

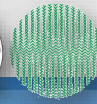
Promoting Diversity in Evolutionary Optimization: Why and How

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1

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2

2

Outline

- Generic EA
- Divergence of character in natural and artificial evolution
- Background (diversity and similarity, ...)
- Mechanisms for promoting diversity
- Hints and tips
- Conclusion

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3

3

Real world (Galapagos)



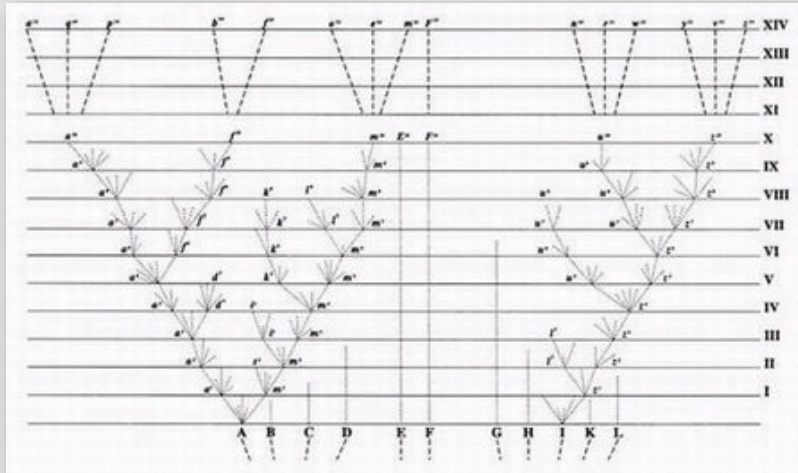
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Darwin's tree of life



The only illustration in *On the Origin of Species by Natural Selection* (1859)

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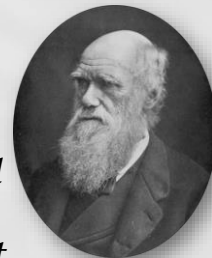
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Divergence of character

- *"Great diversity of forms in nature"*
- *"The principle, which I have designated by this term, is of high importance, and explains, as I believe, several important facts"*
 - *"The principle of divergence causes differences, at first barely appreciable, to steadily to increase, and the breeds to diverge in character, both from each other and from their common parent"*
 - *"The varying descendants of each species try to occupy as many and as different places as possible in the economy of nature"*



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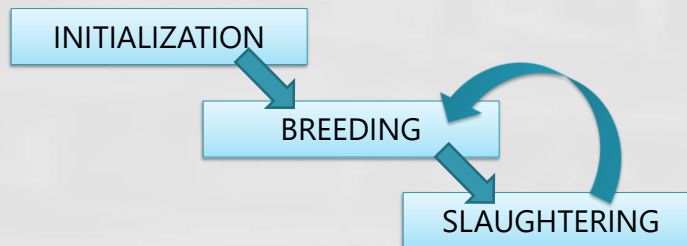
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Evolutionary computation

- A rough idea about “what” an evolutionary algorithm is



- Note: Optimization, not artificial life!

7

Evolutionary algorithms



8

Premature convergence

- I.e., the tendency of an algorithm to converge towards a point where it was not supposed to converge to in the first place
- Probably an oxymoron
- Holland's "Lack of speciation"
- EAs general inability to exploit environmental niches

9

divergence of character

vs.

premature convergence

10

Divergence of character

- “The basic point of the principle of divergence is **simplicity itself**: the more the coinhabitants of an area differ from each other in their ecological requirements, the less they will compete with each other; therefore natural selection will tend to favor any variation toward greater divergence.”



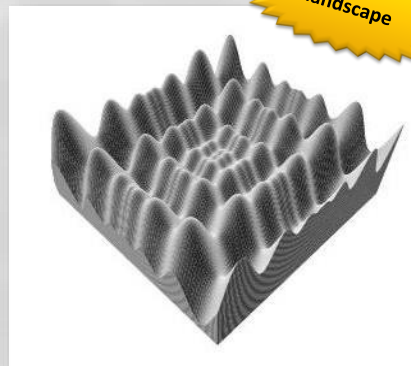
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Environment vs. Fitness function

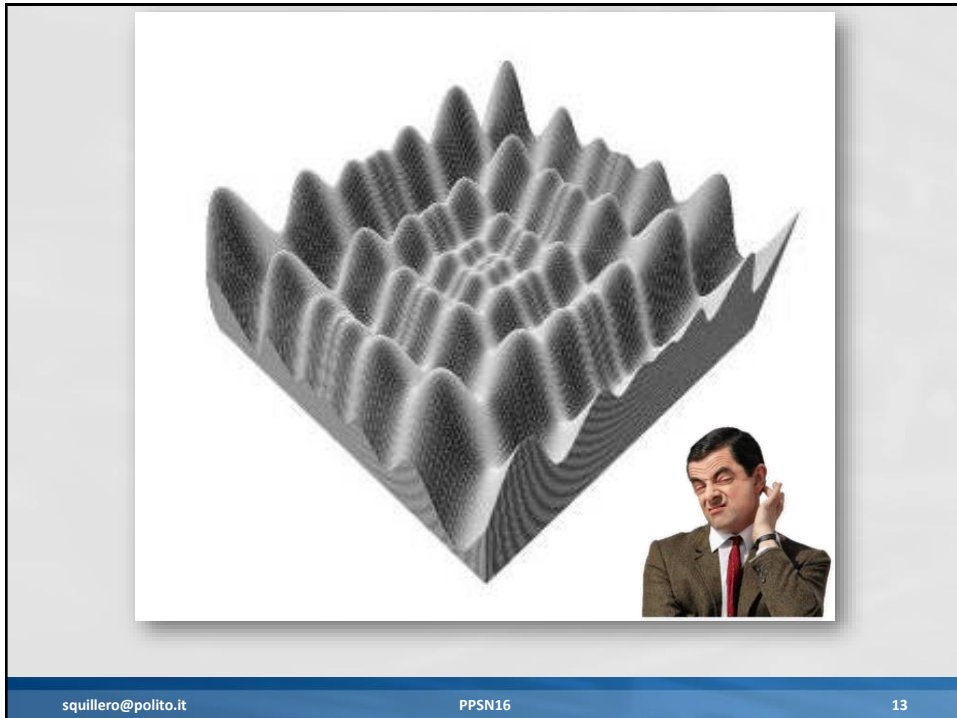


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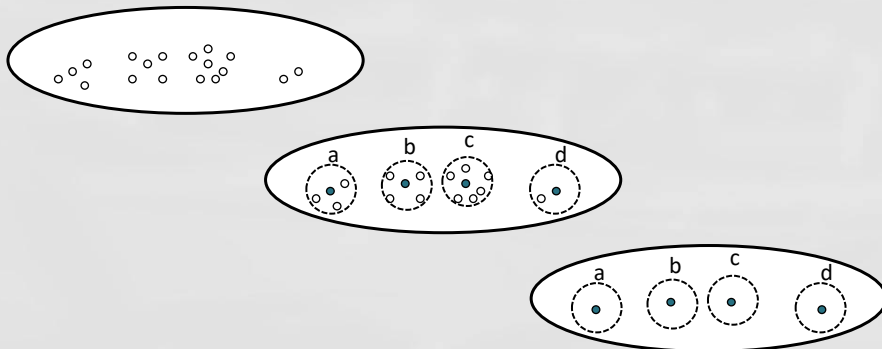
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Niches

- Niche: subspace in the environment with a finite amount of physical resources that can support different types of life



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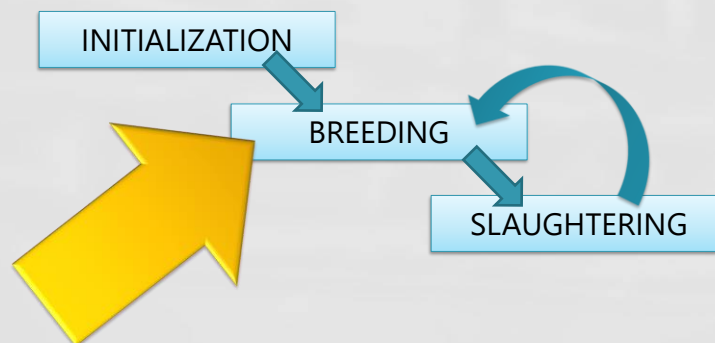
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Niches

- Niches favor the divergence of character
- Niches and speciation
- How to create “niches” in EAs since the environment is missing?

15

A Very Generic EA



16

Exploration vs. Exploitation

• Recombination

- mixes together two or more solutions to create the offspring
- associated with the idea of *exploration*

• Mutation

- performs a (usually small) change in an individual
- associated with the idea of *exploitation*



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17

Exploration vs. Exploitation

- When all parents are very similar, the effectiveness of recombination is limited
- The ability to explore remote parts of the search space is impaired
- “Conventional wisdom suggests that increasing diversity should be generally beneficial”



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18

Exploration vs. Exploitation

- When all parents are **very similar**, the effectiveness of reproduction is limited
- **what is the definition of "similar"?**
- The ability to explore remote parts of the search space is impaired
- "Conventional wisdom suggests that increasing **diversity** should be generally beneficial"

and the definition of "diversity"?



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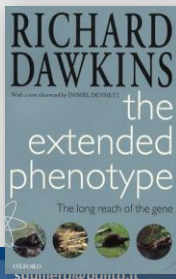
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19

Levels in biology

- **Genotype**: the genetic constitution of an organism
- **Phenotype**: the composite of the organism's observable characteristics or traits
- **Fitness**: individual's ability to propagate its genes (well, almost)



Richard Dawkins

The Extended Phenotype: The Long Reach of the Gene
Oxford University Press, 1982 (revised ed. 1999)

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20

Levels in EC (a modest proposal)

- **Fitness**: how well the candidate solution is able to solve the target problem
- **Genotype**: the internal representation of the individual, i.e., what is directly manipulated by genetic operators
- **Phenotype**: the candidate solution that is encoded in the genotype
 - the intermediate form in which the genotype needs to be transformed into for evaluating fitness
 - if genotype can be directly evaluated: genotype and phenotype coincide

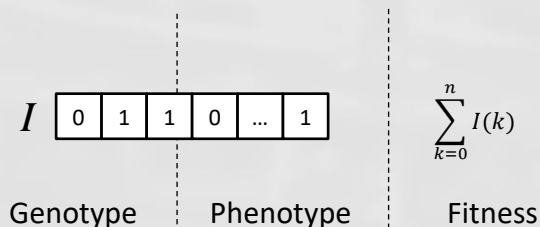
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21

Levels in EC (GA)



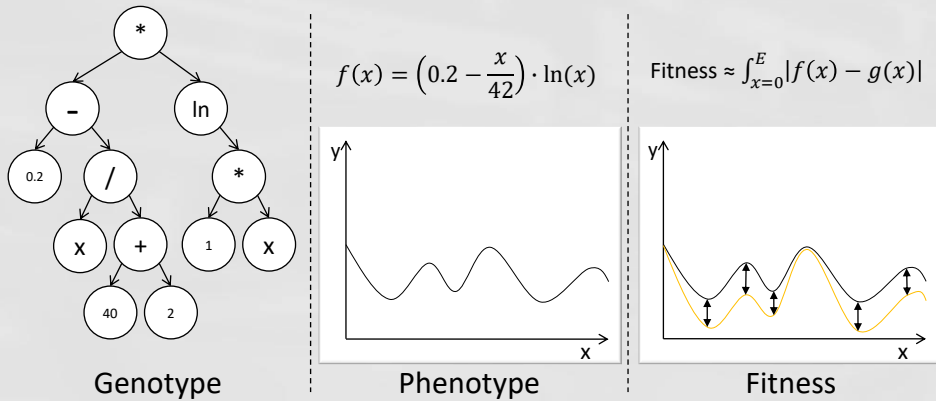
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Levels in EC (GP & LGP)



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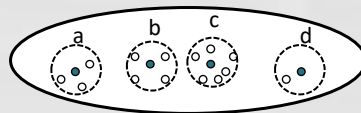
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23

Niches in EA

- Niching: grouping similar individual
 - similar spatial positions (i.e., islands)
 - similar genotypes (i.e., niching)
 - similar phenotypes
- Several approaches are based on niching



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24

Detecting clones

- Detecting whether two individuals are clones, i.e., identical, is often an easy task at any level



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25

Measuring diversity

- Diversity \Rightarrow distance metric: *how far* the individual is
 - from (a subset of) the whole population
 - from a single individual
- Diversity \Rightarrow property of the population
- But, at what level?
 - Phenotype
 - Genotype
 - Fitness



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Measuring diversity

- Different fitness values imply different phenotypes, different phenotypes imply different genotypes

$$F_x \neq F_y \Rightarrow P_x \neq P_y \Rightarrow G_x \neq G_y$$

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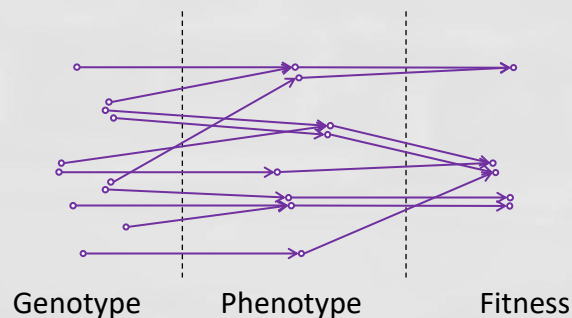
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Measuring diversity

- What about “diversity”?
- Locality principle
- Rechenberg’s *strong causality*



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28

Measuring diversity

- Fitness
 - Usually trivial
- Phenotype
 - Usually ad-hoc
- Genotype
 - Different genotypes in the population
 - GP subtree frequency
 - Edit distance (a.k.a., Levenshtein distance)
 - Entropy and free energy

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29

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What has been proposed
to alleviate it?

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30

End goal vs. Means goal

- The **end goal** in optimization is reaching better solutions in less time
- Promoting diversity has often been seen as the key factor to improve performances
- Promoting diversity is a mere **means goal** (yet a quite important one)
- No distinction is made here whether the means goal is
 - preserve existing diversity
 - increase diversity

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31

How diversity is promoted (practice)

- Fitness scaling
- Fitness holes
- Tweaking selection mechanism
- Adding selection mechanism
- Multiple populations
- Population topologies
- ...

In theory there is no
difference between
theory and practice



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How diversity is promoted (theory)

- A methodology for promoting diversity alters the selection probability of individuals

$$\bar{p}_{x|\Psi} = p_{x|\Psi} \cdot \xi(x, \Psi)$$

- **Mere definition:** we do not imply that a mechanism operates *explicitly* on the selection operators
- **But** the *effects* on selection probabilities are assessed to classify it

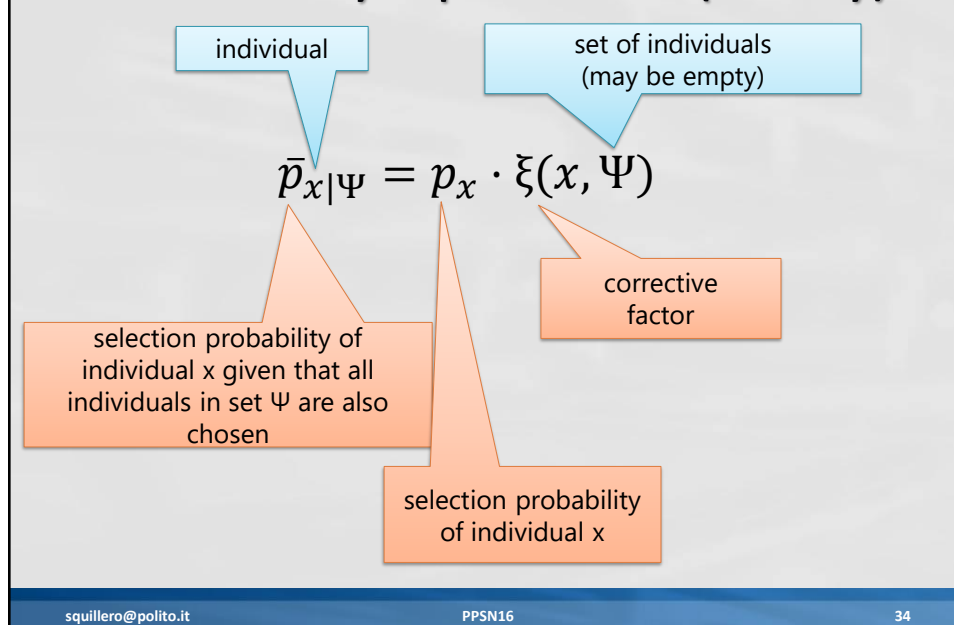
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How diversity is promoted (theory)



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34

Relevant characteristic

- Lineage (**LIN**)
- Phenotype (**PHE**)
- Genotype (**GEN**)
- ~~Fitness~~ (used as a proxy for either phenotype or genotype)

$$\xi(x, \Psi)$$

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35

Lineage-based methodologies

- The value of $\xi(\circ)$ does not depend on individual structure nor behavior, but it can be determined considering circumstances of its birth (e.g., time, position)
- LBMs can be applied to any kind of problem, even in addition to other diversity preservation methods

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36

Genotype-based methodologies

- Particularly effective when it is possible to define a sensible distance between genotypes
- Often used to
 - avoid overexploitation of peaks in the fitness landscape
 - promote the generation of new solutions very far from the most successful ones
 - preserve variability in the gene pool

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37

Phenotype-based methodologies

- Usually impractical
- Sometimes fitness distance can be used as a proxy for phenotype distance (multi objective EAs, or many objective EAs)

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38

38

Type of selection

- Parent selection (α or ϵ)
 - Usually non-deterministic
- Survival selection (ω or μ)
 - Usually deterministic

$$\bar{p}_{x|\Psi}$$

Diversity Promotion in EA



Information Sciences

Volume 329, 1 February 2016, Pages 782–799

Special issue on Discovery Science



Divergence of character and premature convergence: A survey of methodologies for promoting diversity in evolutionary optimization ☆

Giovanni Squillero · , Alberto Tonda^b

DOI: [10.1016/j.ins.2015.09.056](https://doi.org/10.1016/j.ins.2015.09.056)



Island model

• Recipe [$LIN \alpha \omega$]

- The population is partitioned into sub-populations
- Only local interactions are allowed
- Periodically, individuals are moved between sub-populations (**migrants**)

• Rationale

- Different populations may explore different parts of the search space
- ... but global interactions can be useful



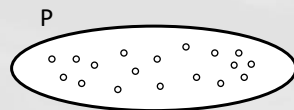
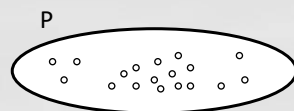
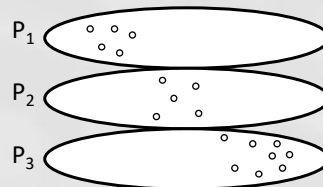
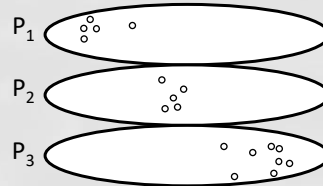
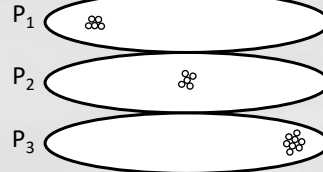
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Island model

 $T = t_0$  $T = t_k$  $T = t_N$ 

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42

Segregation

• Recipe [$\text{LIN} \alpha \omega$]

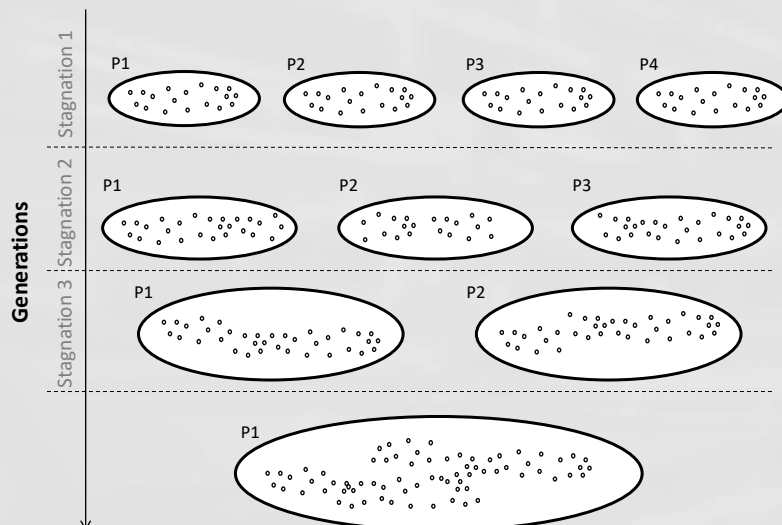
- The population is partitioned into N sub-populations
- Only local interactions are allowed
- Upon stagnation, the N sub-populations are merged into $N-1$ sub-populations

• Rationale

- Same as island models
- The selective pressure decreases during evolution

43

Segregation



44

Hierarchical fair competition

• Recipe [**PHE** $\alpha\omega$]

- The population is partitioned into sub-populations with similar fitness
- Only local interactions are allowed
- The offspring is promoted or demoted according to fitness
- New random individuals are constantly generated

• Rationale

- Hard niching with implicit neighborhood
- Reduce competition between newborns and already optimized individuals (ladder)

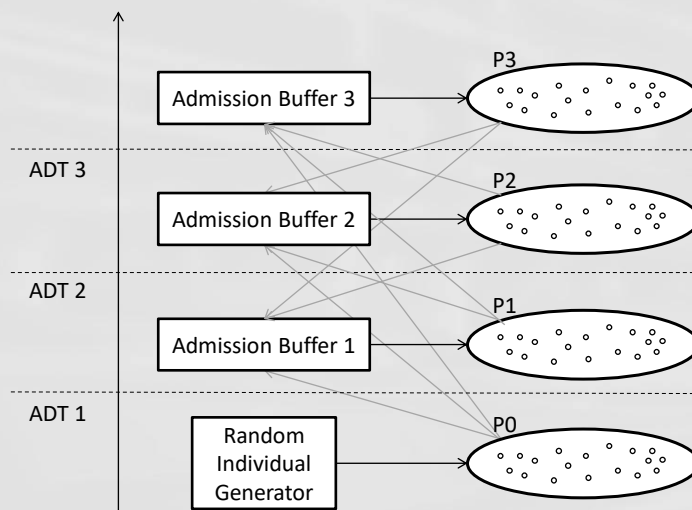
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Hierarchical fair competition



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46

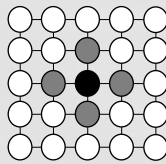
Cellular EA

• Recipe [**LIN** $\alpha\omega$]

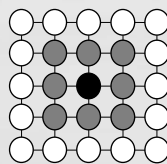
- Fixed topology (lattice)
- Only interactions between neighbors are allowed

• Rationale

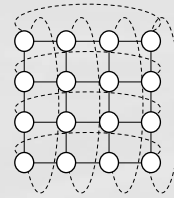
- Limiting interaction could defer the takeover of the population by clones of the fittest individual



Linear-5 (L5)



Compact-9 (C9)



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47

Deterministic crowding

• Recipe [**LIN** $\epsilon\omega$]

- Offspring compete against parents for survival

• Rationale

- Flexible niching with implicit neighborhood
- Parents and offspring occupy the same niche
- No need for evaluating the similarity

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48

Allopatric selection

- Recipe [**LIN** $\alpha\omega$]
 - The whole offspring compete for survival
- Rationale
 - Flexible niching with implicit neighborhood
 - No need for evaluating the similarity
 - Genetic operators that create large offspring can be exploited without the risk for the offspring to invade the population

49

Fitness Sharing

- Recipe [**GEN** $\alpha\omega$]
 - Scale down individual fitness

$$\bar{f}(I_k) = \frac{f(I_k)}{\sum_i sh(I_k, I_i)}$$
 - with $sh(x, y)$ depending on the distance between the individuals, and is 0 beyond a fixed radius
- Rationale
 - Flexible niching with explicit neighborhood
 - Reduce attractiveness of densely populated area

50

Clearing

- Recipe [GEN $\alpha\omega$]
 - Inside niches of a certain radius, the best k individuals retain their fitness while the rest are zeroed
- Rationale
 - Flexible niching with explicit neighborhood
 - Set a hard limit to population density

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51

Standard crowding

- Recipe [GEN $\alpha\omega$]
 - New individuals replace the most similar individual in a random niche of size CF
- Rationale
 - Flexible niching with explicit neighborhood
 - Favor novelty (generational approach)

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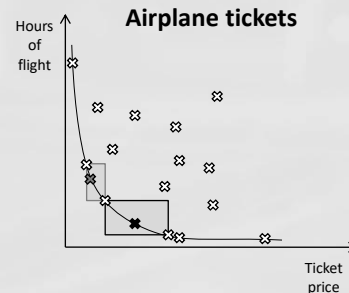
Crowded-comparison operator

• Recipe [PHE $\alpha\omega$]

- Estimate the *free territory* around solutions and favor solutions less crowded regions

• Rationale

- Smart implementation of artificial niches
- Requires a strong correlation between phenotype and fitness
- NSGA-III introduces ϵ -domination (adaptive discretization)



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53

Reference points partitioning

• Recipe [GEN $\alpha\omega$]

- Population is partitioned using in clusters centered around a set of reference points
- Reference points are initially chosen by the user, then can be dynamically updated
- New individuals compete for survival inside their own niche

• Rationale

- Flexible niching with explicit neighborhood

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54

Vector evaluated genetic algorithm

• Recipe [PHE ~~α~~]

- Divide the mating pool in N parts, each one filled with individual selected on their i -th component of the fitness
- Alternative: select on a weighted sum, but use different weight sets for the different parts

• Rationale

- Increase the push towards specialization

• Caveats

- Only applicable to MOEAs, or when using an aggregate fitness

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Lexicase selection

• Recipe [PHE ~~α~~]

- Before selection, re-arrange the components of the fitness
- Compare individual fitnesses lexicographically

• Rationale

- Increase the push towards specialization

• Caveats

- Only applicable when using an aggregate fitness

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Restricted tournament selection

- Recipe [GEN $\alpha\omega$]
 - New individuals compete with the most similar individual in a random niche of size CF
- Rationale
 - Flexible niching with explicit neighborhood

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Sequential niching

- Recipe [GEN $\alpha\omega$]
 - The most promising points in the search space after each run are altered so to become less interesting in further executions
- Rationale
 - Avoid over exploitation

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Gender

- Recipe [**LIN**/**GEN** $\alpha\omega$]
 - Add gender to individual and enforce sexual reproduction
 - More than two sexes are possible, with different mutation probabilities
 - Gender might be part of the genome or not
- Rationale
 - Prevent crossover between clones
 - Limit interactions between related individuals

59

Tarpeian method

- Recipe [**PHE** $\alpha\omega$]
 - Randomly kill individual who don't adhere to given standards
- Rationale
 - Note: originally used to prevent bloat
 - Creating dynamic and non-deterministic *fitness holes* may have several beneficial effects, including to promote diversity

60

Diversifiers

- Recipe [**GEN** $\alpha\omega$]
 - Detect less populated areas in the search space and try to generate random inhabitants
- Rationale
 - Increase variability in the gene pool regardless the fitness
 - Require a reliable distance metric

61

Random immigrants

- Recipe [**PHE** $\alpha\omega$]
 - Periodically insert random individuals in the population
- Rationale
 - Try to introduce novelty
- Caveats
 - Newborns may need to be artificially kept alive when competing against already optimized individuals

62

Extinction

• Recipe [PHE $\alpha\omega$]

- Upon convergence (or periodically) remove a significant part of the population
- Then fill up the population with the offspring of the survivors and/or random individuals

• Rationale

- A gust of fresh air: already optimized individuals are not enough to occupy the whole population and newborns may start exploring new regions

• Caveat

- Fitness variability used as phenotype variability

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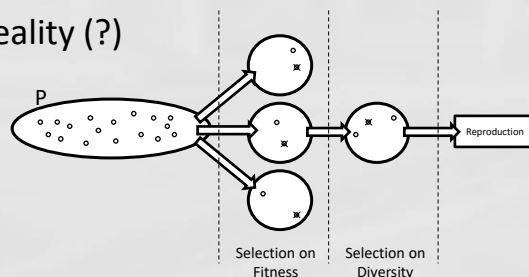
Two-level diversity selection

• Recipe [GEN $\alpha\omega$]

- Select three individuals using fitness, then pick the two with maximum distance for reproduction

• Rationale

- Exploit a reliable distance metric to increase the efficacy of crossover
- Not so far from reality (?)



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GDEM – Genetic Diversity Evaluation Method

- Recipe [**GEN** $\alpha\omega$]
 - Add diversity as an explicit goal and go MO
- Rationale
 - Modify the domination criteria
 - Need a reliable diversity metric
- Historical note
 - See: *Find Only and Complete Undominated Sets* (FOCUS)

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Delta entropy and pseudo entropy

- Recipe [**GEN** $\alpha\omega$]
 - With a certain probability select individuals on their ability to increase the global entropy of the population instead of fitness
- Rationale
 - Not-so-fit individual with peculiar traits should be preserved
 - Measuring the entropy of the population is easier than defining a distance function

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66

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67

Hints and Tips

- Do you really need to promote diversity?
 - Several problems in EA are caused by ill-designed fitness functions
 - Check whether the locality principle holds true
 - Check what happen with multistart



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68

Hints and Tips

- Do you really need to promote diversity?
- Use extinction (20m)
 - Simple n' easy



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69

69

Hints and Tips

- Do you really need to promote diversity?
- Use extinction (20m)
- Use lexicase selection (20m)
 - Simple n' easy
 - Only useful for aggregate fitness (combination of several components)



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70

Hints and Tips

- Do you really need to promote diversity?
- Use extinction (20m)
- Use lexicase selection (20m)
- Use an island model (2h)
 - Far better than multistart (if migrations are properly handled)
 - Only useful if different experiments yield different results



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71

Hints and Tips

- Do you really need to promote diversity?
- Use extinction (20m)
- Use lexicase selection (20m)
- Use an island model (2h)
- Use fitness holes (20h)
 - Tweak selection operator(s)
 - Only useful if a global (and efficient) diversity measure is available



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72

72

Hints and Tips

- Do you really need to promote diversity?
- Use extinction (20m)
- Use lexicase selection (20m)
- Use an island model (2h)
- Use fitness holes (20h)
- Use real niching (2-20d)
 - Only useful if the distance between genotypes is meaningful



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Conclusion



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75

75

More Materials & Bibliography

- **MPDEA**: GECCO Workshop on Measuring and Promoting Diversity in Evolutionary Algorithms
- mpdea@polito.it
- <https://github.com/squillero/mpdea>



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76

76