

Algorithmes Évolutionnaires (M2 MIAGE IA²)

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Séance 2 Théorie

Plan

- Les AE comme processus aléatoires
- Parallélisme implicite et théorème des schémas
- L'hypothèse des « blocs de construction »
- Quand la *fitness* est trompeuse
- Convergence

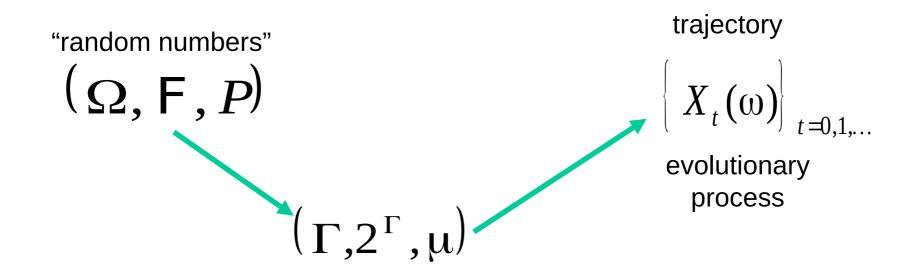
Some notation

- Space of "genotypes" (individuals): Γ
- Space of "phenotypes" (candidate solutions): Φ
- Decoding function: M: $\Gamma \rightarrow \Phi$
- Fitness function: f: $\Gamma \rightarrow [0, +\infty)$, f(y) = F[c(M(y))]
- For all $\gamma, \kappa \in \Gamma, f(\gamma) > f(\kappa)$ iff $c(M(\gamma)) < c(M(\kappa))$
- Space of populations: Γ*
- The composition of a population is described by its share function q: $\Gamma \to [0,\,1]$
- Share of individual y in the population: q(y)



 $(\Gamma, 2^{\Gamma}, \mu)$ probability space

 $x \in \Gamma^{(n)}$ a sample of size *n*



The Schema Theorem

- Originally developed by Holland for simple GAs
 - Bit string representation
 - Uniform mutation
 - One-point crossover
 - Fitness-proportionate selection
- Historically the first justification of *why* EAs work
- Provides interesting insights into *how* EAs work
- Can be extended to other types of
 - Encoding
 - Genetic operators
 - Selection strategies

Definition of Schema

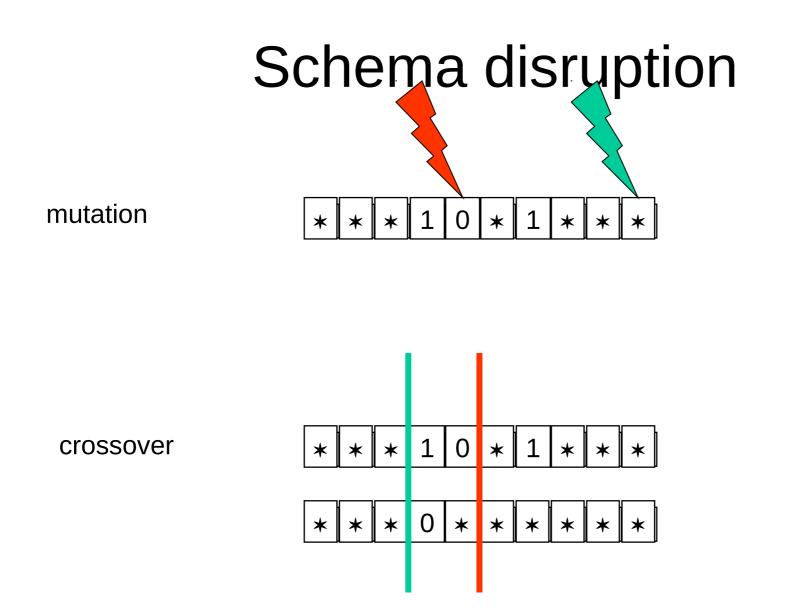
- A schema is a subset $S \subseteq \Gamma$
- Represented by a template string
- Symbols in {0, 1, *}

order of a schema: o(S) = # fixed positions defining length $\delta(S) =$ distance between first and last fixed position

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Properties of schemata

- A schema S matches 2^{1 o(S)} distinct strings
- A string of length / is matched by 2[/]
 - schemata



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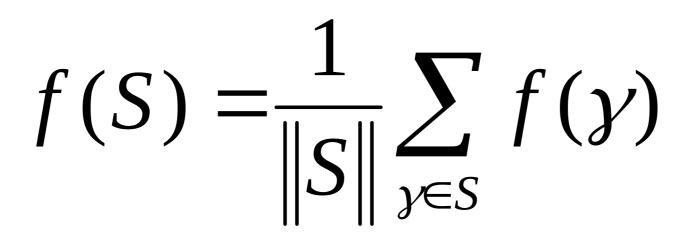
"Implicit Parallelism"

In a population of *n* individuals of length *l* $2^{\prime} \leq \#$ schemata processed $\leq n2^{\prime}$ n^{3} of which are processed usefully (Holland 1989)

(i.e. they are not disrupted by crossover and mutation)

Absolute fitness of a schema

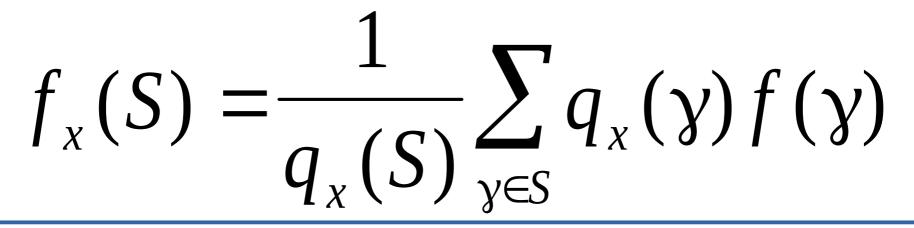
- Expected fitness for any of the matched strings
- Cannot be directly observed by the GA



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Relative fitness of a schema

- Expected fitness of an individual in a population *x*...
- ... Given that it belongs to a schema S



The Schema Theorem

 ${X_t}_{t=0,1,\dots}$ populations at times *t*

suppose that
$$\frac{f_{X_t}(S) - f(X_t)}{f(X_t)} = c \quad \text{is constant}$$

t

$$E[q_{X_t}(S)|X_0] \ge q_{X_0}(S)(1+c)^t \left(1 - p_{cross} \frac{\delta(S)}{l-1} - o(S)p_{mut}\right)^t$$

i.e. short, low-order above-average schemata increase exponentially!

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The Schema Theorem (proof)

$$E[q_{X_{t}}(S)|X_{t-1}] \ge q_{X_{t-1}}(S) \frac{f_{X_{t-1}}(S)}{f(X_{t-1})} P_{surv}[S] = q_{X_{t-1}}(S)(1+c)P_{surv}[S]$$

$$P_{surv}[S] = 1 - p_{cross} \frac{\delta(S)}{1 - l} - p_{mut} o(S)$$

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Discussion of the Schema Theorem

- Hypothesis: ratio between schema fitness and average fitness of population is constant.
- This is only true "locally", i.e., within few generations.
- Therefore, the exponential growth of aboveaverage schemata happens only locally.
- However, the growth of high-fitness schemata has an effect on the average fitness in the population.
- Soon this effect compensates and balances the growth.

Punctuated Equilibria

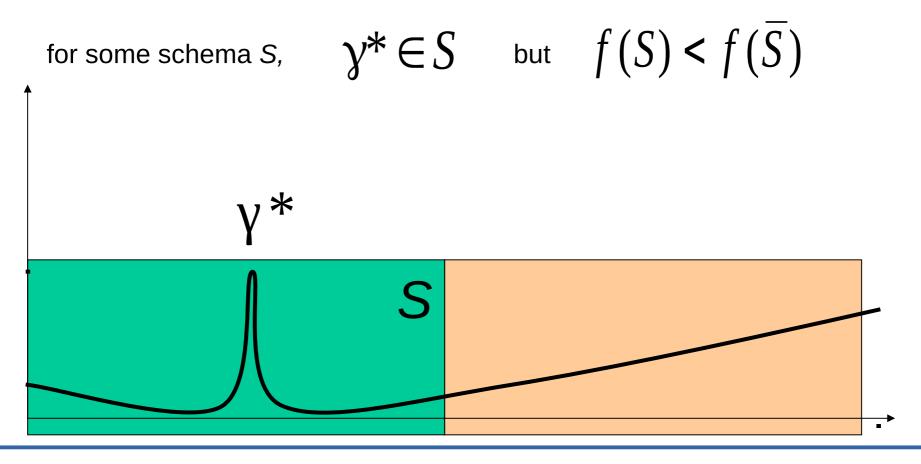
- Evolution proceeds in leaps and bounds
- Long periods of equilibrium
- Punctual events cause revolutionary changes
- Innovation
- Good match with observations of natural evolution

The Building Blocks Hypothesis

"An evolutionary algorithm seeks nearoptimal performance through the juxtaposition of short, low-order, highperformance schemata — the building blocks"

Deception

i.e. when the building block hypothesis does not hold:



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Deception: Example

$$S_{1} = 111^{*******}$$

$$S_{2} = ********11$$

$$S = 111^{******11}$$

$$S = 000^{*****00}$$

$$\gamma^* = 1111111111$$

$$f(S_1) = 3$$

 $f(S_2) = 2$
 $f(S) = 5$
 $f(\overline{S}) = 10$

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Critical success factors

- Solution encoding
- Fitness function
- Prior knowledge of the problem (if available)
- Trial and error (otherwise)
- Rules of thumb
- Analogy with other problems

Convergence properties of EAs

- Does my evolutionary algorithm converge to the optimal solution?
- How fast does my evolutionary algorithm find the optimal solution?

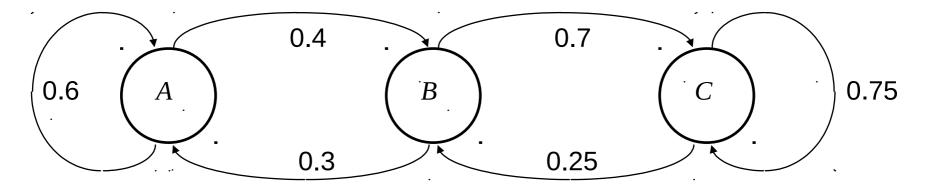
Markov Chains

A stochastic process

 $X_t(\omega)$

is a Markov chain if and only if its state depends only on the previous state, i.e., for all t,

$$P[X_{t} = x | X_{0}, X_{1}, \dots, X_{t-1}] = P[X_{t} = x | X_{t-1}]$$

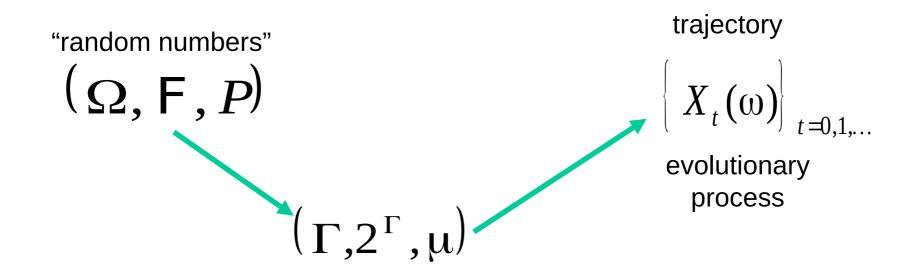


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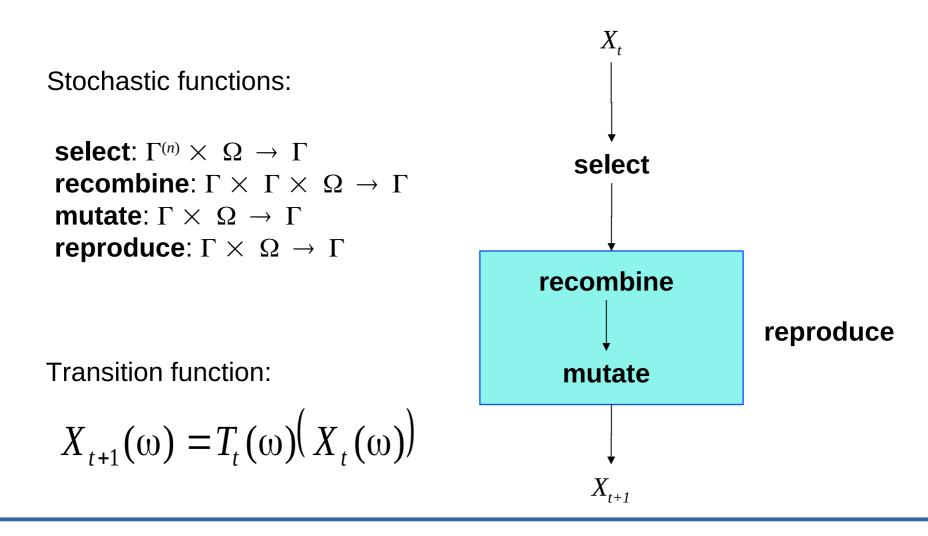


 $(\Gamma, 2^{\Gamma}, \mu)$ probability space

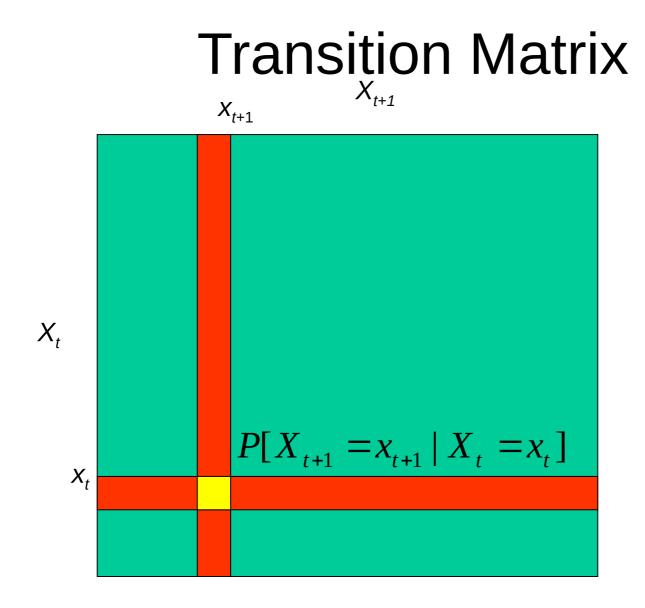
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Abstract Evolutionary Algorithm



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Convergence to Optimum

Theorem: if $\{X_t(\omega)\}_{t=0, 1, ...}$ is *monotone*, *homogeneous*, x_0 is given, $\forall y$ in **reach** $(x_0) \exists y \in \Gamma^{(n)}{}_O$ reachable, then

$$\lim_{t\to\infty} P[X_t \in \Gamma_O^{(n)} | X_0 = x_0] = 1.$$

Theorem: if **select**, **mutate** are *generous*, the neighborhood structure is *connective*, transition functions $T_t(\omega)$, t = 0, 1, ... are i.i.d. and *elitist*, then

$$\lim_{t\to\infty} P[X_t \in \Gamma_O^{(n)}] = 1.$$

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Speed of Convergence

- Still an open problem
- Very difficult
- Partial attempts
- Complexity classes

