# 5. Epistasis. Linkage identification. Optimization by model fitting.

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- ✓ Epistasis, short example
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# **Introduction to Epistasis**

# **Black-box Optimization and Genetic Algorithms**

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## In BBO, we need to specify:

- ✓ representation of candidate solution
- ✓ objective function (gives the quality of candidate solution)
  - can have almost any form (nondifferentiable, discontinuous, multimodal, noisy, ...)

## How to solve BBO problems?

- ✓ Algorithm applicable in BBO scenario cannot require any other knowledge about the objective function
- ✔ Hill-climbing, simulated annealing, taboo, GAs, ...

## GAs are popular:

- ✓ they are easy to use
- ✓ appliable without prior knowledge
- ✓ easy to parallelize

## GAs are great, but not perfect!!!

## GA works well...

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Summary

## Problem $f_1$ :

- ✓ defined over 40-bit strings
- ✓ quality of the worst solution:  $f_1(x^{\text{worst}}) = 0$ .
- ✓ quality of the best solution:  $f_1(x^{\text{opt}}) = 40$ .
- ✓ best solution:  $x^{\text{opt}} = (1111...1)$ .

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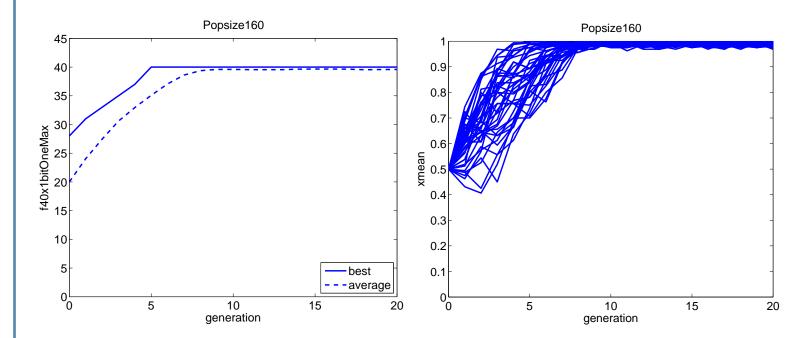
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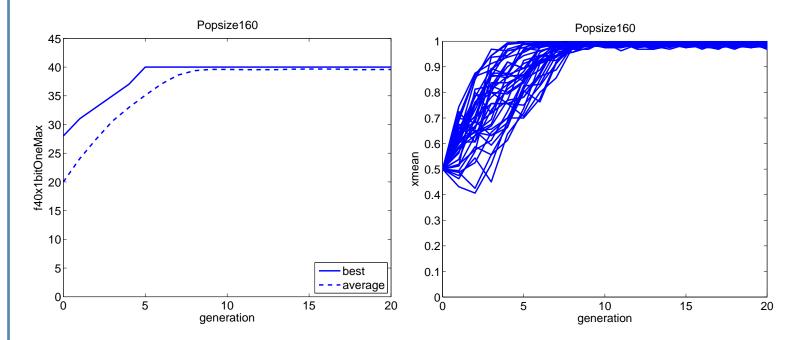
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Problem  $f_1$  contains no epistatic interactions among design variables.

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## Problem $f_2$ :

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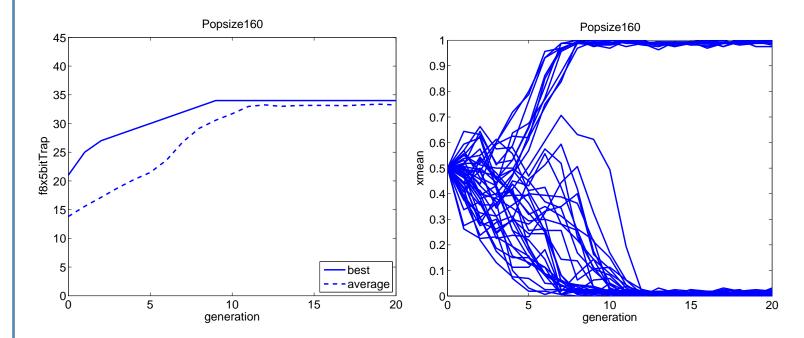
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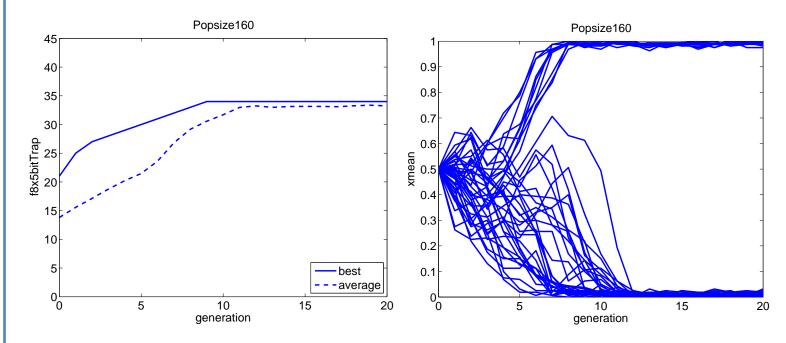
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Problem  $f_2$  contains some interactions among variables, *GA* is not aware of them and works with the individual bits as if they were truly independent of each other.

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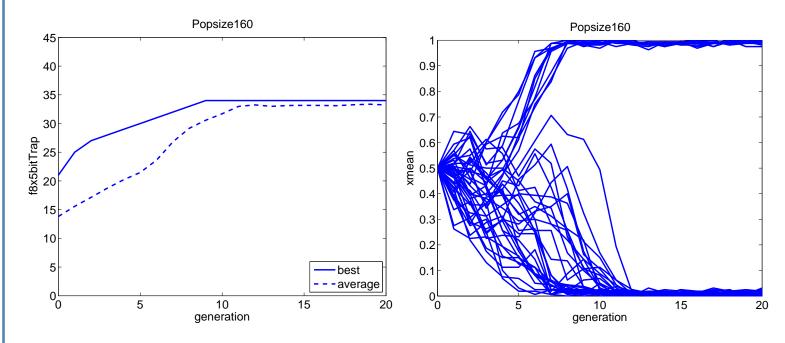
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Problem  $f_2$  contains some interactions among variables, *GA* is not aware of them and works with the individual bits as if they were truly independent of each other.

None of the above mentioned problem characteristics is important to judge if the GA will work well!!!

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# GA works again...

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#### GA works again...

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## Still solving $f_2$ :

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Instead uniform crossover

✓ let us allow the crossover only after each 5th bit

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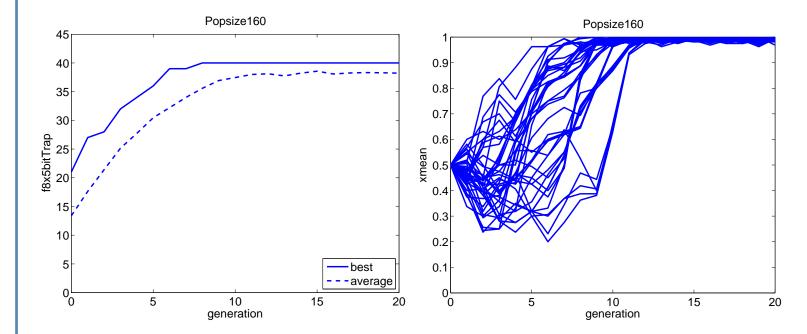
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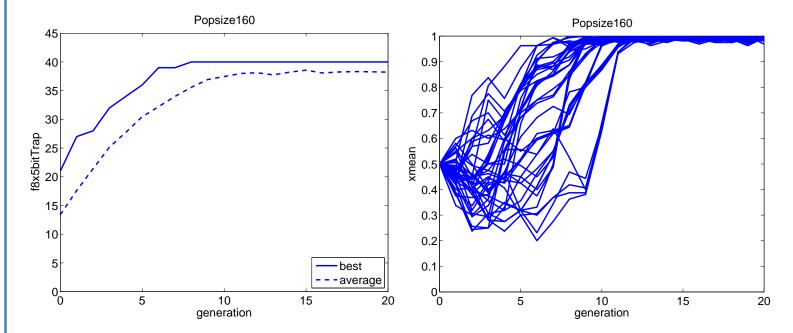
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### Instead uniform crossover

✓ let us allow the crossover only after each 5th bit



Problem  $f_2$  contains some interactions among variables and *GA* knows about them.

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Summary

### **Epistasis:**

✓ Effects of one gene are dependent on (influenced, conditioned by) other genes.

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### Epistasis

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Summary

## **Epistasis:**

✓ Effects of one gene are dependent on (influenced, conditioned by) other genes.

### Linkage:

✓ Tendency of certain loci or alleles to be inherited together.

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Summary

## **Epistasis:**

✓ Effects of one gene are dependent on (influenced, conditioned by) other genes.

## Linkage:

✓ Tendency of certain loci or alleles to be inherited together.

### Other names:

- ✓ dependencies
- ✓ interdependencies
- ✓ interactions

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### Linkage:

✓ Tendency of certain loci or alleles to be inherited together.

### Other names:

- ✓ dependencies
- ✓ interdependencies
- ✓ interactions

## When optimizing the following functions, which of the variables are linked together?

$f = x_1 + x_2 + x_3$	(1)
$f = 0.1x_1 + 0.7x_2 + 3x_3$	(2)
$f = x_1 x_2 x_3$	(3)
$f = x_1 + x_2^2 + \sqrt{x_3}$	(4)
$f = \sin(x_1) + \cos(x_2) + e^{x_3}$	(5)
$f = \sin(x_1 + x_2) + e^{x_3}$	(6)

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Summary

Which of the semestral projects contain interactions? Why?✓ Circles in the square

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Which of the semestral projects contain interactions? Why?

- ✔ Circles in the square
- ✓ Japanese puzzle

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Which of the semestral projects contain interactions? Why?

- ✔ Circles in the square
- ✓ Japanese puzzle
- ✓ MTSP

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Which of the semestral projects contain interactions? Why?

- ✔ Circles in the square
- ✓ Japanese puzzle
- ✓ MTSP
- ✓ Image compression

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Which of the semestral projects contain interactions? Why?

- ✔ Circles in the square
- ✓ Japanese puzzle
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- ✓ Image compression

### Notes:

✓ Almost all real-world problems contain interactions among design variables.

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Which of the semestral projects contain interactions? Why?

- ✔ Circles in the square
- ✓ Japanese puzzle
- ✓ MTSP
- ✓ Image compression

### Notes:

- ✓ Almost all real-world problems contain interactions among design variables.
- ✓ The "amount" and "type" of interactions depend on the representation and the objective function.

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Which of the semestral projects contain interactions? Why?

- ✔ Circles in the square
- ✓ Japanese puzzle
- ✓ MTSP
- ✓ Image compression

### Notes:

- ✓ Almost all real-world problems contain interactions among design variables.
- ✓ The "amount" and "type" of interactions depend on the representation and the objective function.
- ✓ Sometimes, by a clever choice of the representation and the objective function, we can get rid of the interactions.

# **Linkage Identification Techniques**

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Problems:

✓ How to detect dependencies among variables?

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### Problems:

- ✓ How to detect dependencies among variables?
- ✓ How to use them?

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Summary

### Problems:

- ✓ How to detect dependencies among variables?
- ✓ How to use them?

Techniques used for image identification:

- 1. Indirect detection along genetic search (messy GAs)
- 2. Direct detection of fitness changes by perturbation
- 3. Model-based approach: classification
- 4. Model-based approach: distribution estimation (EDAs)

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Linkage identification by non-linearity check (LINC)

LINC: Example

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# **Perturbation Techniques of Linkage Identification**

## LINC

✓ Developed by Munetomo [MG98], based on gene expression messy GA [Kar95]

## Linearity and nonlinearity

- ✓ consider a bitstring  $\mathbf{x} = (x_1, x_2, ..., x_D)$
- ✓ perturb loci i, j, and both i and j

$$\begin{array}{l} \mathbf{x} = (x_1, \dots, x_i, \dots, x_j, \dots, x_D) \\ \mathbf{x}_i = (x_1, \dots, \bar{x}_i, \dots, x_j, \dots, x_D) \\ \mathbf{x}_j = (x_1, \dots, x_i, \dots, \bar{x}_j, \dots, x_D) \\ \mathbf{x}_{ij} = (x_1, \dots, \bar{x}_i, \dots, \bar{x}_j, \dots, x_D) \end{array} \begin{array}{l} \Delta f_i = f(\mathbf{x}_i) - f(\mathbf{x}) \\ \Delta f_j = f(\mathbf{x}_j) - f(\mathbf{x}) \\ \Delta f_{ij} = f(\mathbf{x}_{ij}) - f(\mathbf{x}) \end{array}$$

~

If  $\Delta f_{ij} = \Delta f_i + \Delta f_j$ ,

- ✓ fitness is linear, interaction did not show up,
- ✓  $x_i$  and  $x_j$  needn't be part of the same BB.

If  $\Delta f_{ij} \neq \Delta f_i + \Delta f_j$ ,

- ✓ fitness is nonlinear, interaction exists,
- $x_i$  and  $x_j$  are members of the same BB,
- ✓ add i to j's linkage set and j to i's linkage set.

- [Kar95] H. Kargupta. Search, polynomial complexity, and the fast messy genetic algorithm. Technical Report IlliGAL Report No. 95008, Illinois Genetic Algorithm Laboratory, Urbana Champaign, Illinois, 1995.
- [MG98] Masaharu Munetomo and David E. Goldberg. Identifying linkage by nonlinearity check. Technical Report IlliGAL Report No. 98012, Illinois Genetic Algorithm Laboratory, Urbana Champaign, Illinois, 1998.

## LINC: Example

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## 3-bit Trap Function:

$u(\mathbf{x})$	0	1	2	3
$f(\mathbf{x})$	0.6	0.3	0	1

✓ Let us consider string  $\mathbf{x} = (010)$  and bits 1 and 2:

x = (010)				
$\mathbf{x}_1 = (110)$	$\Delta f_1$	= f(110) - f(010)	= 0 - 0.3	= -0.3
$\mathbf{x}_2 = (000)$	$\Delta f_2$	= f(000) - f(010)	= 0.6 - 0.3	= 0.3
$\mathbf{x}_{12} = (100)$	-	=f(100)-f(010)	= 0.3 - 0.3	= 0

 $\Delta f_{12} = \Delta f_1 + \Delta f_2 \longrightarrow i$  and *j* needn't be part of the same BB

✓ The same string  $\mathbf{x} = (010)$  and bits 1 and 3:

x = (010)				
$\mathbf{x}_1 = (110)$	$\Delta f_1$	= f(110) - f(010)	= 0 - 0.3	= -0.3
$\mathbf{x}_3 = (011)$	$\Delta f_3$	=f(011) - f(010)	= 0 - 0.3	= -0.3
$\mathbf{x}_{13} = (111)$	$\Delta f_{13}$	=f(111) - f(010)	= 1 - 0.3	= 0.7

 $\Delta f_{13} \neq \Delta f_1 + \Delta f_3 \longrightarrow i \text{ and } j \text{ are part of the same BB}$ 

Analysis of the string 010 suggests linkage between 1st and 3rd bit only. Additional string (e.g. 011) would suggest that all three bits are linked together.

# Linkage identification by non-monotonicity detection (LIMD)

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(LINC) LINC: Example

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### LIMD:

- ✓ Developed by Munetomo [MG99a, MG99b]
- ✓ Not all nonlinearities detected by LINC are bad for GAs (or hill-climber)
- ✓ Consider 4 bitstrings (as in case of LINC):  $\mathbf{x}$ ,  $\mathbf{x}$ <sub>*i*</sub>,  $\mathbf{x}$ <sub>*j*</sub>, and  $\mathbf{x}$ <sub>*ij*</sub>
- ✓ Reorder them so that  $\mathbf{x} = \arg\min(f(\mathbf{x}), f(\mathbf{x}_i), f(\mathbf{x}_i), f(\mathbf{x}_{ij}))$
- ✔ Monotonicity constraints:
  - $\bigstar \quad f(\mathbf{x}) < f(\mathbf{x}_i) < f(\mathbf{x}_{ij})$
  - $\bigstar \quad f(\mathbf{x}) < f(\mathbf{x}_j) < f(\mathbf{x}_{ij})$
- ✓ If monotonicity is violated, add *i* to *j*'s linkage set and *j* to *i*'s linkage set.

- [MG99a] Masaharu Munetomo and David E. Goldberg. Identifying linkage groups by nonlinearity/non-monotonicity detection. In *Proceedings of the 1999 Genetic and Evolutionary Computation Conference*, pages 433–440, 1999.
- [MG99b] Masaharu Munetomo and David E. Goldberg. Linkage identification by non-monotonicity detection for overlapping functions. *Evolutionary Computation*, 7(4):377–398, 1999.

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### For each pair of variables, we know if the two variables are linked. So what?

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For each pair of variables, we know if the two variables are linked. So what?

- ✓ Construct linkage groups (decompose the problem)
- ✓ Use modified crossover: always transfer the bits in the linkage group together
- ✓ Use modified mutation: allow mutations of the whole linkage group

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What if the problem cannot be decomposed sufficiently?

- ✓ No clearly separated groups of bits
- ✓ Solution: tightness detection

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What if the problem cannot be decomposed sufficiently?

- ✓ No clearly separated groups of bits
- ✓ Solution: tightness detection

### Algorithm:

- 1. Given: linkage groups from LIMD (or LINC) procedure
- 2. Tightness definition:

 $tightness(i,j) = \frac{n(i \wedge j)}{n(i \vee j)} \in \langle 0,1 \rangle,$ 

### where

 $n(i \land j)$  is the number of linkage groups that contain both *i* and *j*, and  $n(i \lor j)$  is the number of linkage groups that contain *i* or *j*.

3. If  $tightness(i, j) < \delta$ , remove locus *j* from linkage set of locus *i* and vice versa.

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Descriptive vs. Generative Models Learnable Evolution Model (LEM)

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# **Optimization by Model Fitting**

### **Modeling the Interactions**

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Descriptive vs. Generative Models Learnable Evolution Model (LEM)

Summary

What kind of "models" should we use?

- ✓ The perturbation techniques
  - ★ extract *global information* from the population, and
  - ✗ use modified recombination operators that use the learned information, i.e. they used modified sampling process.

### **Modeling the Interactions**

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  - ✗ use modified recombination operators that use the learned information, i.e. they used modified sampling process.
- ✓ Classification techniques can be used to
  - ✗ build a model distinguishing the "good" individuals from the "bad" individuals, and
  - ★ sample offspring from the "good" areas (more often than from the "bad" ones).

### **Modeling the Interactions**

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- ✓ Classification techniques can be used to
  - build a model distinguishing the "good" individuals from the "bad" individuals, and
  - ★ sample offspring from the "good" areas (more often than from the "bad" ones).
- ✓ Probability distribution models can be used to
  - ★ build a model describing the distribution of "good" individuals, and
  - **×** sample offspring from that model.

# **Optimization by Model Fitting: The Algorithm**

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Fitting: The Algorithm

Descriptive vs. Generative Models Learnable Evolution Model (LEM)

```
Algorithm 1: General Evolutionary Scheme
 1 begin
          \mathcal{M}^{(0)} \leftarrow \texttt{InitializeModel()}
 2
        X^{(0)} \leftarrow \texttt{Sample}(\mathcal{M}^{(0)})
 3
        f^{(0)} \leftarrow \texttt{Evaluate}(X^{(0)})
 4
          g \leftarrow 1
 5
           while not TerminationCondition() do
 6
                   \{\mathcal{S}, \mathcal{D}\} \leftarrow \texttt{Select}(X^{(g-1)}, f^{(g-1)})
 7
                  \mathcal{M}^{(g)} \leftarrow \texttt{Update}(g, \mathcal{M}^{(g-1)}, X^{(g-1)}, f^{(g-1)}, \mathcal{S}, \mathcal{D})
 8
                  X_{\text{Offs}} \leftarrow \texttt{Sample}(\mathcal{M}^{(g)})
 9
                  f_{\text{Offs}} \leftarrow \texttt{Evaluate}(X_{\text{Offs}})
10
               \{X^{(g)}, f^{(g)}\} \leftarrow \texttt{Replace}(X^{(g-1)}, X_{\text{Offs}}, f^{(g-1)}, f_{\text{Offs}})
11
                  g \leftarrow g + 1
12
```

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- ✓ given a point, they are able to
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- ✓ given a point, they are able to
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Many models are *descriptive* in the sense that

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- ✓ *Generative* models can be used as data generators.

How to turn a descriptive model into generative?

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How to turn a descriptive model into generative?

- ✓ Use (weighted) rejection sampling (see [Luk09, algorithms 115 and 117])Problems with rejection sampling?
  - ✓ In later stages of evolution, the acceptance region becomes very small.
  - ✓ It becomes increasingly difficult to hit that region.

[Luk09] Sean Luke. Essentials of Metaheuristics. 2009. available at http://cs.gmu.edu/~sean/book/metaheuristics/.

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LEM: Evolutionary process guided by machine learning

- ✓ created by Ryszard Michalski [Mic00]
- ✓ originally, alternates between 2 phases:
  - **★** machine learning mode
  - **×** Darwinian evolution mode
- ✓ uses AQ decision rules as the classification model

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#### LEM: Evolutionary process guided by machine learning

- ✓ created by Ryszard Michalski [Mic00]
- ✓ originally, alternates between 2 phases:
  - **×** machine learning mode
  - **★** Darwinian evolution mode
- ✓ uses AQ decision rules as the classification model

LEM3: the most recent implementation of the LEM approach

- ✓ implemented by Janusz Wojtusiak [WM06, Woj07]
- ✓ alternates between more phases including local search, randomization, and representation adjustment
- ✓ good results reported, especially in the initial phases of the search

[Mic00] Ryszard S. Michalski. Learnable evolution model: Evolutionary processes guided by machine learning. *Machine Learning*, 38:9–40, 2000.

[WM06] Janusz Wojtusiak and Ryszard S. Michalski. The LEM3 system for non-darwinian evolutionary computation and its application to complex function optimization. Reports of the Machine Learning and Inference Laboratory MLI 04-1, George Mason University, Fairfax, VA, February 2006.

[Woj07] Janusz Wojtusiak. Handling Constrained Optimization Problems and Using Constructive Induction to Improve Representation Spaces in Learnable Evolution Model. Reports of the machine learning and inference laboratory, Fairfax, VA, November 2007.

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**×** because the assume all variables independent of each other

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- ✓ GAs are not very good for problems containing epistatic interactions among design variables
  - **×** because the assume all variables independent of each other
- ✓ There are methods that identify interactions and create the right linkage
  - ★ Perturbation techniques are the most straightforward
  - ★ Optimization by model fitting is a general approach

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  - ★ Optimization by model fitting is a general approach
- ✓ Using classification models is not so common
  - ★ LEM is a prominent example
  - ✗ another example for the real-valued spaces using elliptic classifiers will be given later in the course

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  - ★ Perturbation techniques are the most straightforward
  - **×** Optimization by model fitting is a general approach
- ✓ Using classification models is not so common
  - **×** LEM is a prominent example
  - ✗ another example for the real-valued spaces using elliptic classifiers will be given later in the course
- ✓ Using probabilistic models of the distribution of promising individuals is very popular:
  - ★ The class of estimation-of-distribution (EDA) algorithms
  - ★ EDAs are covered in the next 2 lectures