

# B. Genetic Algorithms

# Genetic Algorithms

- Developed by John Holland in '60s
- Did not become popular until late '80s
- A simplified model of genetics and evolution by natural selection
- Most widely applied to optimization problems (maximize “fitness”)

# Assumptions

- Existence of fitness function to quantify merit of potential solutions
  - this “fitness” is what the GA will maximize
- A mapping from bit-strings to potential solutions
  - best if each possible string generates a legal potential solution
  - choice of mapping is important
  - can use strings over other finite alphabets

# Outline of Simplified GA

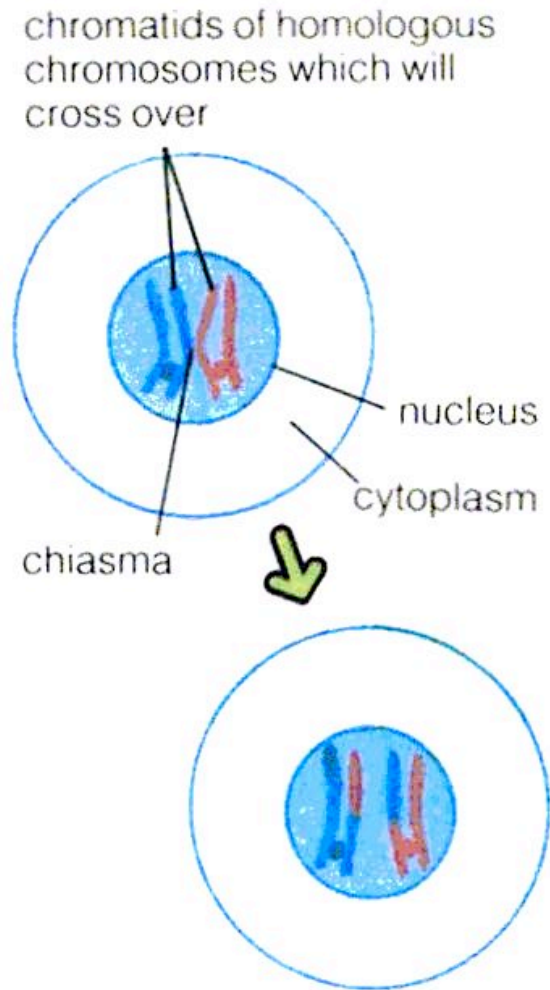
1. Random initial population  $P(0)$
2. Repeat for  $t = 0, \dots, t_{\max}$  or until converges:
  - a) create empty population  $P(t + 1)$
  - b) repeat until  $P(t + 1)$  is full:
    - 1) select two individuals from  $P(t)$  based on fitness
    - 2) optionally mate & replace with offspring
    - 3) optionally mutate offspring
    - 4) add two individuals to  $P(t + 1)$

# Fitness-Biased Selection

- Want the more “fit” to be more likely to reproduce
  - always selecting the best  
⇒ premature convergence
  - probabilistic selection ⇒ better exploration
- Roulette-wheel selection: probability  $\propto$  relative fitness:

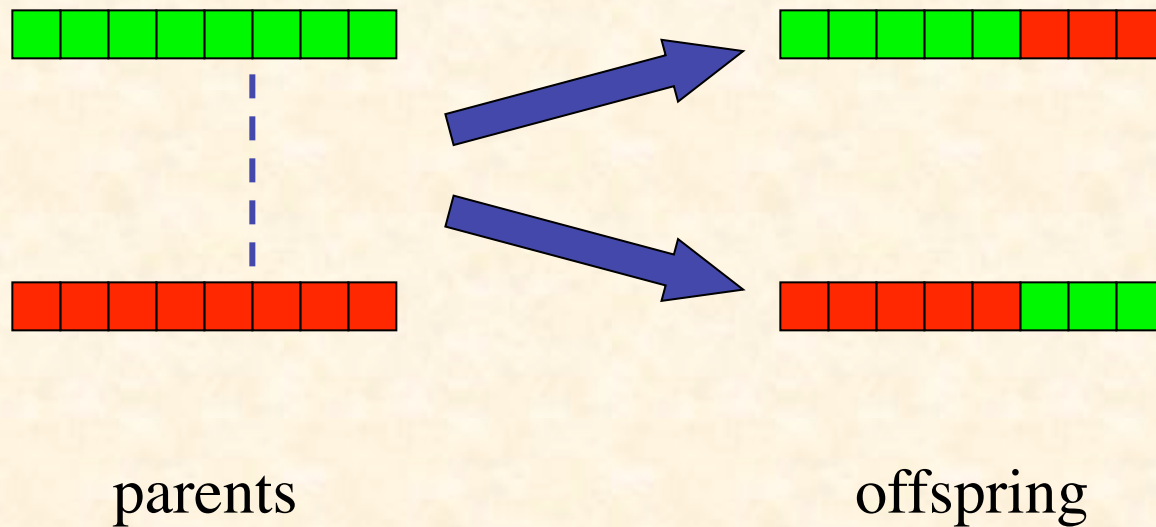
$$\Pr\{i \text{ mates}\} = \frac{f_i}{\sum_{j=1}^n f_j}$$

# Crossover: Biological Inspiration

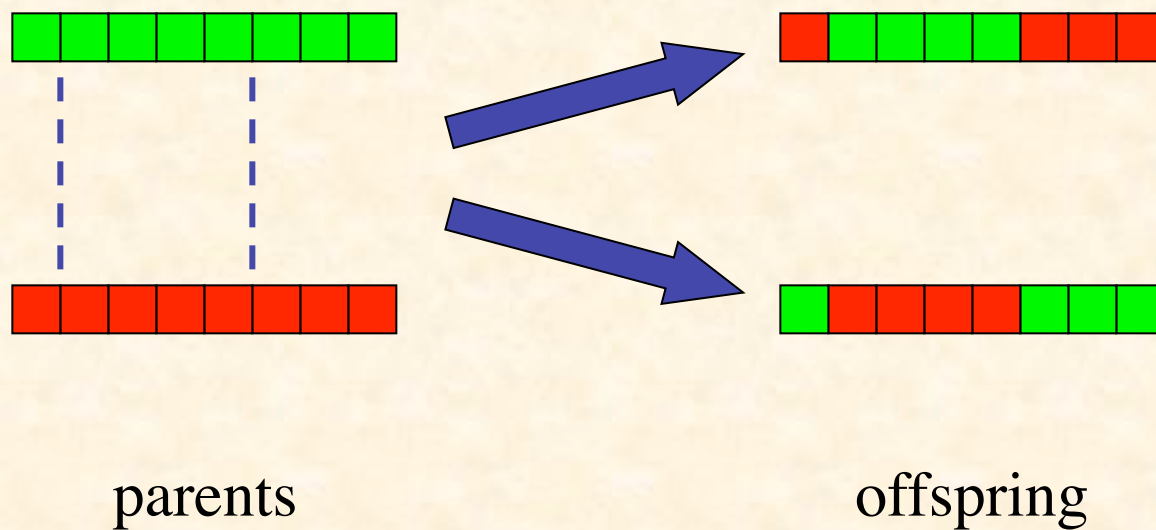


- Occurs during meiosis, when haploid gametes are formed
- Randomly mixes genes from two parents
- Creates genetic variation in gametes

# GAs: One-point Crossover

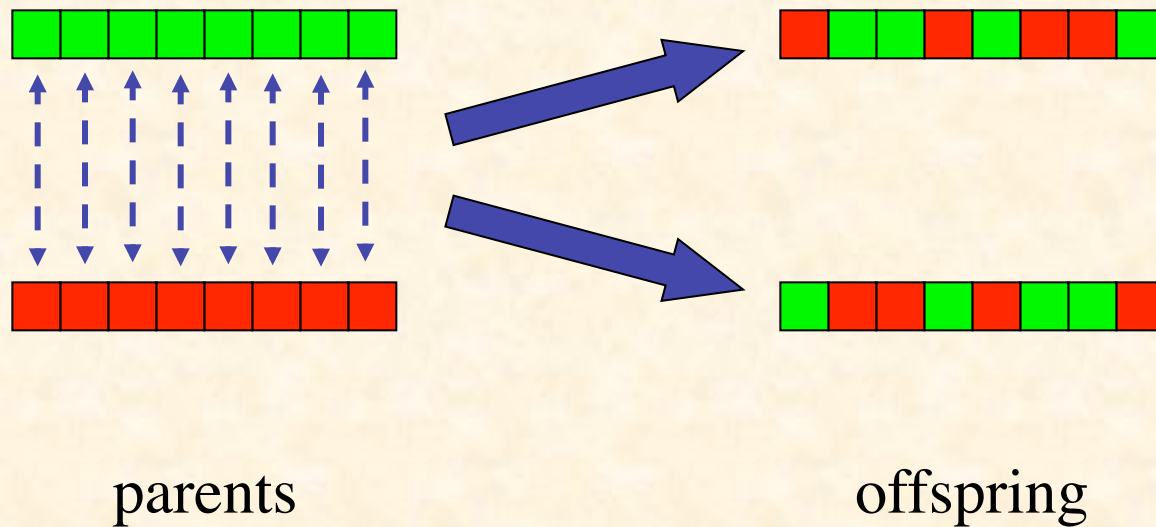


# GAs: Two-point Crossover



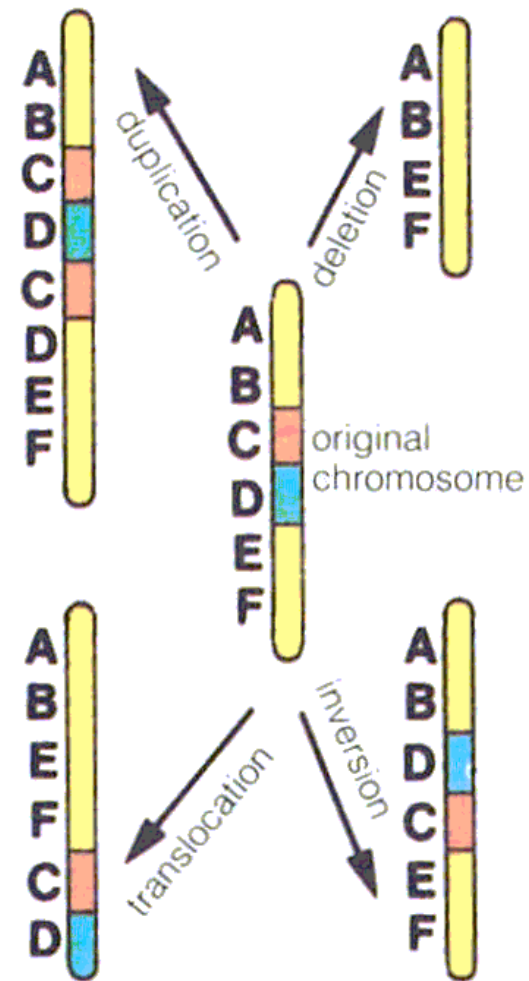


# GAs: $N$ -point Crossover



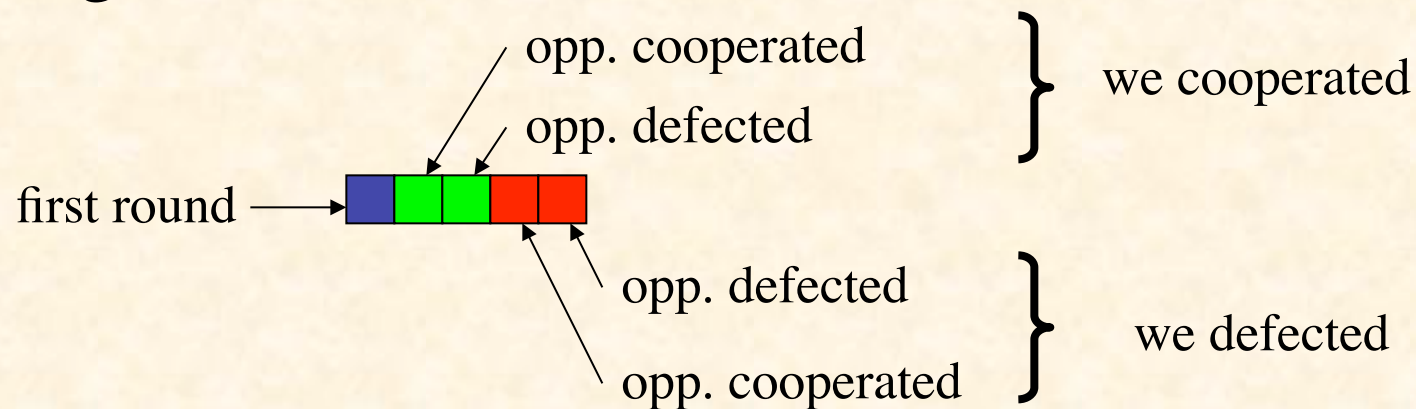
# Mutation: Biological Inspiration

- **Chromosome mutation**  $\Rightarrow$
- **Gene mutation**: alteration of the DNA in a gene
  - inspiration for mutation in GAs
- In typical GA each bit has a low probability of changing
- Some GAs models rearrange bits

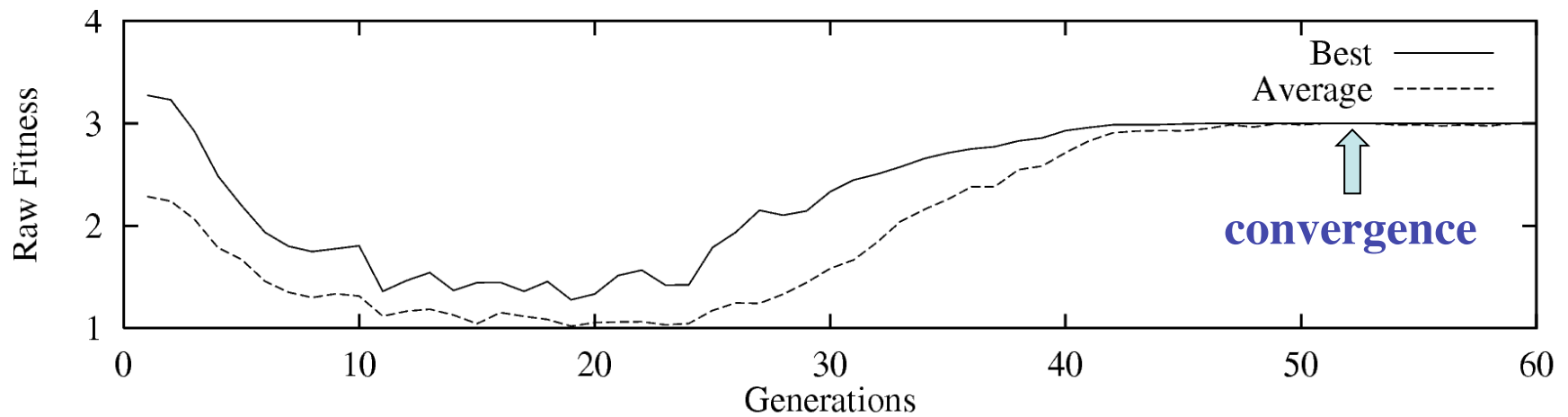


# Example: GA for IPD

- Genetic strings encode strategy
  - for first round
  - based on self's & opponent's action on  $r$  previous rounds
  - hence  $2^{2r} + 1$  bits
- E.g., for  $r = 1$ :



# Typical Result



**Figure 20.4** Average and best raw fitness scores for the IPD-playing GA

Figure from *The Computational Beauty of Nature: Computer Explorations of Fractals, Chaos, Complex Systems, and Adaptation*. Copyright © 1998–2000 by Gary William Flake. All rights reserved. Permission granted for educational, scholarly, and personal use provided that this notice remains intact and unaltered. No part of this work may be reproduced for commercial purposes without prior written permission from the MIT Press.

# The Red Queen Hypothesis



“Now, *here*, you see, it takes all the running *you* can do, to keep in the same place.”  
— *Through the Looking-Glass and What Alice Found There*

- *Observation*: a species probability of extinction is independent of time it has existed
- *Hypothesis*: species continually adapt to each other
- Extinction occurs with insufficient variability for further adaptation

# Reading

- Read Flake, ch. 17, “Competition & Cooperation”

# Demonstration of GA: Finding Maximum of Fitness Landscape

Run Genetic Algorithms — An Intuitive  
Introduction  
by Pascal Glauser

[www.glauserweb.ch/gentore.htm](http://www.glauserweb.ch/gentore.htm)

# Demonstration of GA: Evolving to Generate a Pre-specified Shape (Phenotype)

Run Genetic Algorithm Viewer

[www.rennard.org/alife/english/gavgb.html](http://www.rennard.org/alife/english/gavgb.html)



# Demonstration of GA: Eaters Seeking Food

<http://math.hws.edu/xJava/GA/>

# Morphology Project

by Michael “Flux” Chang

- Senior Independent Study project at UCLA
  - [users.design.ucla.edu/~mflux/morphology](http://users.design.ucla.edu/~mflux/morphology)
- Researched and programmed in 10 weeks
- Programmed in **Processing** language
  - [www.processing.org](http://www.processing.org)

# Genotype $\Rightarrow$ Phenotype

- Cells are “grown,” not specified individually
- Each gene specifies information such as:
  - angle
  - distance
  - type of cell
  - how many times to replicate
  - following gene
- Cells connected by “springs”
- Run **phenome**:  
[users.design.ucla.edu/~mflux/morphology/gallery/sketches/phenome](http://users.design.ucla.edu/~mflux/morphology/gallery/sketches/phenome)

# Complete Creature

- Neural nets for control (**blue**)
  - integrate-and-fire neurons
- Muscles (**red**)
  - decrease “spring length” when fire
- Sensors (**green**)
  - fire when exposed to “light”
- Structural elements (**grey**)
  - anchor other cells together
- Creature embedded in a fluid
- Run [users.design.ucla.edu/~mflux/morphology/gallery/sketches/creature](http://users.design.ucla.edu/~mflux/morphology/gallery/sketches/creature)

# Effects of Mutation

- Neural nets for control (**blue**)
- Muscles (**red**)
- Sensors (**green**)
- Structural elements (grey)
- Creature embedded in a fluid
- Run

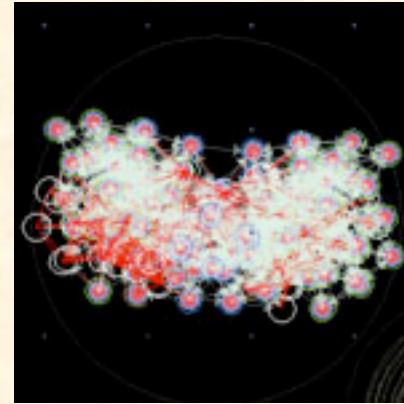
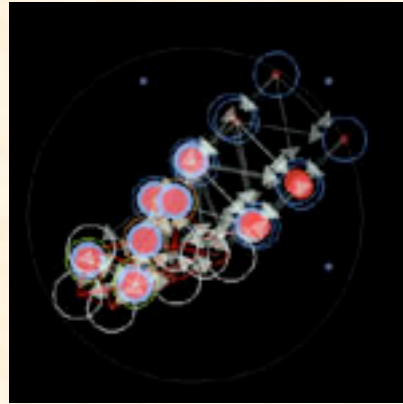
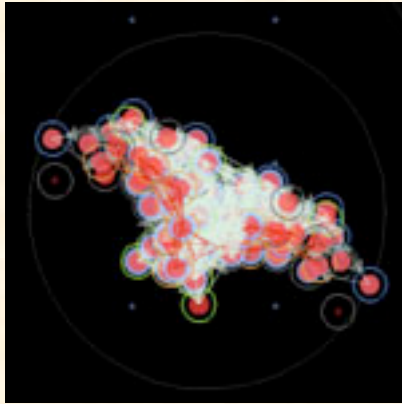
[users.design.ucla.edu/~mflux/morphology/gallery/sketches/creaturepack](http://users.design.ucla.edu/~mflux/morphology/gallery/sketches/creaturepack)

# Evolution

- Population: 150–200
- Nonviable & nonresponsive creatures eliminated
- Fitness based on speed or light-following
- 30% of new pop. are mutated copies of best
- 70% are random
- No crossover



# Gallery of Evolved Creatures



- Selected for speed of movement
- Run

[users.design.ucla.edu/~mflux/morphology/gallery/sketches/creaturegallery](http://users.design.ucla.edu/~mflux/morphology/gallery/sketches/creaturegallery)

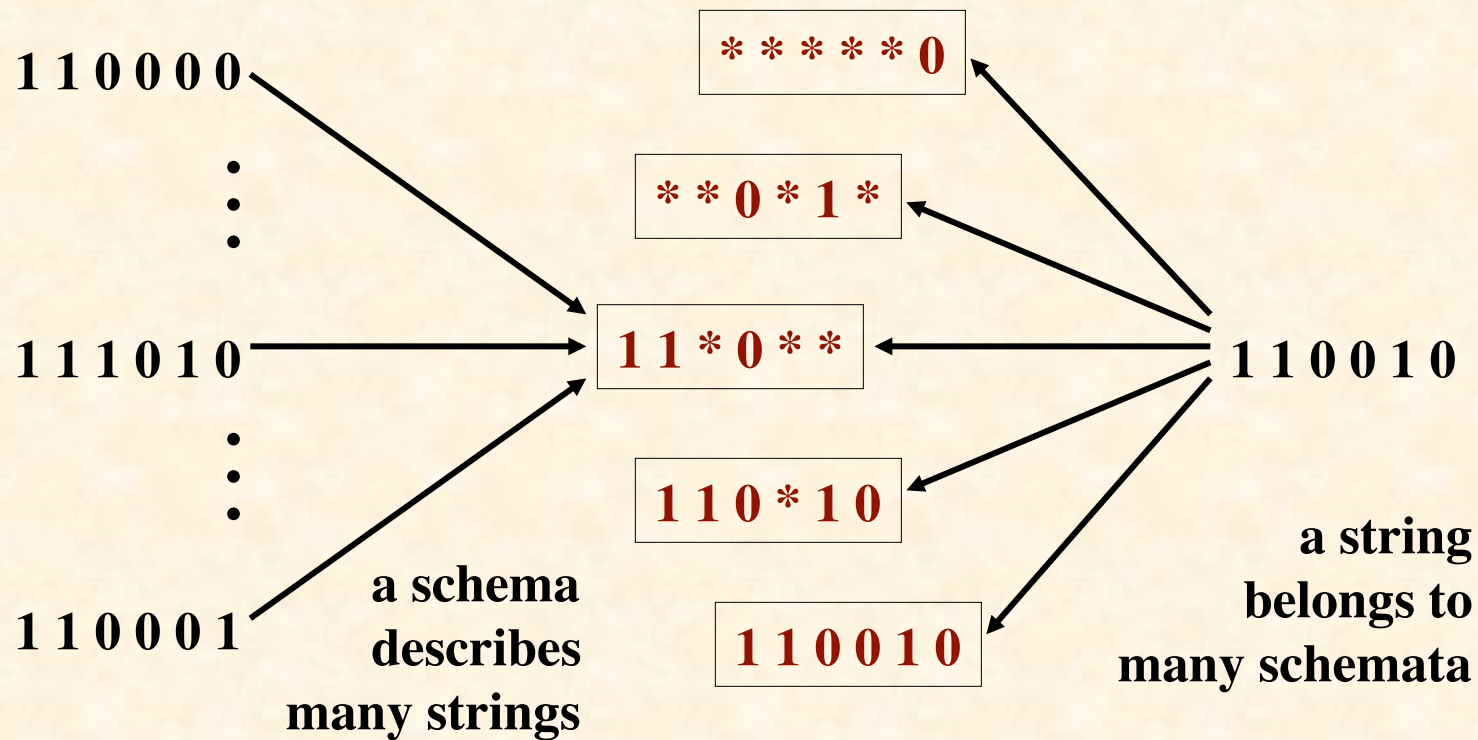
# Why Does the GA Work?

## The Schema Theorem



# Schemata

A **schema** is a description of certain patterns of bits in a genetic string



# The Fitness of Schemata

- The schemata are the **building blocks** of solutions
- We would like to know the average fitness of all possible strings belonging to a schema
- We cannot, but the strings in a population that belong to a schema give an estimate of the fitness of that schema
- Each string in a population is giving information about all the schemata to which it belongs (**implicit parallelism**)

# Effect of Selection

Let  $n$  = size of population

Let  $m(S, t)$  = number of instances of schema  $S$  at time  $t$

String  $i$  gets picked with probability  $\frac{f_i}{\sum_j f_j}$

Let  $f(S)$  = avg fitness of instances of  $S$  at time  $t$

So expected  $m(S, t + 1) = m(S, t) \cdot n \cdot \frac{f(S)}{\sum_j f_j}$

Since  $f_{\text{av}} = \frac{\sum_j f_j}{n}$ ,  $m(S, t + 1) = m(S, t) \frac{f(S)}{f_{\text{av}}}$

# Exponential Growth

- We have discovered:

$$m(S, t+1) = m(S, t) \cdot f(S) / f_{av}$$

- Suppose  $f(S) = f_{av} (1 + c)$
- Then  $m(S, t) = m(S, 0) (1 + c)^t$
- That is, **exponential growth** in above-average schemata

# Effect of Crossover

\*\*1 ... 0\*\*

|← $\delta$ →|

- Let  $\lambda$  = length of genetic strings
- Let  $\delta(S)$  = defining length of schema  $S$
- Probability {crossover destroys  $S$ }:

$$p_d \leq \delta(S) / (\lambda - 1)$$

- Let  $p_c$  = probability of crossover
- Probability schema survives:

$$p_s \geq 1 - p_c \frac{\delta(S)}{\lambda - 1}$$

# Selection & Crossover Together

$$m(S, t + 1) \geq m(S, t) \frac{f(S)}{f_{\text{av}}} \left[ 1 - p_c \frac{\delta(S)}{\lambda - 1} \right]$$

# Effect of Mutation

- Let  $p_m$  = probability of mutation
- So  $1 - p_m$  = probability an allele survives
- Let  $o(S)$  = number of fixed positions in  $S$
- The probability they all survive is  
 $(1 - p_m)^{o(S)}$
- If  $p_m \ll 1$ ,  $(1 - p_m)^{o(S)} \approx 1 - o(S) p_m$

# Schema Theorem: “Fundamental Theorem of GAs”

$$m(S, t + 1) \geq m(S, t) \frac{f(S)}{f_{av}} \left[ 1 - p_c \frac{\delta(S)}{\lambda - 1} - o(S) p_m \right]$$



# The Bandit Problem

- Two-armed bandit:
  - random payoffs with (unknown) means  $m_1, m_2$  and variances  $\sigma_1, \sigma_2$
  - optimal strategy: allocate exponentially greater number of trials to apparently better lever
- $k$ -armed bandit: similar analysis applies
- Analogous to allocation of population to schemata
- Suggests GA may allocate trials optimally

# Goldberg's Analysis of Competent & Efficient GAs

# Paradox of GAs

- Individually uninteresting operators:
  - selection, recombination, mutation
- Selection + mutation  $\Rightarrow$  continual improvement
- Selection + recombination  $\Rightarrow$  innovation
  - fundamental to invention:  
*generation vs. evaluation*
- Fundamental intuition of GAs: the three work well together

# Race Between Selection & Innovation: Takeover Time

- Takeover time  $t^*$  = average time for most fit to take over population
- Transaction selection: population replaced by  $s$  copies of top  $1/s$
- $s$  quantifies selective pressure
- Estimate  $t^* \approx \ln n / \ln s$

# Innovation Time

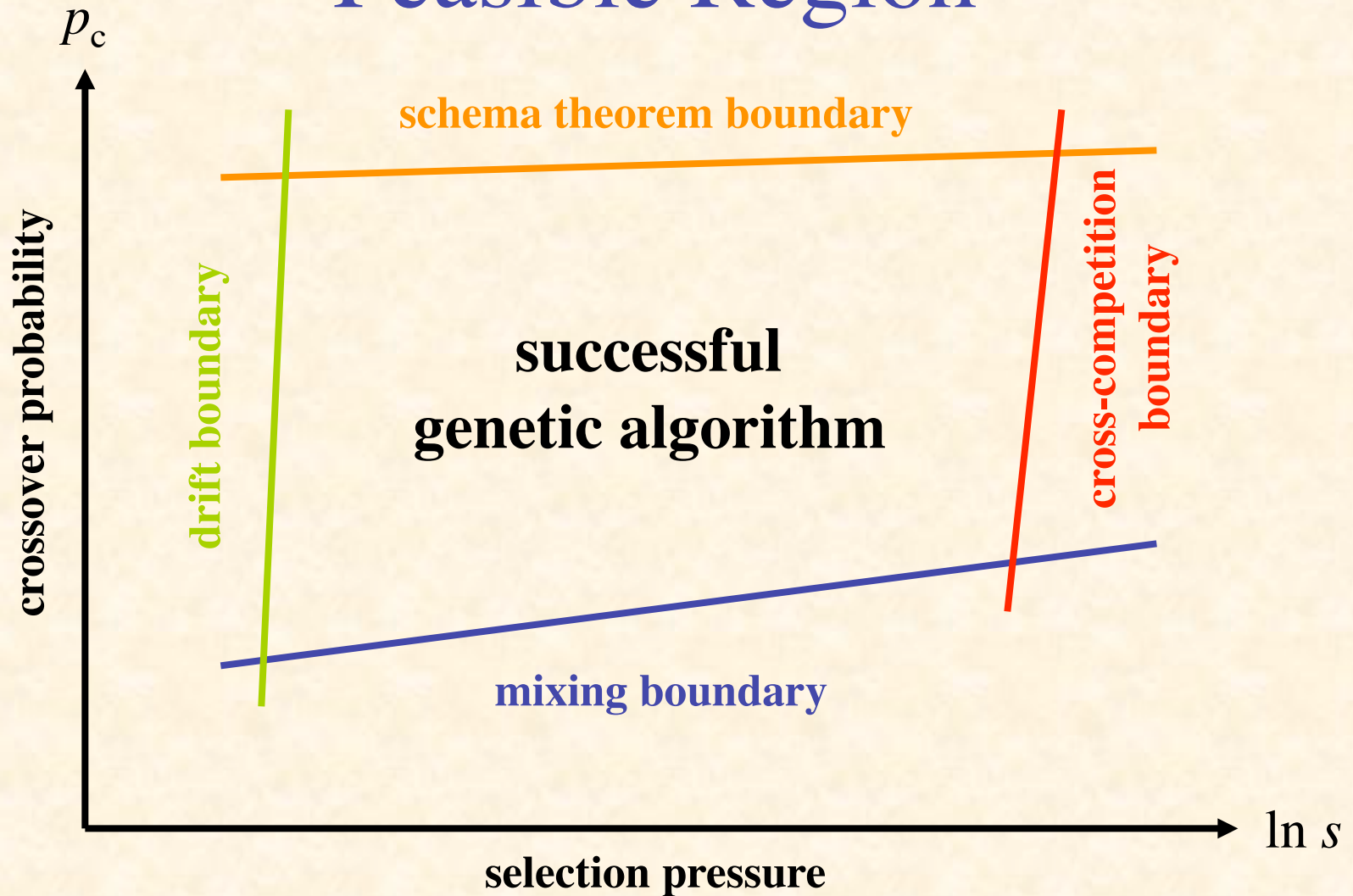
- Innovation time  $t_i$  = average time to get a better individual through crossover & mutation
- Let  $p_i$  = probability a single crossover produces a better individual
- Number of individuals undergoing crossover =  $p_c n$
- Probability of improvement =  $p_i p_c n$
- Estimate:  $t_i \approx 1 / (p_c p_i n)$

# Steady State Innovation

- Bad:  $t^* < t_i$ 
  - because once you have takeover, crossover does no good
- Good:  $t_i < t^*$ 
  - because each time a better individual is produced, the  $t^*$  clock resets
  - *steady state innovation*
- Innovation number:

$$\text{Iv} = \frac{t^*}{t_i} = p_c p_i \frac{n \ln n}{\ln s} > 1$$

# Feasible Region



# Other Algorithms Inspired by Genetics and Evolution

- Evolutionary Programming
  - natural representation, no crossover, time-varying continuous mutation
- Evolutionary Strategies
  - similar, but with a kind of recombination
- Genetic Programming
  - like GA, but program trees instead of strings
- Classifier Systems
  - GA + rules + bids/payments
- and many variants & combinations...



# Additional Bibliography

1. Goldberg, D.E. *The Design of Innovation: Lessons from and for Competent Genetic Algorithms*. Kluwer, 2002.
2. Milner, R. *The Encyclopedia of Evolution*. Facts on File, 1990.

