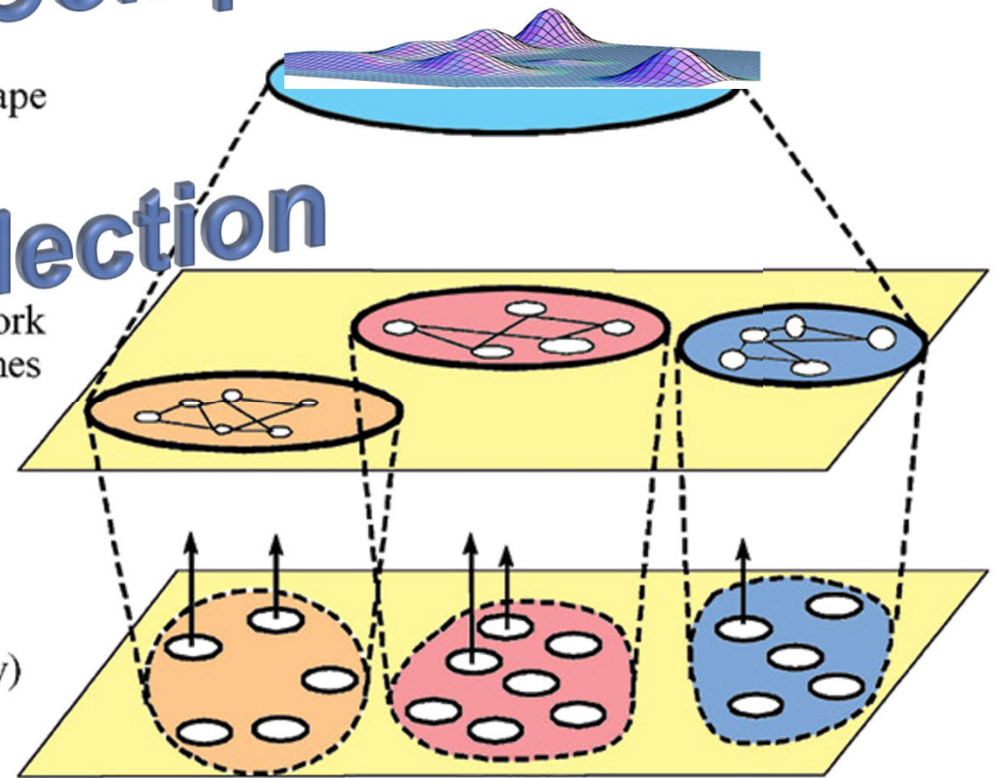


Adding Genomic Complexity And Multi-level Selection

Landscape

Patchwork
of regimes

Niches
(novelty)



key events coming up

- Labs: 35% (ISE-483)
 - Complete 5 (best 4 graded) assignments based on algorithms presented in class
 - Lab 4 : April 8th
 - Evolutionary Algorithms, (Assignment 4)
 - Delivered by SSIE583 Group 2
 - Due April 22nd
 - Lab 5: April 29th
 - Ant Clustering Algorithm, (Assignment 5)
 - Delivered by Group 1
 - Due May 6th
- SSIE – 583 -Presentation and Discussion: 25%
 - Present and lead the discussion of an article related to the class materials
 - Enginet students post/send video or join by Zoom
 - April 25th or April 29th
 - Conrad, M. [1990]. "The geometry of evolution." *Biosystems* 24: 61-81.
 - Mario Franco
 - Stanley, Kenneth O., Jeff Clune, Joel Lehman, and Risto Miikkulainen. "Designing Neural Networks through Neuroevolution." *Nature Machine Intelligence* 1, no. 1 (January 2019): 24–35.
 - Jessica Lasebikan
 - Lindgren, K. [1991]. "Evolutionary Phenomena in Simple Dynamics." In: *Artificial Life II*. Langton et al (Eds). Addison-wesley, pp. 295-312.
 - Akshay Gangadhar
 - Salahshour, Mohammad. "Interaction between Games Give Rise to the Evolution of Moral Norms of Cooperation." *PLOS Computational Biology* 18, no. 9 (September 29, 2022): e1010429
 - Srikanth Iyer
 - Discussion by all



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until now

■ Class Book

- Floreano, D. and C. Mattiussi [2008]. *Bio-Inspired Artificial Intelligence: Theories, Methods, and Technologies*. MIT Press.

- Chapter 7

■ Lecture notes

- Chapter 1: What is Life?
- Chapter 2: The logical Mechanisms of Life
- Chapter 3: Formalizing and Modeling the World
- Chapter 4: Self-Organization and Emergent Complex Behavior
- Chapter 5: Reality is Stranger than Fiction
- Chapter 6: Von Neumann and Natural Selection

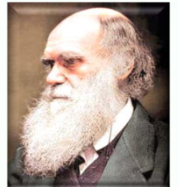
- posted online @ casci.binghamton.edu/academics/i-bic

■ Papers and other materials

- Optional

- Nunes de Castro, Leandro [2006]. *Fundamentals of Natural Computing: Basic Concepts, Algorithms, and Applications*. Chapman & Hall.

- Chapter 5, 7.7, 8.3.1, 8.3.6.



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■ Projects

- Due by **May 8th** in Brightspace, “Final Project Paper” assignment
 - ALIFE 2023
 - Not to submit to actual conference due date (April 3rd , 2024)
 - <https://2024.alife.org/>
 - 8 pages, author guidelines:
 - https://2024.alife.org/call_paper.html
 - MS Word and Latex/Overleaf templates
 - Preliminary ideas **by March 15**
 - Submit to “Project Idea” assignment in Brightspace.
- Individual or group
 - With very definite tasks assigned per member of group

ALIFE 2024

Tackle a real problem using bio-inspired algorithms, such as those used in the labs.

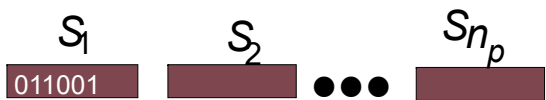


artificial genotype/phenotype mapping

Search algorithms based on the mechanics of Natural Selection
 Based on distinction between a machine and a description of a machine
 Solution alternatives for optimization problems

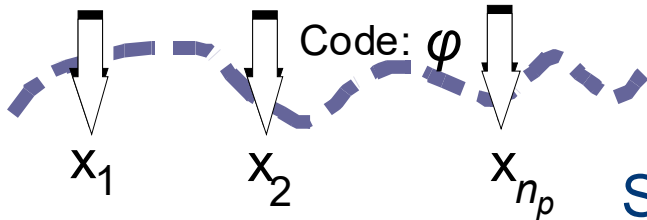
Traditional Genetic Algorithm

Genotype



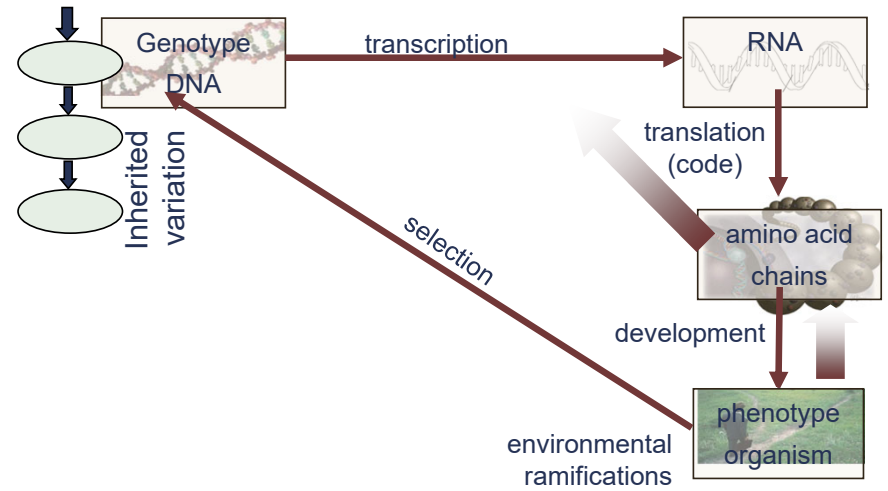
Variation

code



Selection

Phenotype

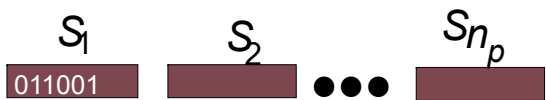


artificial genotype/phenotype mapping

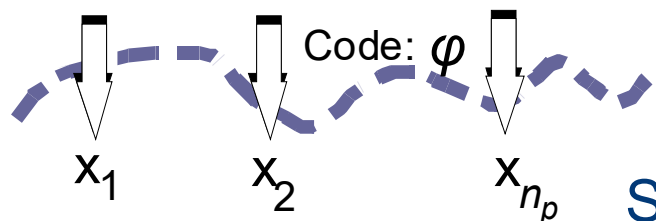
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Traditional Genetic Algorithm

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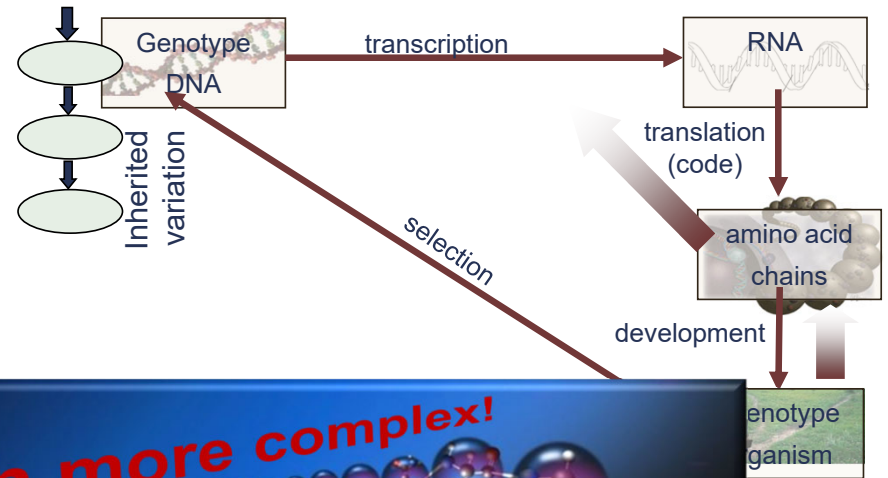
Variation



Selection

Phenotype

code



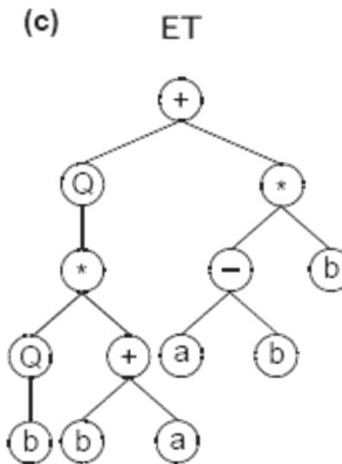
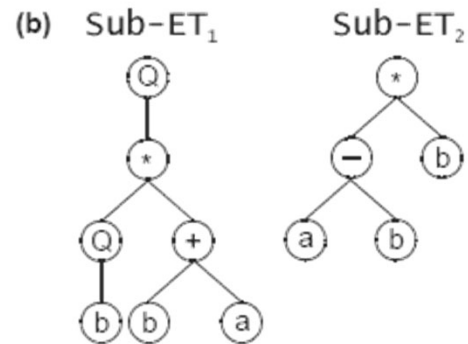
Including a genotype/phenotype map in GP

- Proposed by Candida Ferreira
- Program trees are encoded in fixed-length linear genotypes
- Genotypes
 - Open-reading frame architecture
 - Stop signal not necessarily at end of genotype
 - Non-coding genes are possible
 - Can include genetic operators
 - Genes contain two types of symbols
 - Functions (only at the head) and terminals
 - Multigenic solutions
 - Assembled from non-coding operations between various open-reading frames

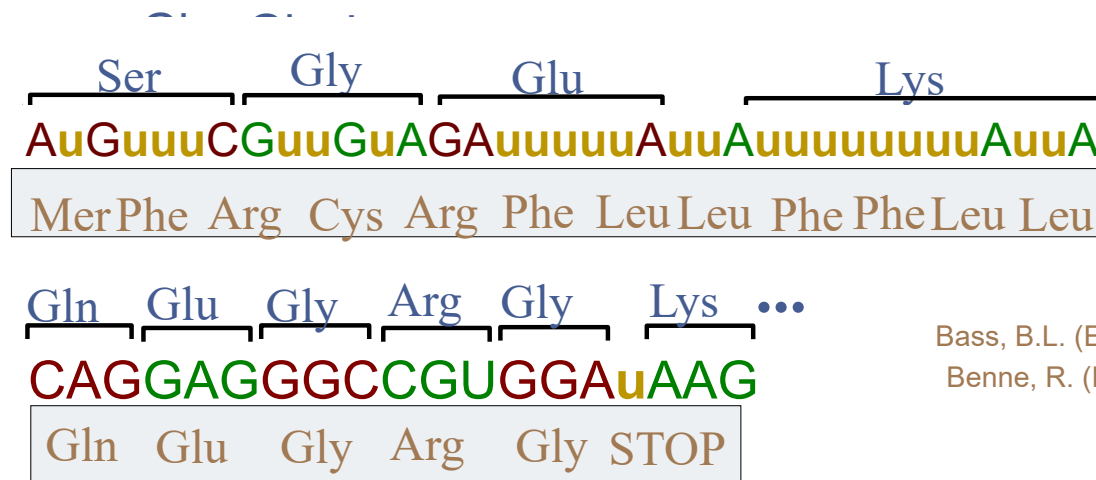
012345678012345678012345678
 -b*babbab*Qb+abbba-*Qabbaba

C. FERREIRA [2001]. Gene Expression Programming: A New Adaptive Algorithm for Solving Problems. *Complex Systems*, 13 (2): 87-129.

(a)
 012345678012345678
 Q*Q+bbaaa*-babaabb



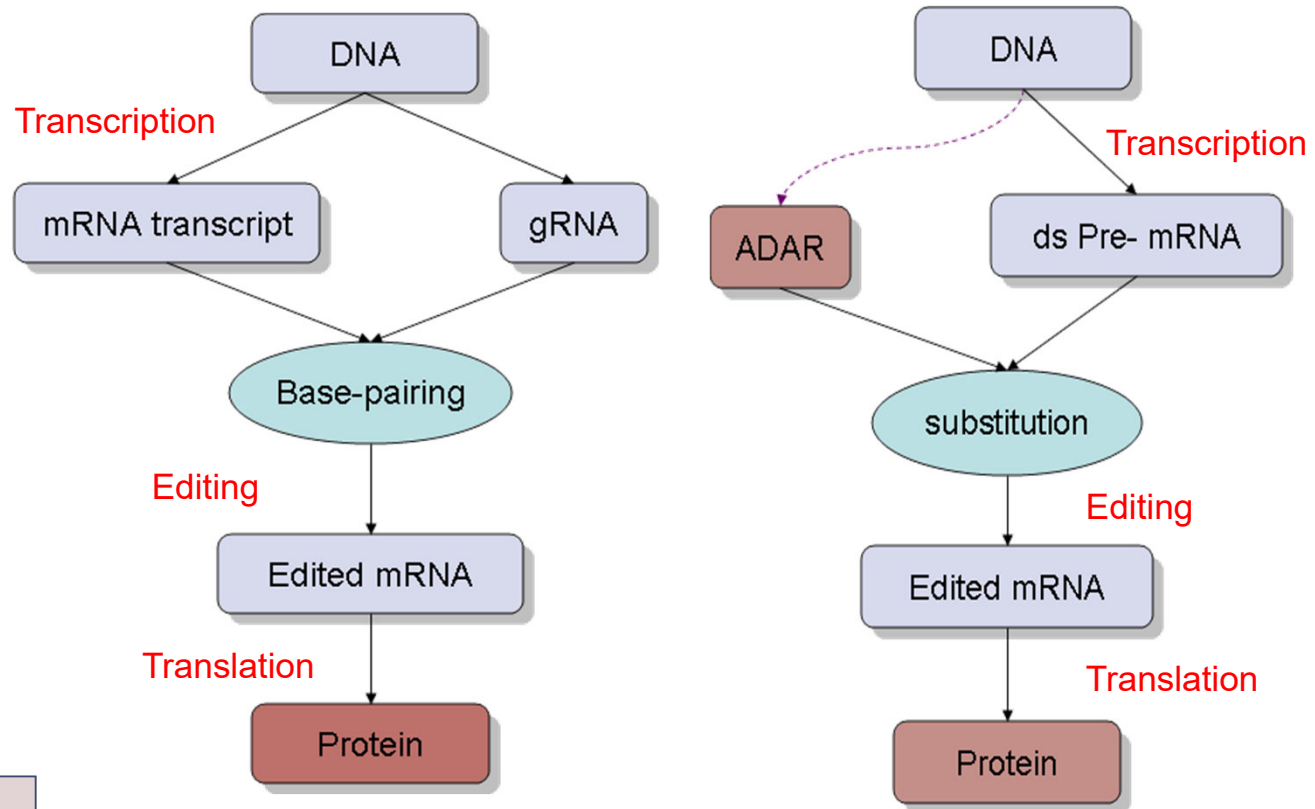
Including transcription regulation and translation



Bass, B.L. (Ed.) (2001). *RNA Editing*.
 Benne, R. (Ed.) (1993). *RNA Editing*

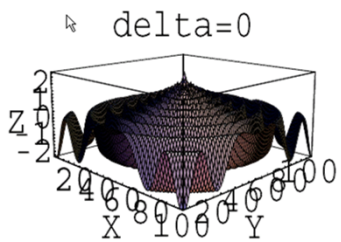
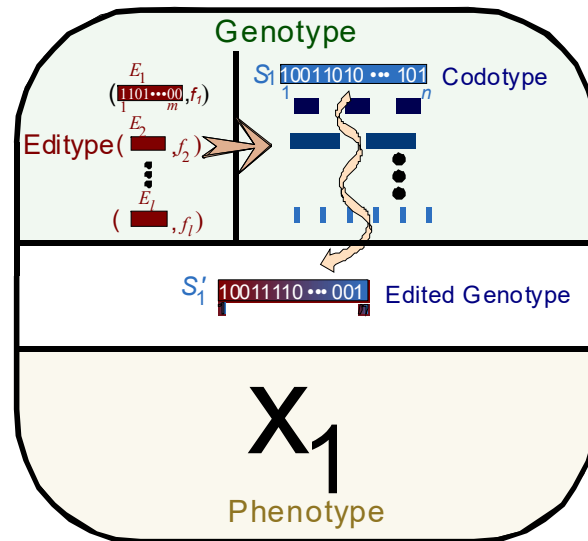
- RNA Editing: post-transcriptional alteration of genetic information
 - can be performed by ncRNA structures and proteins (i.e enzyme cascades).
- U-Insertion/deletion RNA Editing (mitochondria of kinetoplastid protozoa)
 - Involve small guide RNAs (gRNA) complementary to the target mRNA, and editosome (multi-protein complex)
 - gRNA is a template for editing
 - insertion/deletion of Uracil (U) residues, usually in coding (U) regions of mRNA transcripts
 - e.g. creation of open reading frames

U-insertion and A-to-I substitution



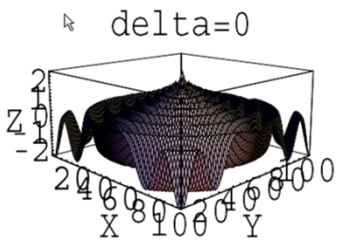
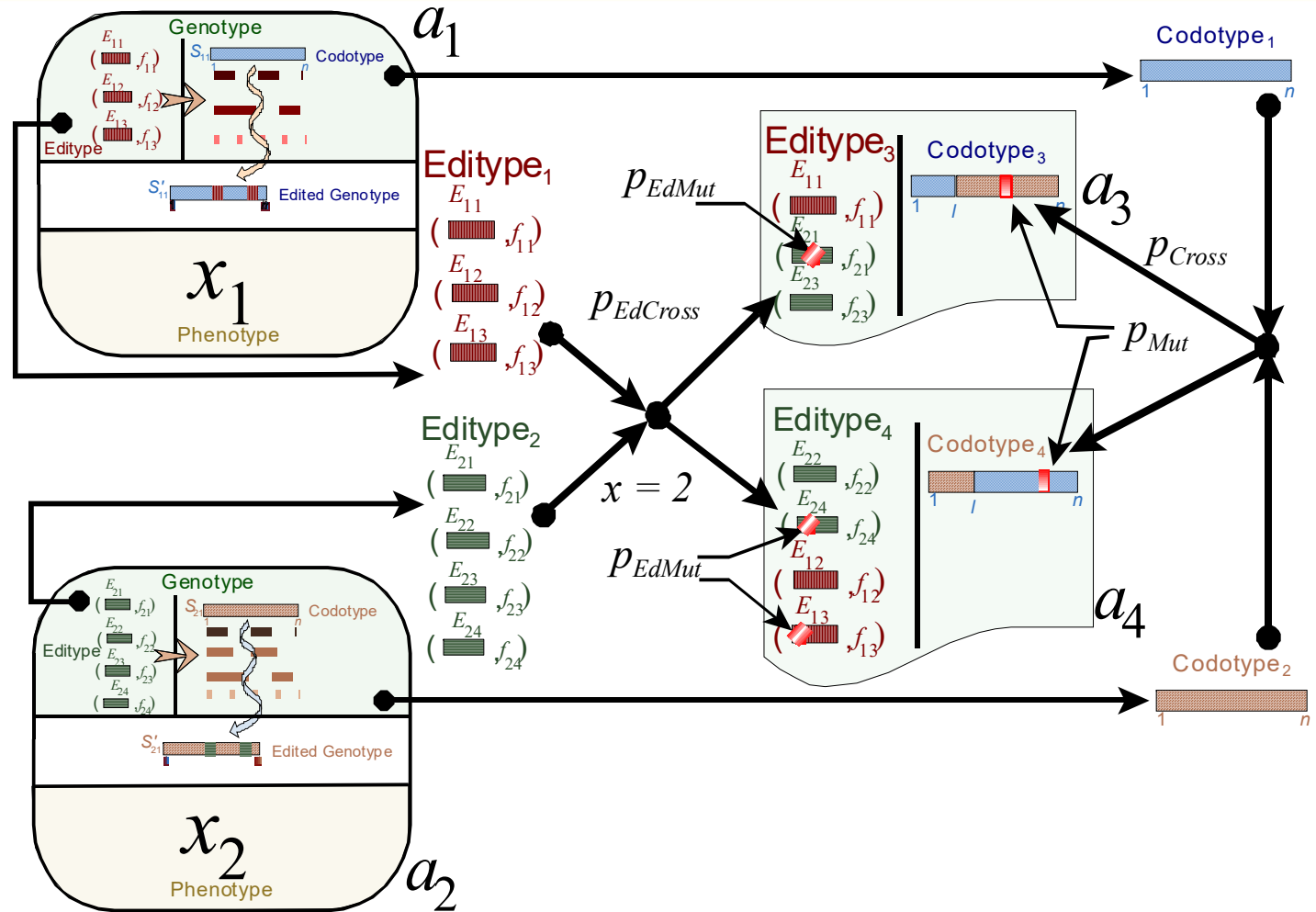
is there a **general principle** at play?

RNA Editing



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RNA Editing



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Simple fitness function

```
s1 = 11111*****; c1 = 10
s2 = *****11111*****; c2 = 10
s3 = *****11111*****; c3 = 10
s4 = *****11111*****; c4 = 10
s5 = *****11111*****; c5 = 10
s6 = *****11111*****; c6 = 10
s7 = *****11111*****; c7 = 10
s8 = *****11111*****; c8 = 10
```

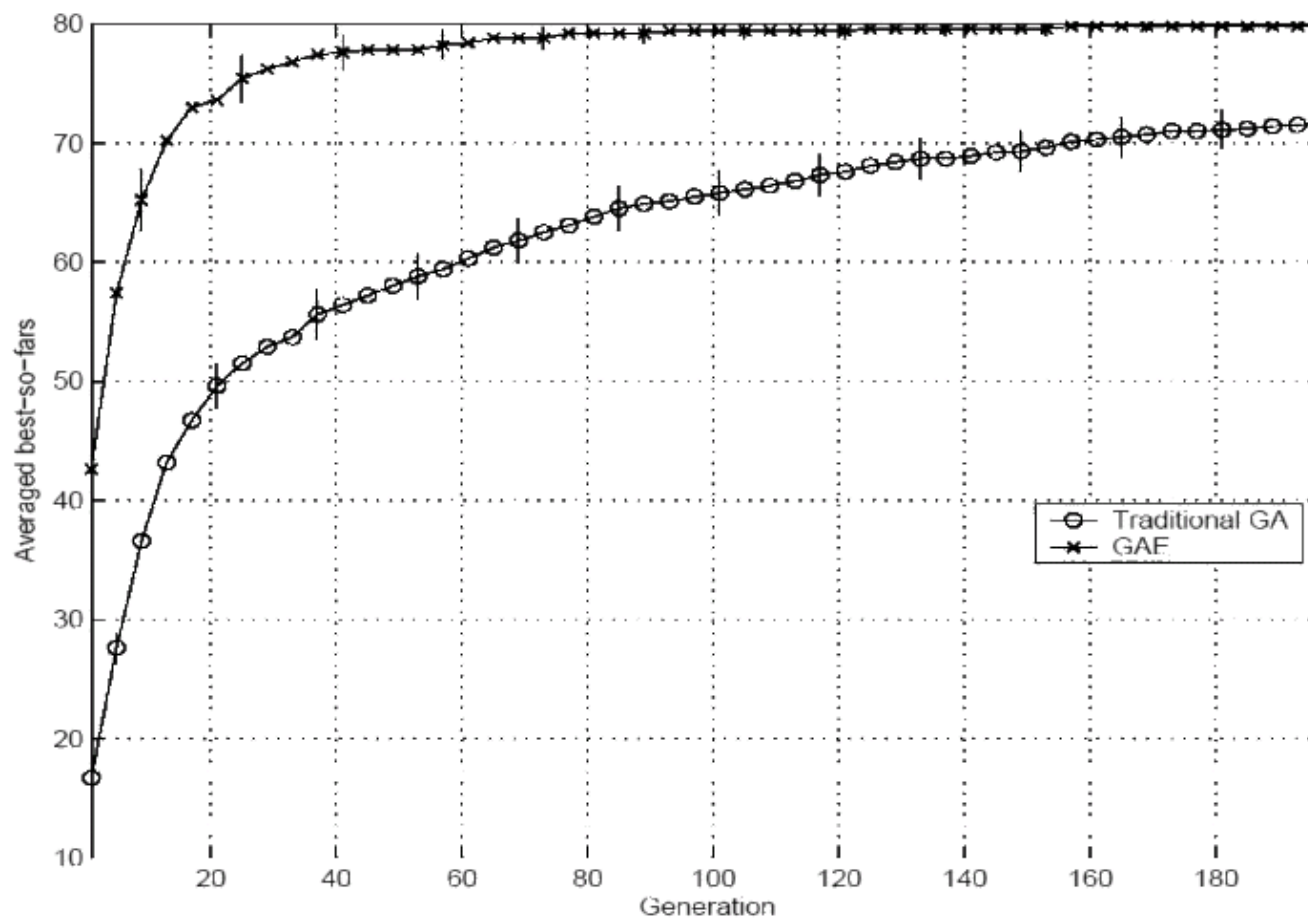
Royal road

$$F(x) = \sum_{s_i \in S} c_i \sigma_{s_i}(x)$$

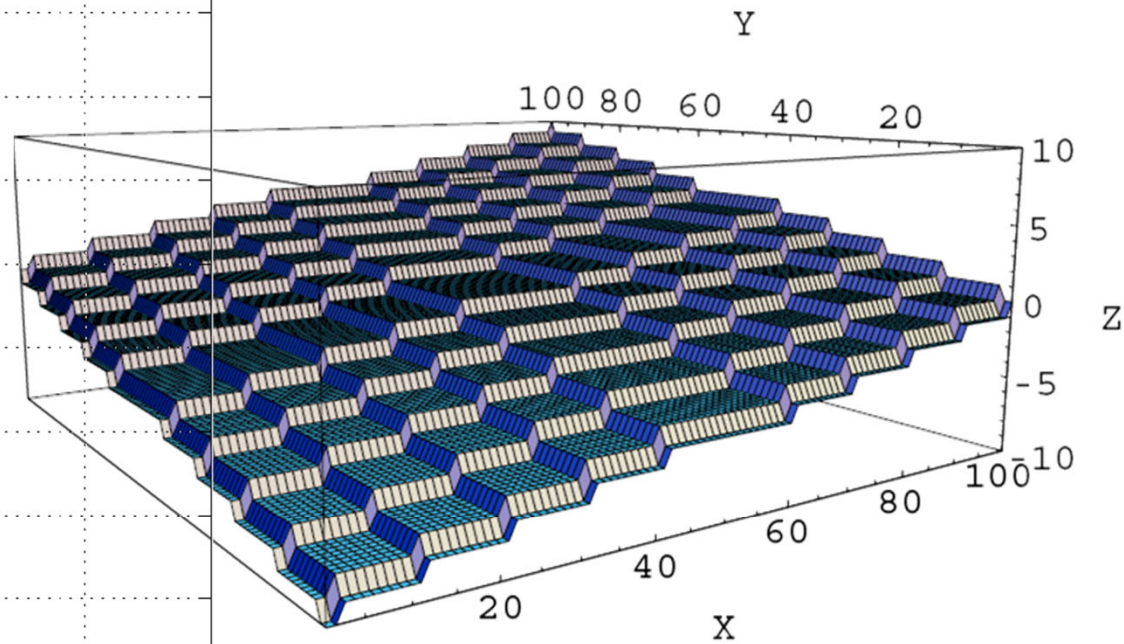
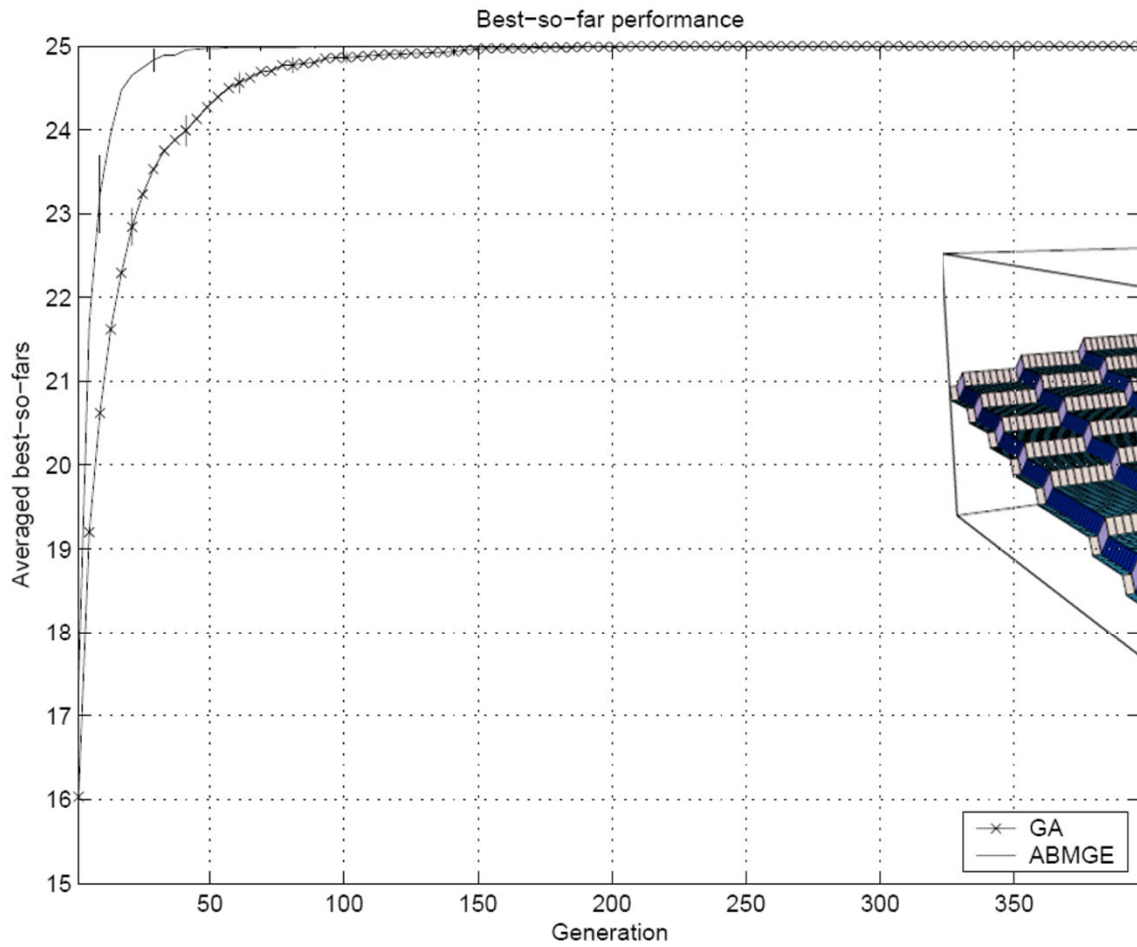
■ Miniature of "Royal Road" function (Forrest and Mitchell, 1993)

- ▶ Schemata $S = (s_1, \dots, s_8)$
- ▶ c_i is a value assigned to each schema s_i
- ▶ $\sigma_{s_i}(x) = 1$ if x is an instance of s_i and 0 otherwise
- ▶ Fitness of the global optimum string (40 1's) is $10 \cdot 8 = 80$

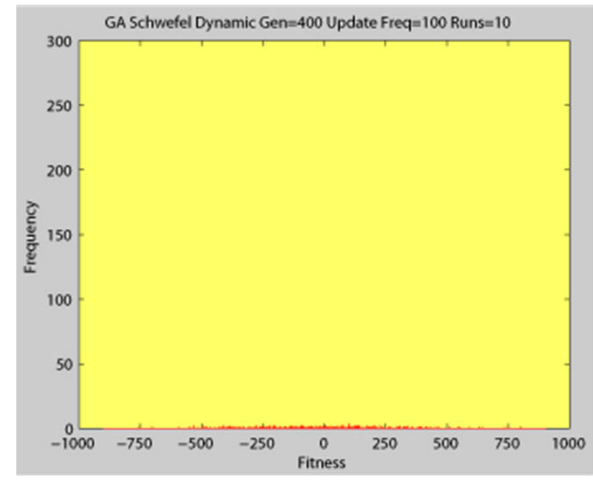
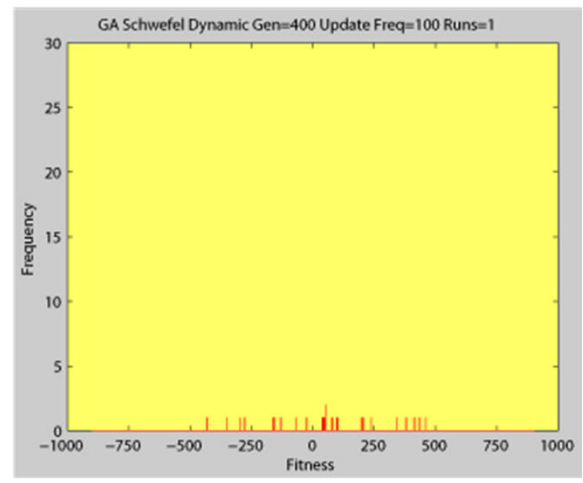
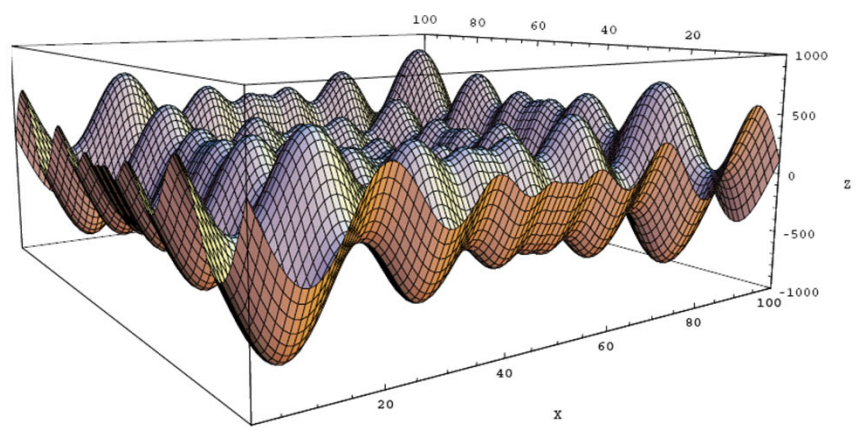
50 runs for small royal road testbed



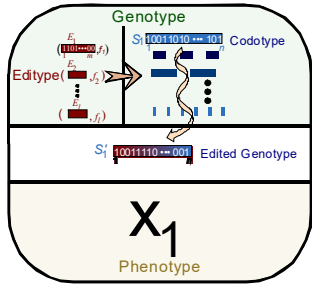
De Jong F3 function



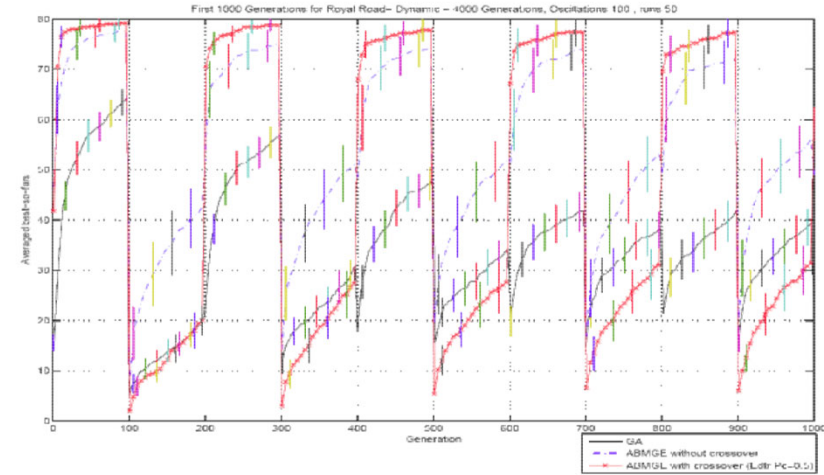
example run: Schwefel function



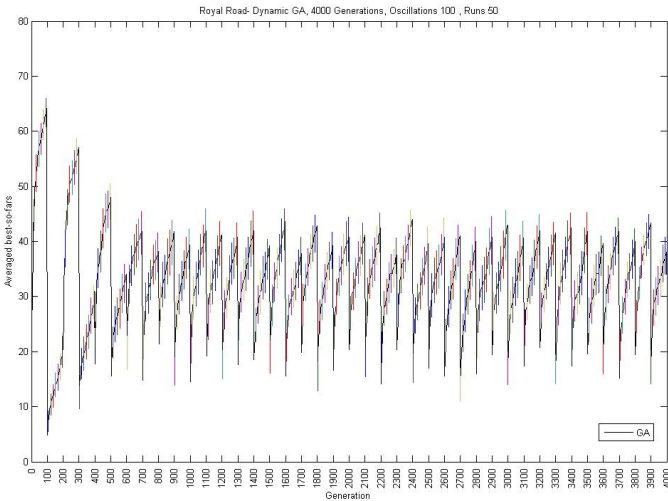
ABMGE on oscillatory fitness landscapes (Royal Road)



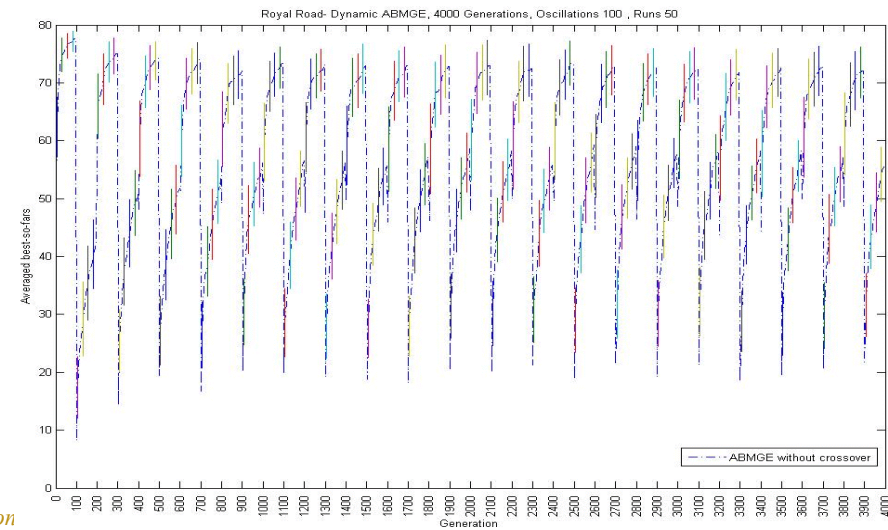
$s_1 = 11111$; $c_1 = 10$
 $s_2 = \text{*****}11111$; $c_2 = 10$
 $s_3 = \text{*****}11111$; $c_3 = 10$
 $s_4 = \text{*****}11111$; $c_4 = 10$
 $s_5 = \text{*****}11111$; $c_5 = 10$
 $s_6 = \text{*****}11111$; $c_6 = 10$
 $s_7 = \text{*****}11111$; $c_7 = 10$
 $s_8 = \text{*****}11111$; $c_8 = 10$



GA



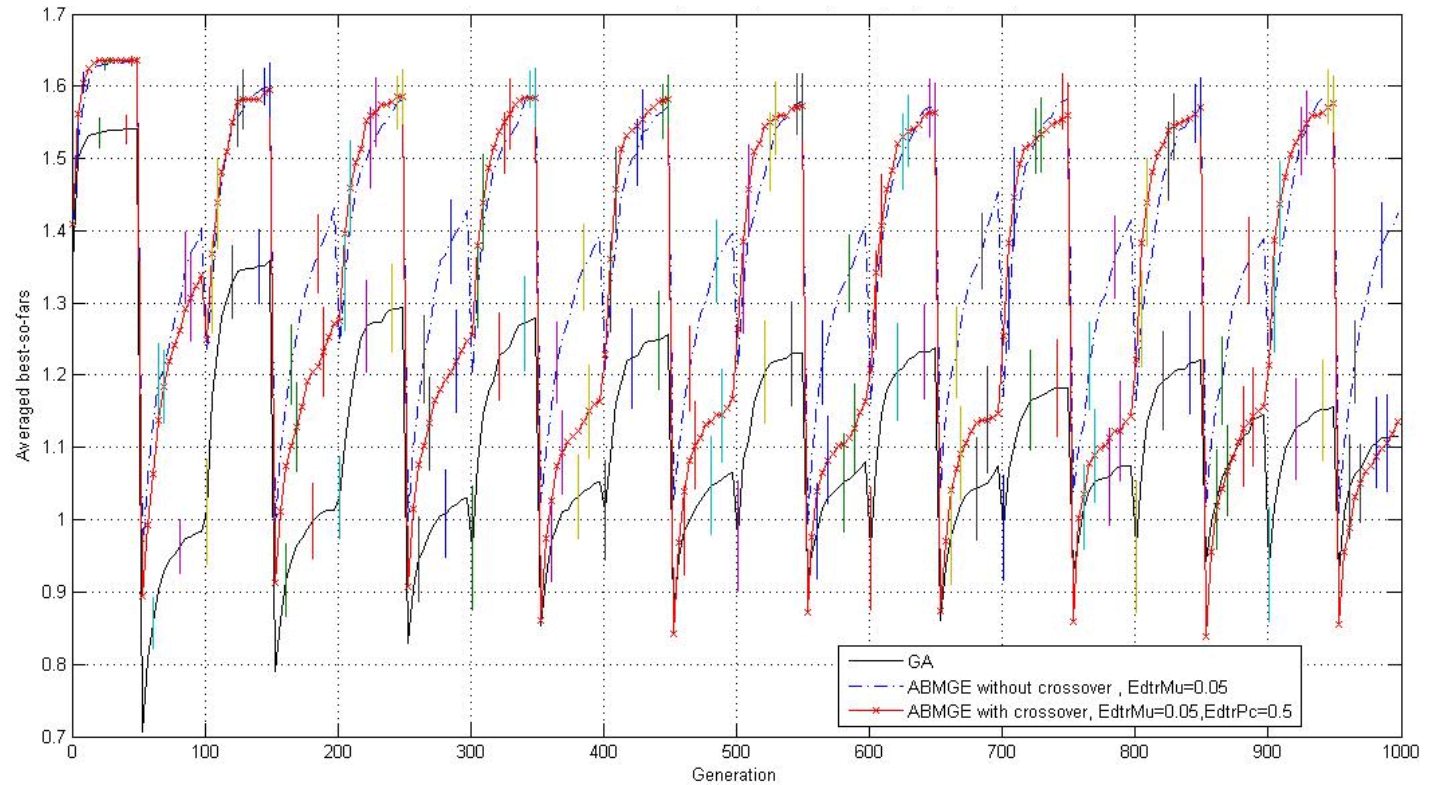
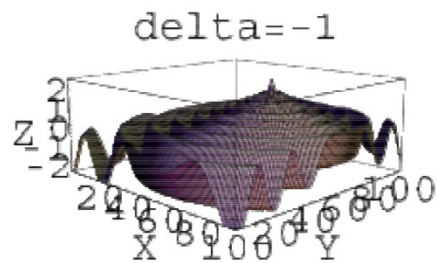
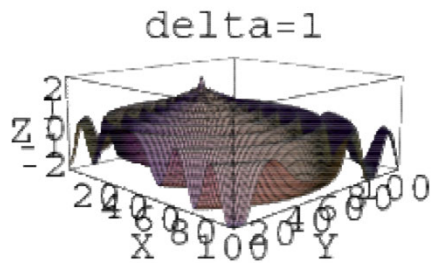
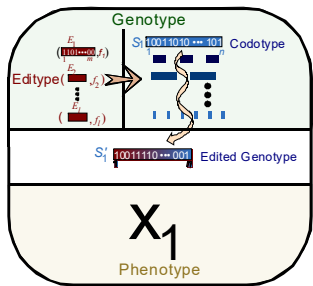
ABMGE



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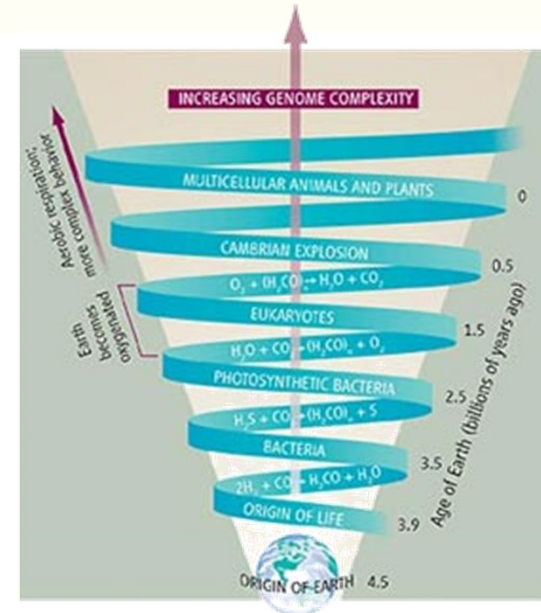
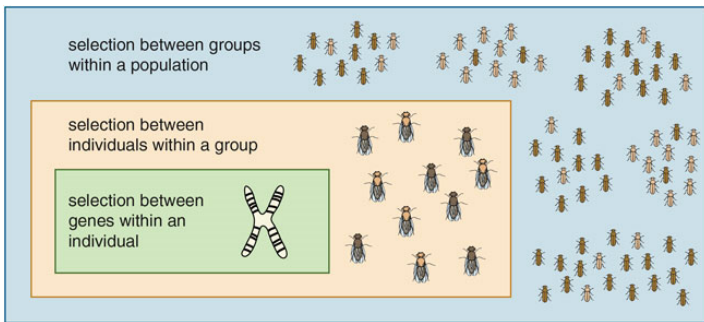
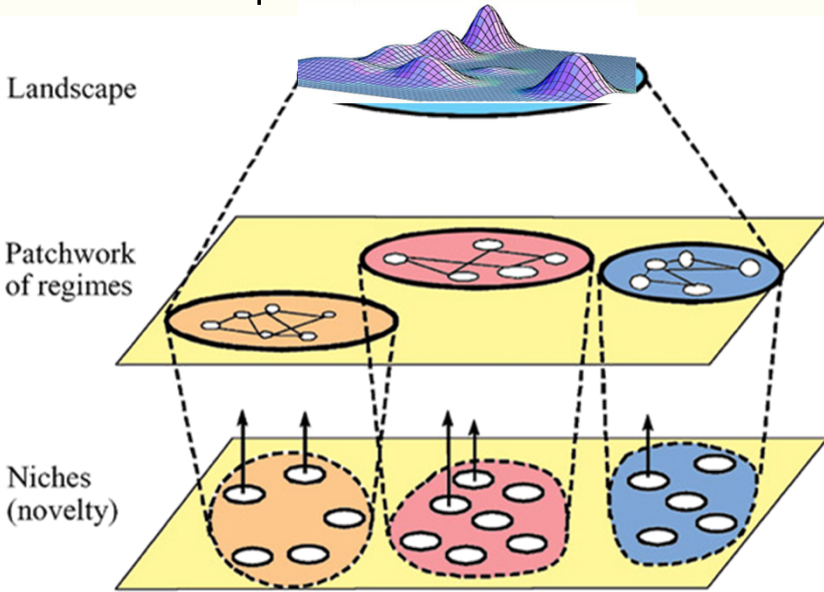
Huang, C-F, et al [2007]. *Evolutionary Computation*

ABMGE on dynamical environments



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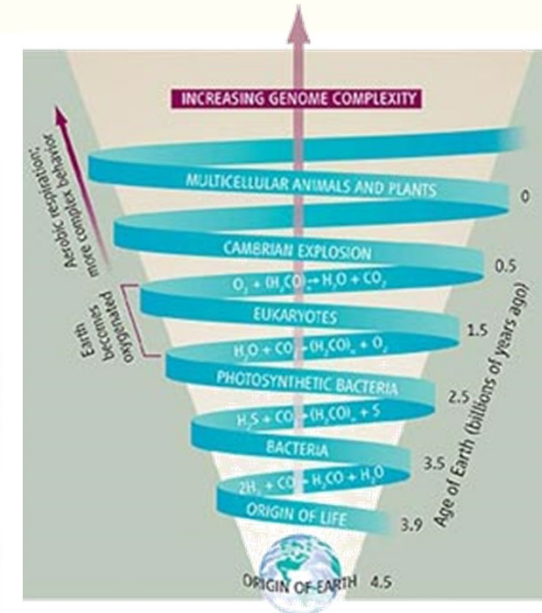
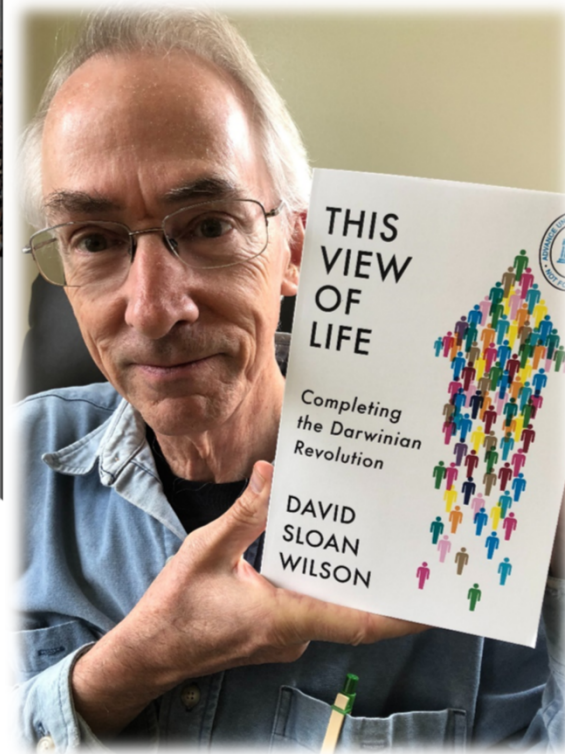
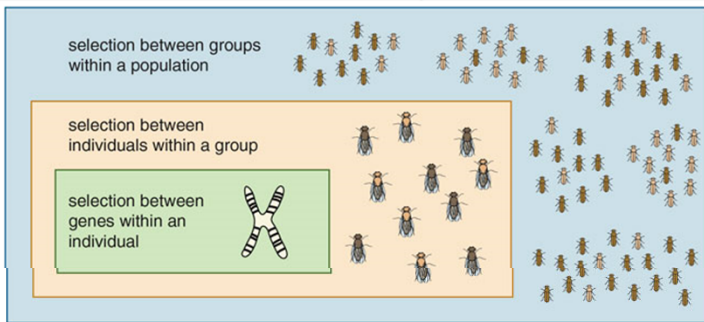
Genomic complexity and multi-level selection



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Wilson, David Sloan, and Edward O. Wilson. "Evolution 'for the Good of the Group' " *American Scientist* 96.5 (2008): 380-389.

Genomic complexity and multi-level selection



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altruism and selection

units of selection

moral men might not do any better than immoral men but tribes of moral men would certainly “have an immense advantage” over fractious bands of pirates. (Charles Darwin)

- **Multilevel selection theory**
 - Selection occurs in multiple levels simultaneously
 - No general-case scenario, each organism on a case-by-case basis
 - David Wilson and E.O. Wilson
- **Experiments with *Pseudomonas fluorescens***
 - Oxygen-exhausting bacteria in liquid
 - Groups with enough altruists survive
- **Kin-selection as special case of group selection**
 - Leading to various, diverse (selectable) groups with high genetic similarity
- **Sociobiology**
 - Selfishness beats altruism within groups. Altruistic groups beat selfish groups.

“Morality is herd instinct in the individual”.
(Friedrich Nietzsche)

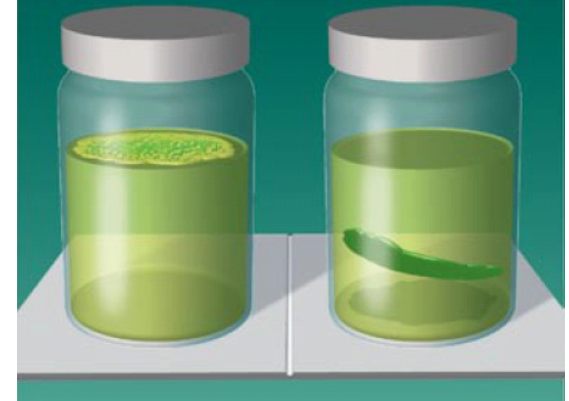
INDIVIDUAL SELECTION

“Altruist” *Pseudomonas fluorescens* bacteria (*below left*) carry a gene for secreting a polymer that enables mats of bacteria to float and thus access oxygen easily; “freeloaders” (*below right*) lack the gene. Producing the polymer costs extra energy, so freeloaders reproduce faster than altruists. Natural selection acting on individuals alone would drive the altruists to extinction.



GROUP SELECTION

But group selection appears to operate, too—at least in the laboratory—favoring mats of *P. fluorescens* bacteria in which some altruists persist. Only mats that include enough polymer-secreting altruists will float and thus survive to reproduce themselves, altruists included (*below left*). Mats in which individual selection leads to too many freeloaders will sink, drowning the entire bacterial colony (*below right*). Such mats leave no progeny.



Sci. American, Jan 2009 (Steve Mirsky)

altruism and selection

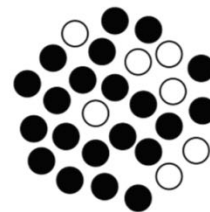
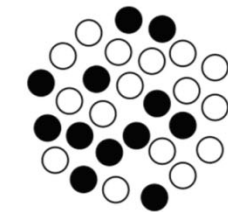
Waring, Timothy M., Michelle Kline Ann, Jeremy S. Brooks, Sandra H. Goff, John Gowdy, Marco A. Janssen, Paul E. Smaldino, and Jennifer Jacquet. "A multilevel evolutionary framework for sustainability analysis." *Ecology and Society* 20, no. 2 (2015).

units of select

moral men might
certainly "have an

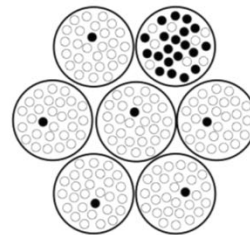
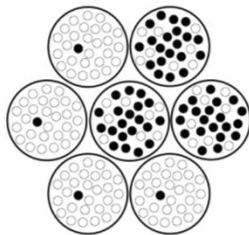
- **Multilevel**
 - Selection
 - No cases
- **Experimental**
 - Oxygen
 - Group
- **Kin-selection**
 - Learning
 - Gene
- **Sociobiology**
 - Selfish
 - Behavior

individual selection



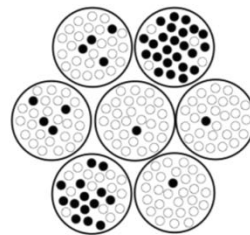
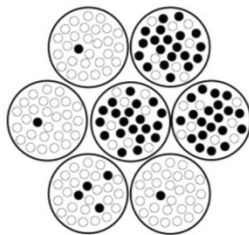
favors selfish individuals

group selection



favors cooperative individuals within selfish groups

multilevel selection

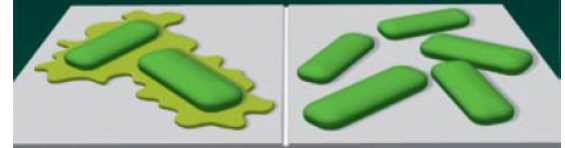


outcomes depend on the balance of both effects

○ cooperative ● non-cooperative ('selfish')

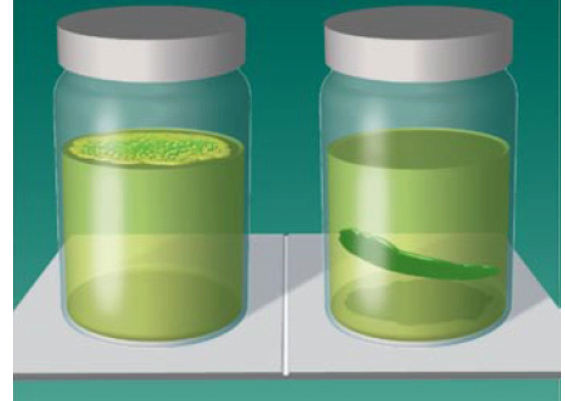
INDIVIDUAL SELECTION

"Altruist" *Pseudomonas fluorescens* bacteria (below left) carry a gene for secreting a polymer that enables mats of bacteria to float and thus access oxygen easily; "freeloaders" (below right) lack the gene. Producing the polymer costs extra energy, so freeloaders reproduce faster than altruists. Natural selection acting on individuals alone would drive the altruists to extinction.



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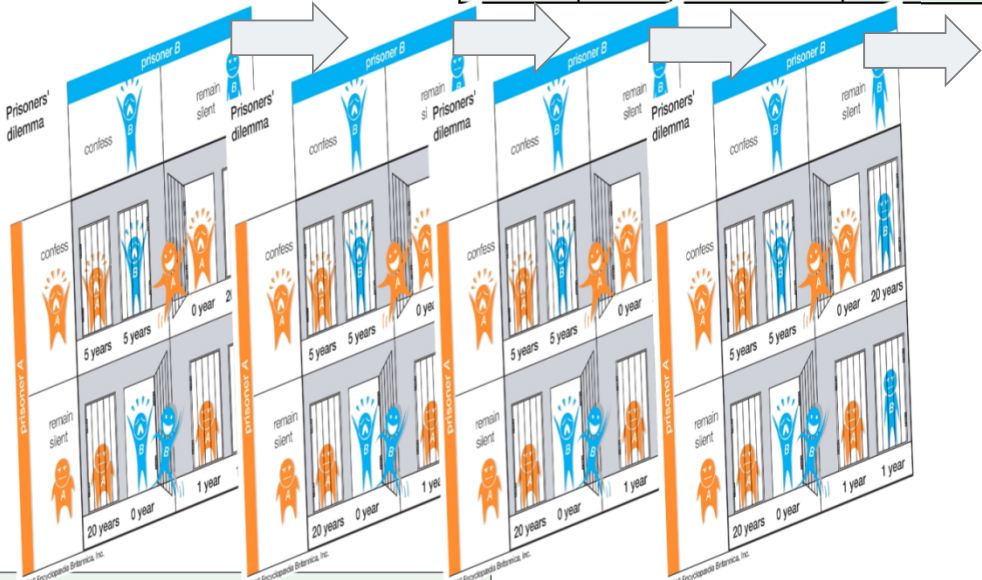
Sci. American, Jan 2009 (Steve Mirsky)

"Morality
(Friedrich Nietzsche)

iterated prisoner's dilemma

encoding

		P2	
		C	D
P1	C	(3,3)	(0,5)
	D	(5,0)	(1,1)



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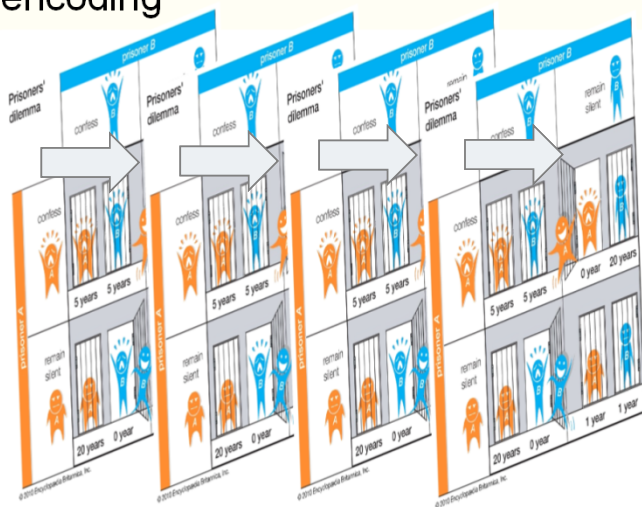
Prisoners' dilemma

		prisoner B	
		confess 	remain silent
prisoner A	confess 	 5 years	 5 years
	remain silent 	 20 years	 1 year

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iterated prisoner's dilemma

encoding

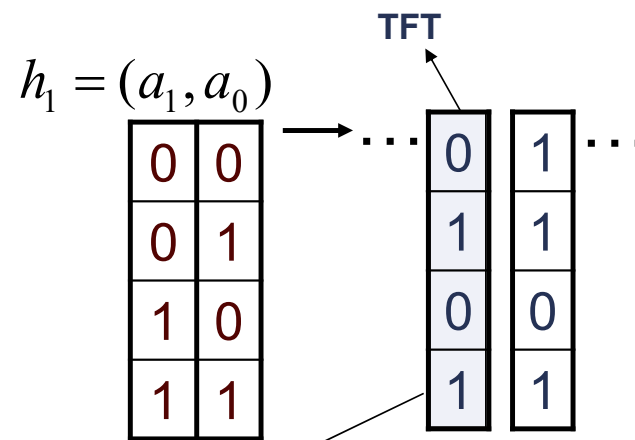


		P2	
		C	D
P1	C	(3,3)	(0,5)
	D	(5,0)	(1,1)

agent opponent

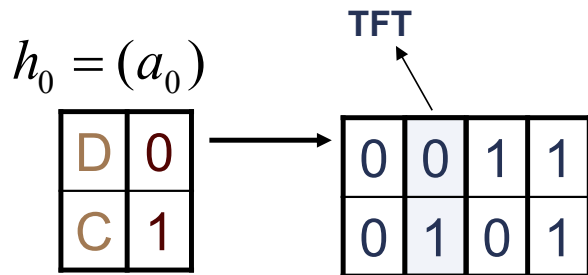
$$h_m = (a_{m-1}, \dots, a_1, a_0)$$

Lindgren's iterated game for agents with memory



Used in the evolutionary search by GA (tournament selection)

16 possible strategies (genotype=4 bits)



4 possible strategies (genotype=2 bits)

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memory 0 strategies

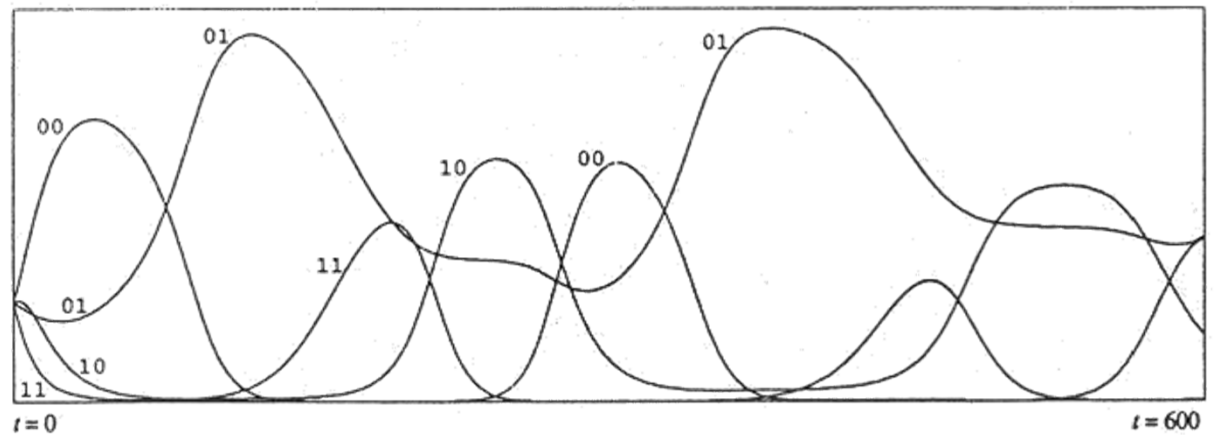
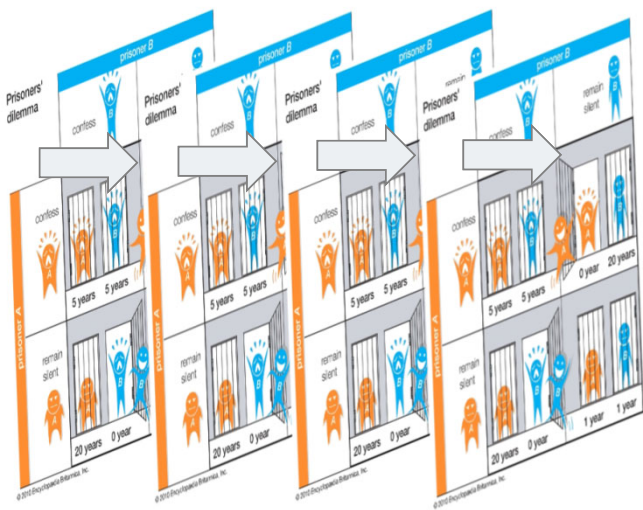
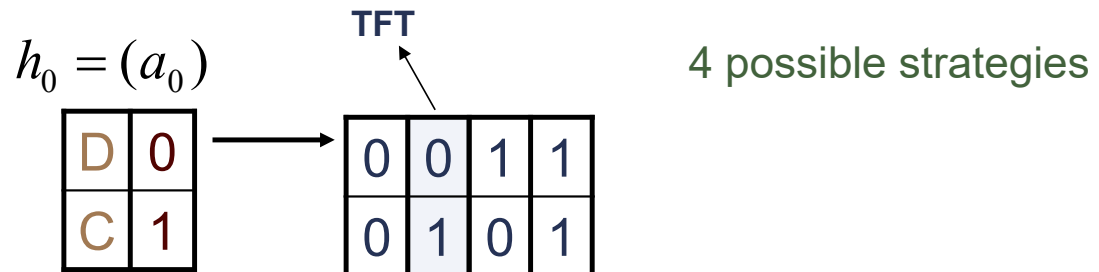


FIGURE 1 The evolution of a population of strategies starting with equal fractions of the memory one strategies [00], [01], [10], and [11] is shown for the first 600 generations. The fractions of different strategies are shown as functions of time (generation).

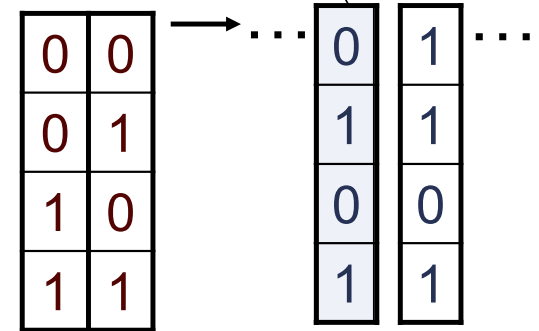


higher memory rules

16 possible strategies (genotype=4 bits)

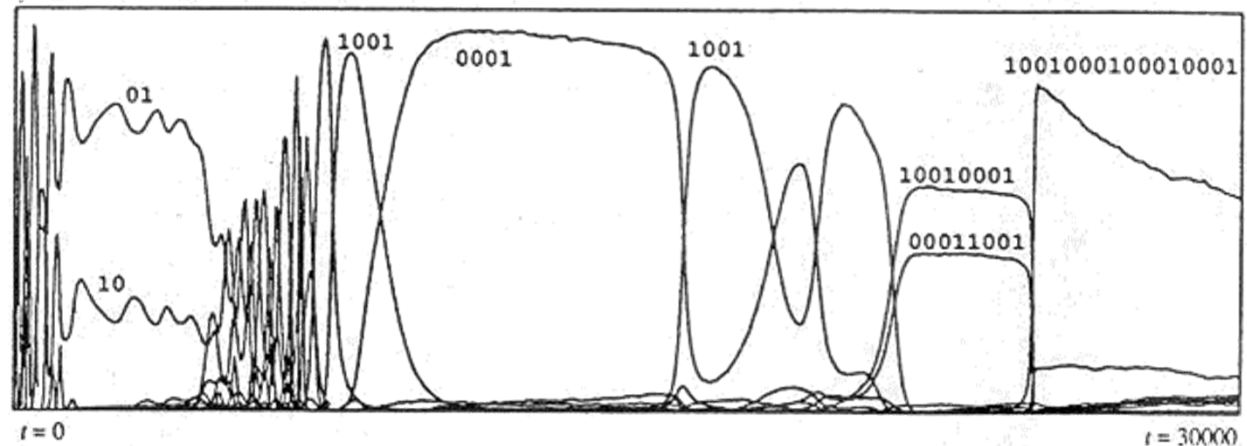
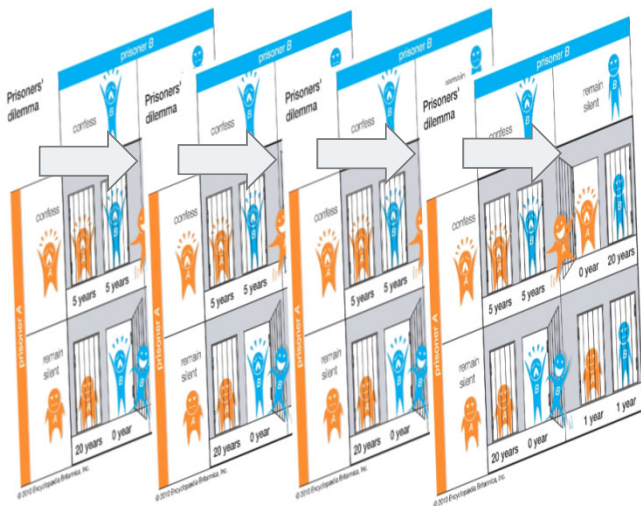
$$h_1 = (a_1, a_0)$$

TFT



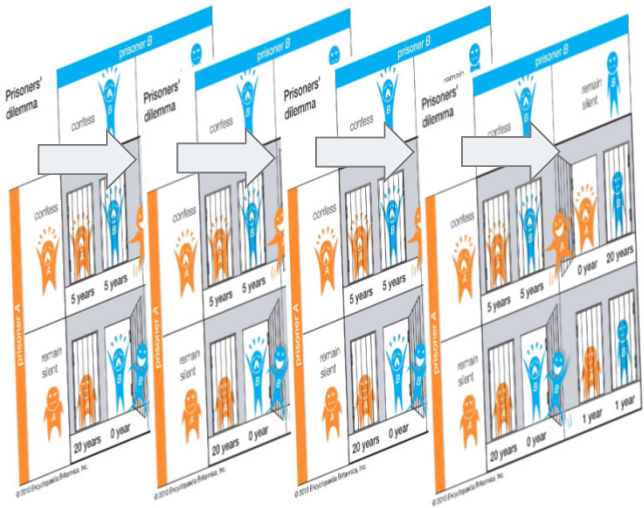
Used in the evolutionary search by GA (tournament selection)

GA uses variable length genotype



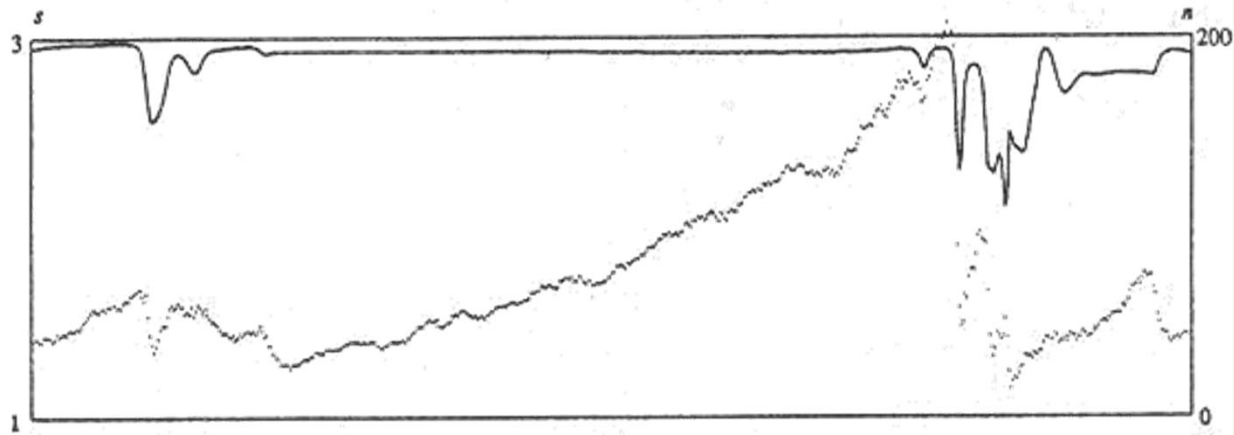
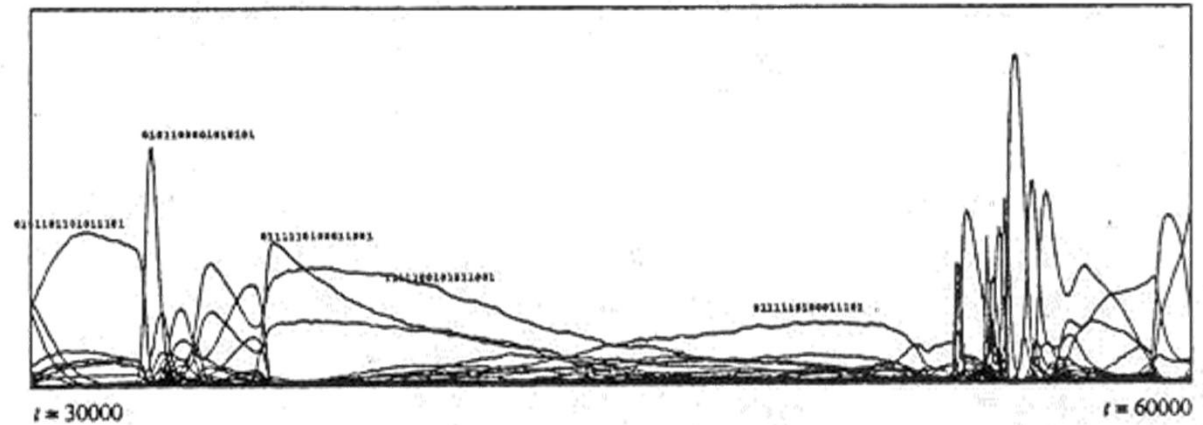
rocha@binghamton.edu
casci.binghamton.edu/academics/i-bic

Lindgren, Kristian. "Evolutionary phenomena in simple dynamics." *Artificial life II* (1991): 295-312.



Punctuated equilibria and complex evolutionary dynamics

(b)



readings

■ Class Book

- Floreano, D. and C. Mattiussi [2008]. *Bio-Inspired Artificial Intelligence: Theories, Methods, and Technologies*. MIT Press.
 - Chapter 7

■ Lecture notes

- Chapter 1: What is Life?
- Chapter 2: The logical Mechanisms of Life
- Chapter 3: Formalizing and Modeling the World
- Chapter 4: Self-Organization and Emergent Complex Behavior
- Chapter 5: Reality is Stranger than Fiction
- Chapter 6: Von Neumann and Natural Selection
 - posted online @ casci.binghamton.edu/academics/i-bic

■ Papers and other materials

● Optional

- Nunes de Castro, Leandro [2006]. *Fundamentals of Natural Computing: Basic Concepts, Algorithms, and Applications*. Chapman & Hall.
 - Chapter 5, 7.7, 8.3.1, 8.3.6,

