distance related factor

[see lecture 10 – fitness sharing is used also for multiobjective optimization]

Related idea: clearing [Petrowski, 1997] = the elements in the same niche are eliminated (by setting their fitness to 0)

```
Pseudo-code for Clearing technique
function Clearing(\sigma, C) /* Function definition*/
begin
Sort_Fitness (P) /* Sort the population in descending or-
der according to their fitness*/
 for i = 0 to (Np - 1)
                                               Niche radius
       if (Fitness(P[i]) > 0)
        n Win = 1
        for j = (i + 1) to(Np - 1)
         if (Fitness(P[j]) > 0 AND Distance(P[i], P[j]) < \sigma)
           ifn_Win < C \leftarrow
           n_Win = n_Win + 1
           else
                                             Niche size
           Fitness (P[i]) = 0
           endif
         endif
        endfor
       endif
  endfor
 end
```

[S. Das et al. Real-parameter evolutionary multimodal optimization – a survey, SWEVO, 2011]

Implicit speciation: crowding

Main idea:

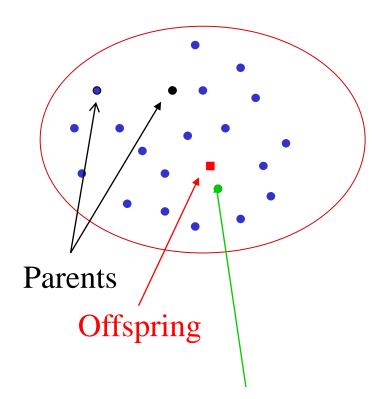
- Elements belonging to different species should not compete between them
- The selection process involve similar elements → this allows to maintain the existing diversity in the population

Implementation:

- A new element replaces the closest old element in a population sample (the ratio between the population sample and the population size is called crowding factor, CF); CF=1 → global crowding
- The distance between elements is computed in the search space

Implicit speciation: crowding

Idea of crowding



- Particularities
 - preserves the population diversity
 - encourages the species formation
 - the final population consists of elements concentrated around the optima
- Advantage:
 - simple
- Disadvantage:
 - global character of crowding (O(m²))

Element to be replaced

Implicit speciation: crowding

- Implementation of deterministic crowding [Mahfoud, 1995]
- to be applied for m/2 times (m=pop size)

Pseudo-code for deterministic crowding

- 1. Select two parents p_1 , p_2 randomly with no replacement.
- 2. Perform a crossover between them yielding offspring c_1 , c_2 .
- 3. Apply mutation operator to generate c'_1 , c'_2 .
- 4. if $[d(p_1, c_1') + d(p_2, c_2') \le d(p_1, c_2') + d(p_2, c_1')]$
 - if $f(c_1') \ge f(p_1)$ replace p_1 with c_1'
 - if $f(c_2') \ge f(p_2)$ replace p_2 with c_2'

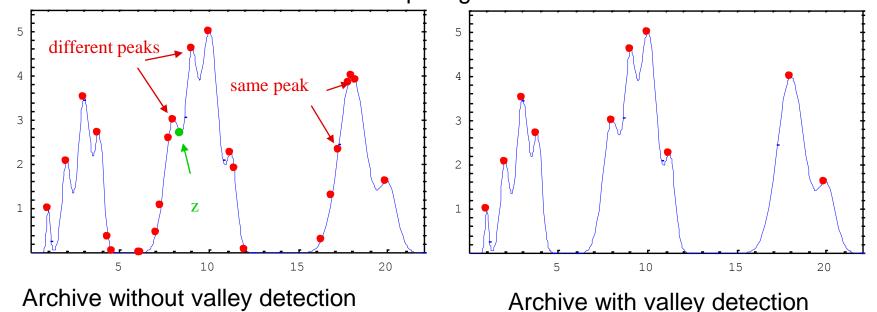
else

- if $f(c_2') \ge f(p_1)$ replace p_1 with c_2'
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Archiving

- Hill-valley function [Ursem, Multinational Evolutionary Algorithms, 1999]:
 - If there exists c in (0,1) such that z=cx+(1-c)y implies f(z)<f(x) and f(z)<f(y)</p>
 - then there exists a valley between x and y → they belong to different "peaks" → both of them may remain in the archive
 - The decision is based on computing z for some values of c



Explicit speciation

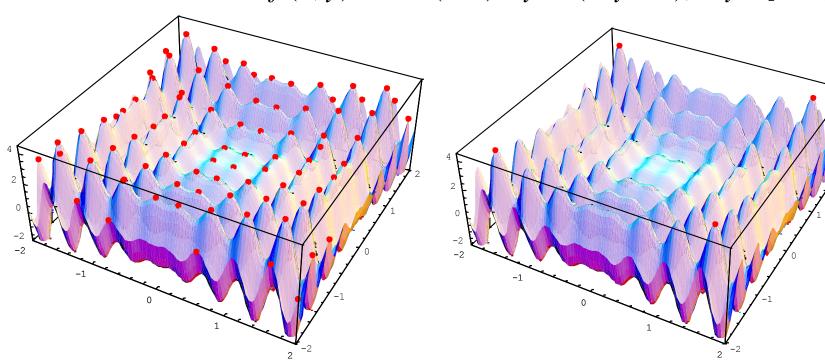
Main idea:

- the population is explicitly divided into sub-populations
- each sub-population aims to identify one or several optima
- Question: which is the influence of the communication between subpopulations?
 - Example: Multipopulation Multiresolution DE (2004)
 - Notations:
 - m = number of elements in the subpopulation
 - s = number of subpopulations
 - g = Number of generations
 - f = frequency of migration (generations between 2 migration stages)
 - p_m= migration probability

Explicit speciation

Test function: multi-peaks

$$f(x, y) = x \sin(4\pi x) - y \sin(4\pi y + \pi), \ x, y \in [-2, 2]$$

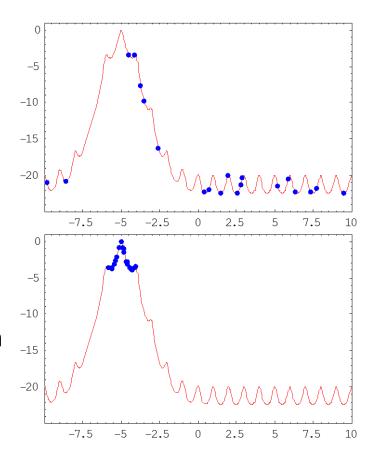


m=5, s=50, g=50, e=20, f=0, p_m =0 (91 elements in the archive)

m=5, s=50, g=100, f=10, e=20, p_m =0.5 (4 elements in the archive)

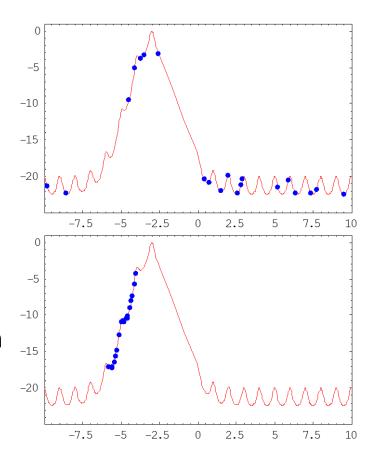
- Dynamic optimization:
- Aim: track a changing optimum
- Difficulty: inability to track the optimum
- Cause: the population lost its diversity
- Solution: stimulating the population diversity

$$f(t, x^*(t)) \ge f(t, x), x \in D^n$$



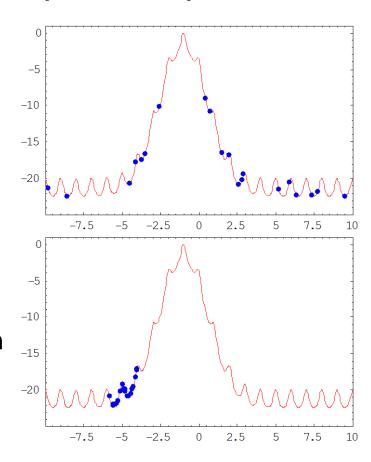
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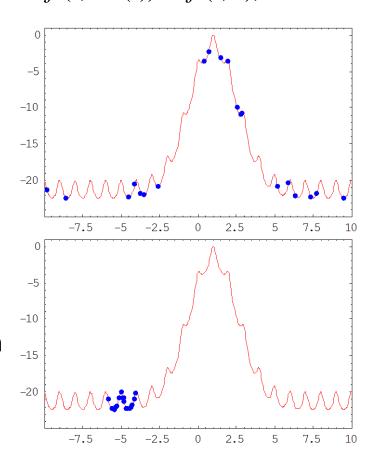
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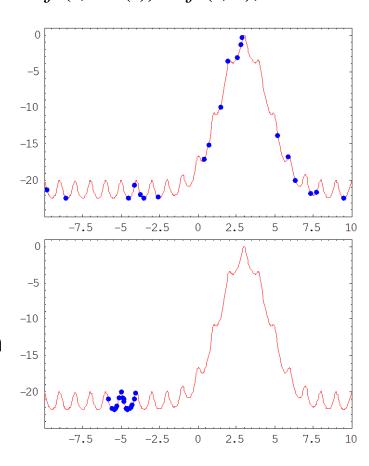
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Dynamic optimization - characteristics

- Types of optimum changes:
 - continuous trajectory of the optimum (smooth changes)
 - discontinuous trajectory of the optimum (jumps of the optimum position)

Main questions:

- Time-linkage: the future behaviour depends on the current state?
- Detectability: can be the changes detected?
- Predictability: can be the changes predicted? are the changes periodic?

Dynamic optimization – problems

- Practical applications:
 - dynamic scheduling
 - dynamic resource allocation
 - dynamic risk minimization
 - other problems with parameters changing in time
- Test problems
 - Moving Peaks Benchmark MPB [Branke, 1999]
 - Dynamic Knapsack Problem time dependent weights and profits
 - Dynamic TSP time dependent costs

- Reactive approaches: trigger a diversity increasing mechanism when a change is detected
 - hypermutation (high mutation rate)
 - random immigrants (random elements are introduced in the population)
 - memory useful in the case of recurrent or periodic changes
 - Implicit (diploid elements)
 - Explicit (store the best elements at various stages of the evolution and re-use them when a change is detected) – similar to archiving

Remarks:

- Appropriate for discontinuous changes of the optima
- How can be detected a change?
 - Use of some detectors (elements of the population which are periodically re-evaluated) - if their current fitness value is different from the previous one then a changed occurred

Diploid elements:

- usually applied for genetic algorithms with binary encoding
- each element ,X_i, from the population consists of two chromosomes C1(i) and C2(i)
- when the fitness of X_i is evaluated only the dominant components are taken into account

```
If C1(i,j)= C2(i,j)
then X_i (j)=C1(i,j) // same value of component j in both chromosomes
Else
if rand(0,1)<Pd(j,g) then X_i (j)=C1(i,j)
else X_i (j)=C2(i,j)
```

Diploid elements:

The selection of the dominant component is based on the dominance probability vector Pd(j,g) which is adjusted at each generation:

$$Pd(j,g+1)=(1-alpha)*Pd(j,g)+alpha*Pbest(j,g)$$

- Variants:
 - Pbest(g) = mask corresponding to the best element in the population
 i.e. Pbest(j,g)=1 if the component j from the best element is taken from C1
 [S. Yang, Dominanc Learning in Diploid GA for DOP, 2006]
 - Pbest(g) = frequency vector computed based on the best vectors identified during the evolutionary process

- Proactive approaches: maintain the population diversity by:
 - diminishing the selection pressure (sharing, crowding)
 - directly stimulating the diversity (periodically inserting random elements)

Remarks

 the proactive approaches are appropriate for continuously changing optima