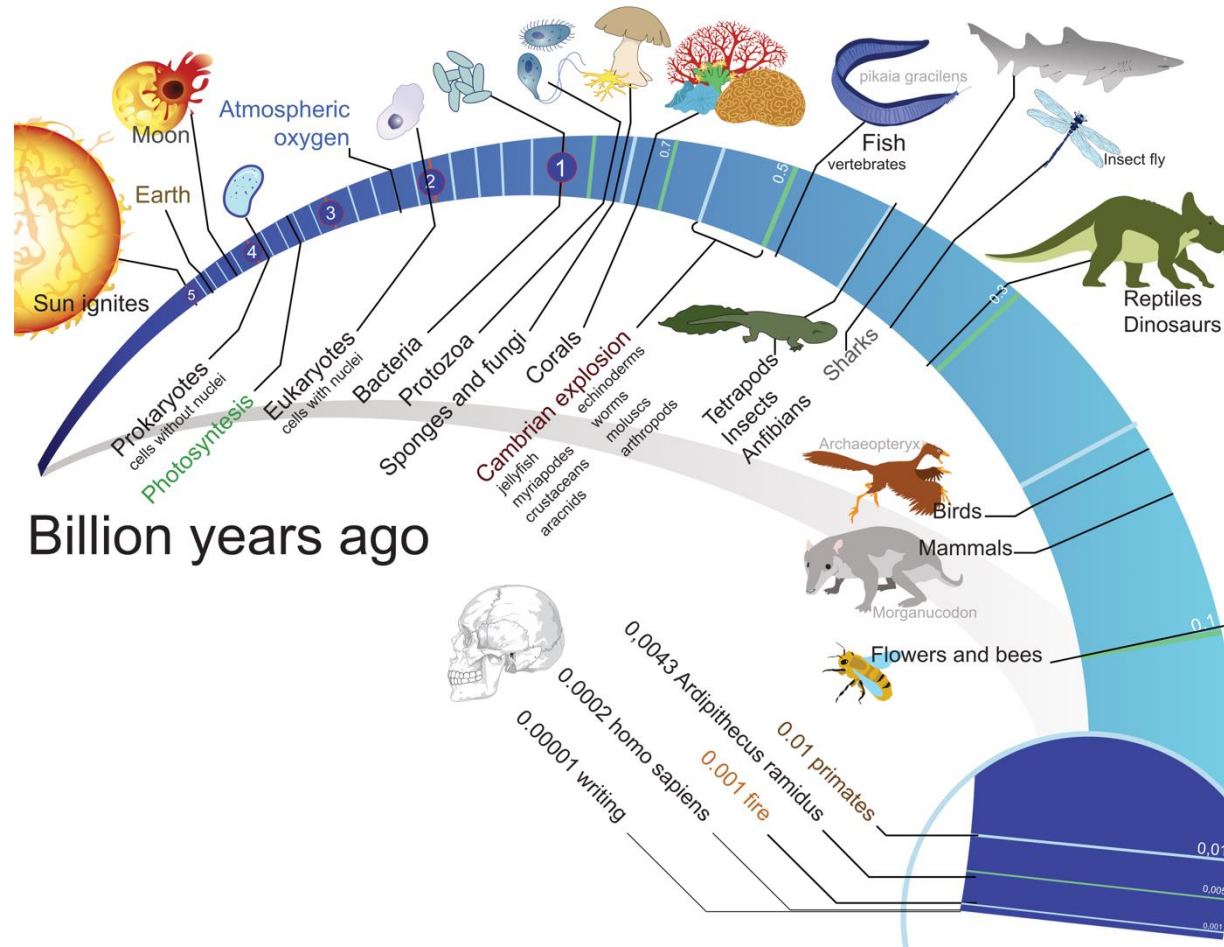


Quality Diversity Optimization

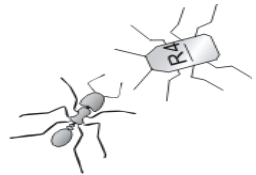


Evolution is diverse



What you will learn in this lecture

- Concept of premature convergence
- Population diversity: genotype vs. phenotype
- Quality Diversity algorithms
 - Open-ended evolution
 - Novelty search
 - Map elites



Examples of Evolutionary Algorithms

Genetic Algorithms (GA) - Holland, 1975

Binary genotypes, crossover and mutation

Genetic Programming (GP) - Koza, 1992

Tree-based genotypes, crossover and mutations

Steady-State GA (SSGA) – Whitley et al., 1988

Gradual replacement: Best individuals replace worst individuals

Differential Evolution (DE) – Storn & Prince, 1996

As SSGA, but with differential factor

Evolutionary Strategies (ES) - Rechenberg, 1973

Real-valued genotypes, mutation step(s) encoded in genotype

Covariance Matrix Adaptation ES (CMA-ES) – Hansen & Ostermeier, 2001

Evolutionary Strategies with correlated and adaptive mutations

Non-dominated Sorting GA (NSGA)– Srinivas, Deb, 1998

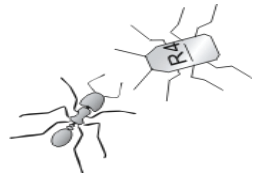
Multi-objective evolutionary optimization

Viability Evolution (VIE)– Maesani, Mattiussi, Floreano, 2014

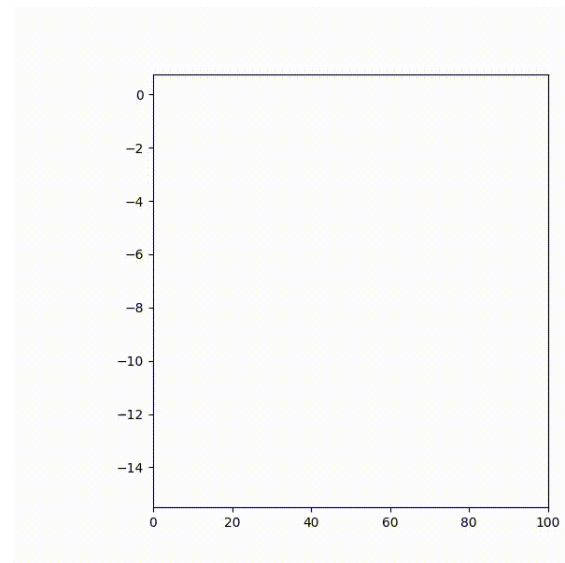
Evolution without fitness ranking and diversity preservation

MAP Elites – Mouret and Clune, 2015

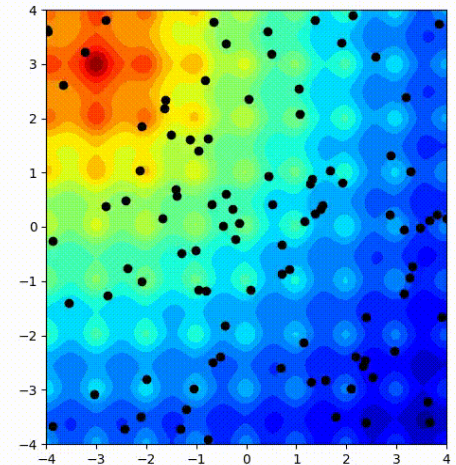
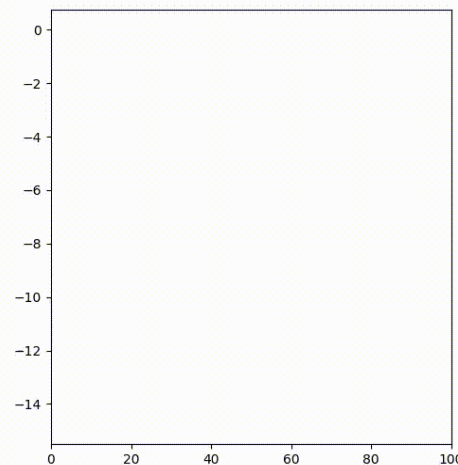
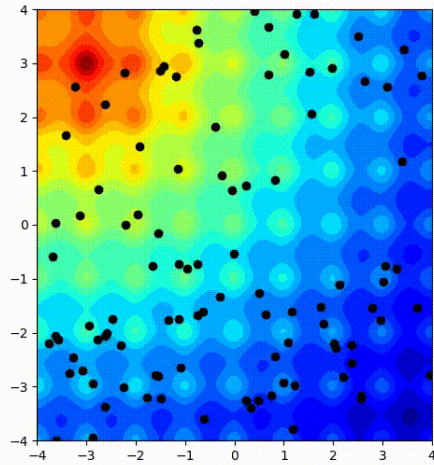
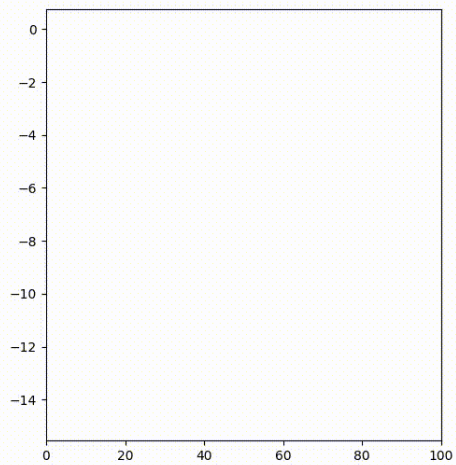
Preserve diversity by making similar solutions compete with each other



When to stop evolution?



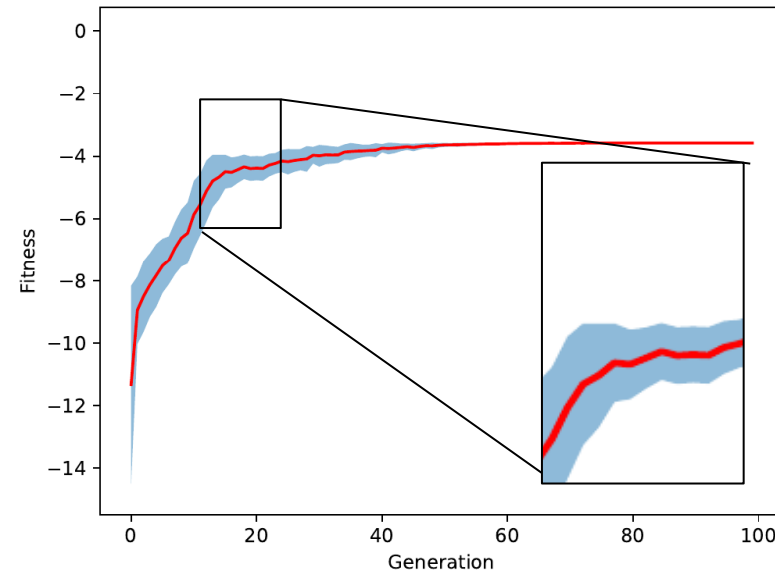
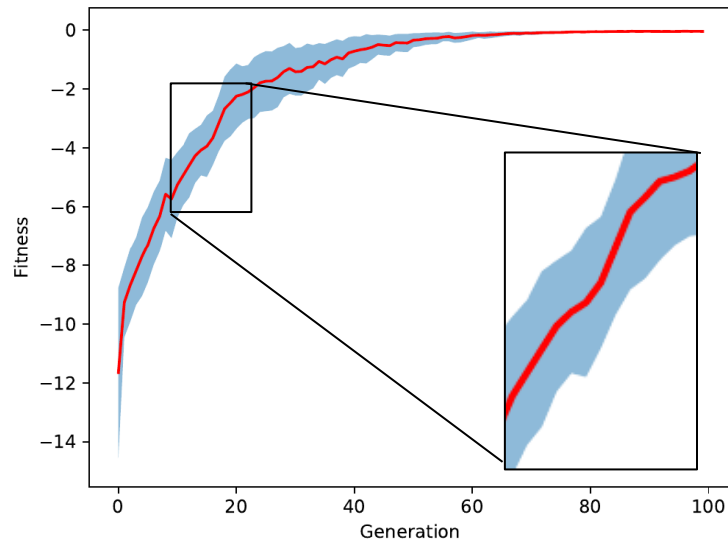
When to stop evolution?



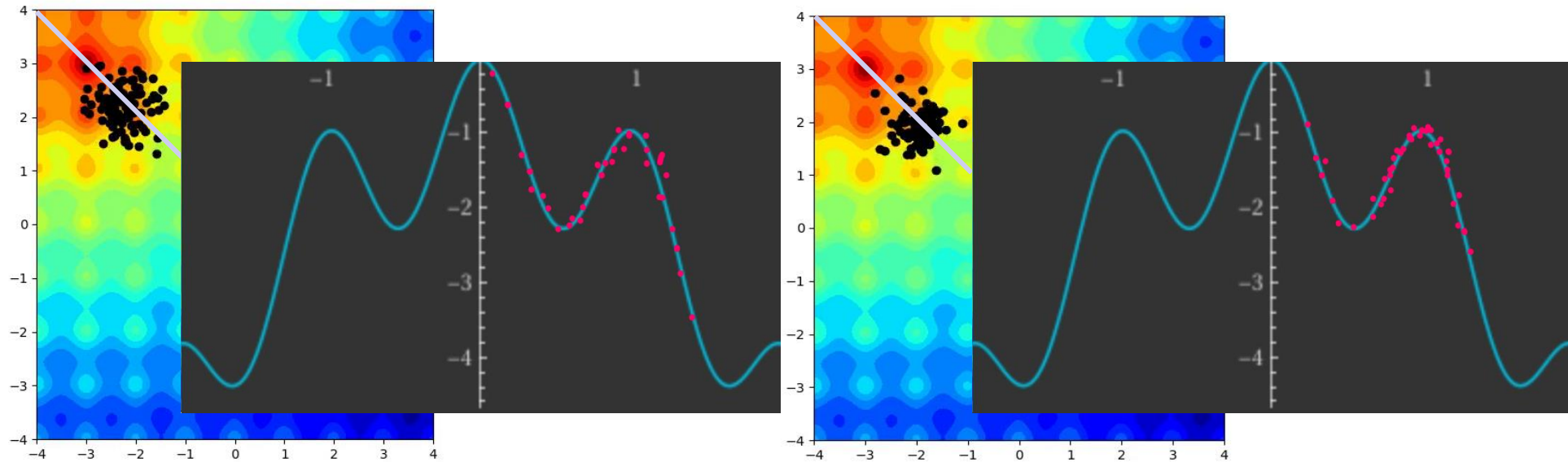
Premature convergence

Premature convergence is the situation where a population loses diversity too quickly and converges to a suboptimal solution (local optimum) before adequately exploring the search space.

- Quick reduction in spread of fitness (sign of convergence)



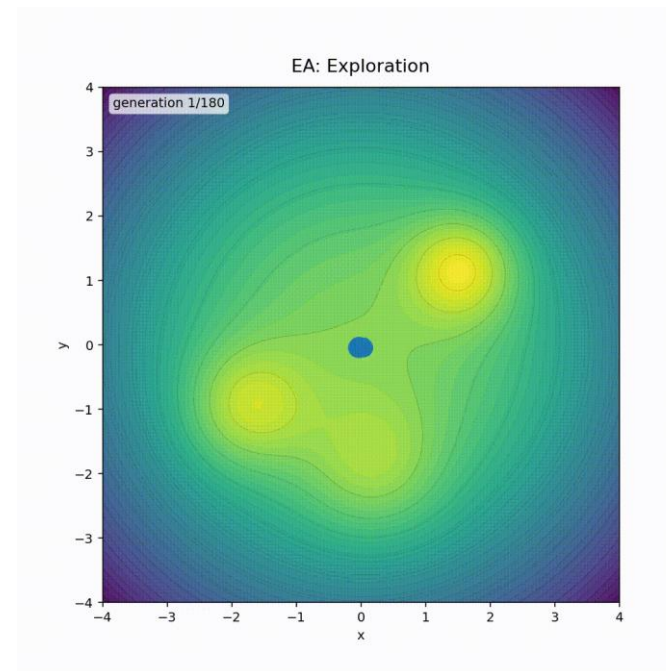
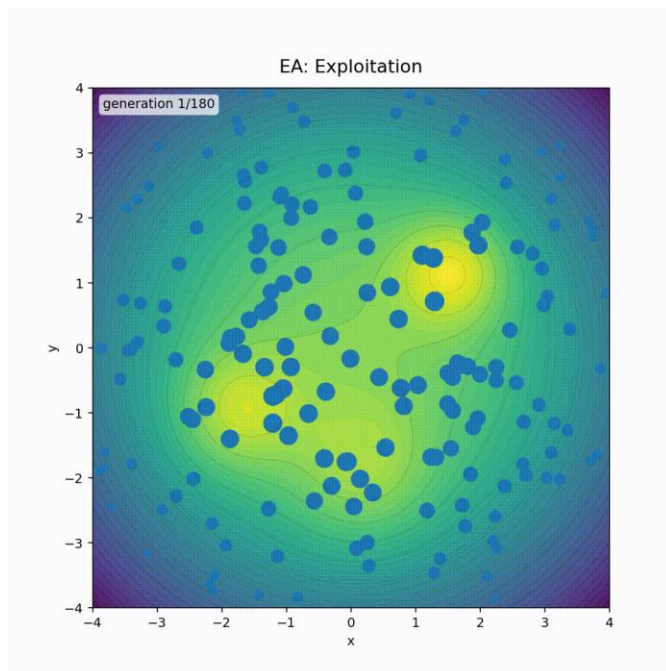
Sensitivity to genotype variation



Exploitation vs. Exploration

The EA balancing act:

- Converge on best solution -> Selection
- Search for better genotypes -> mutation



Genotype diversity

Premature convergence is the situation where a population loses diversity too quickly and converges to a suboptimal solution (local optimum) before adequately exploring the search space.

- Reduced spread in fitness (sign of convergence)
- Reduced spread in genotype
 - Adaptive step size (e.g. CMAES)
 - Co-evolve mutation rate

**MIT
Technology
Review**

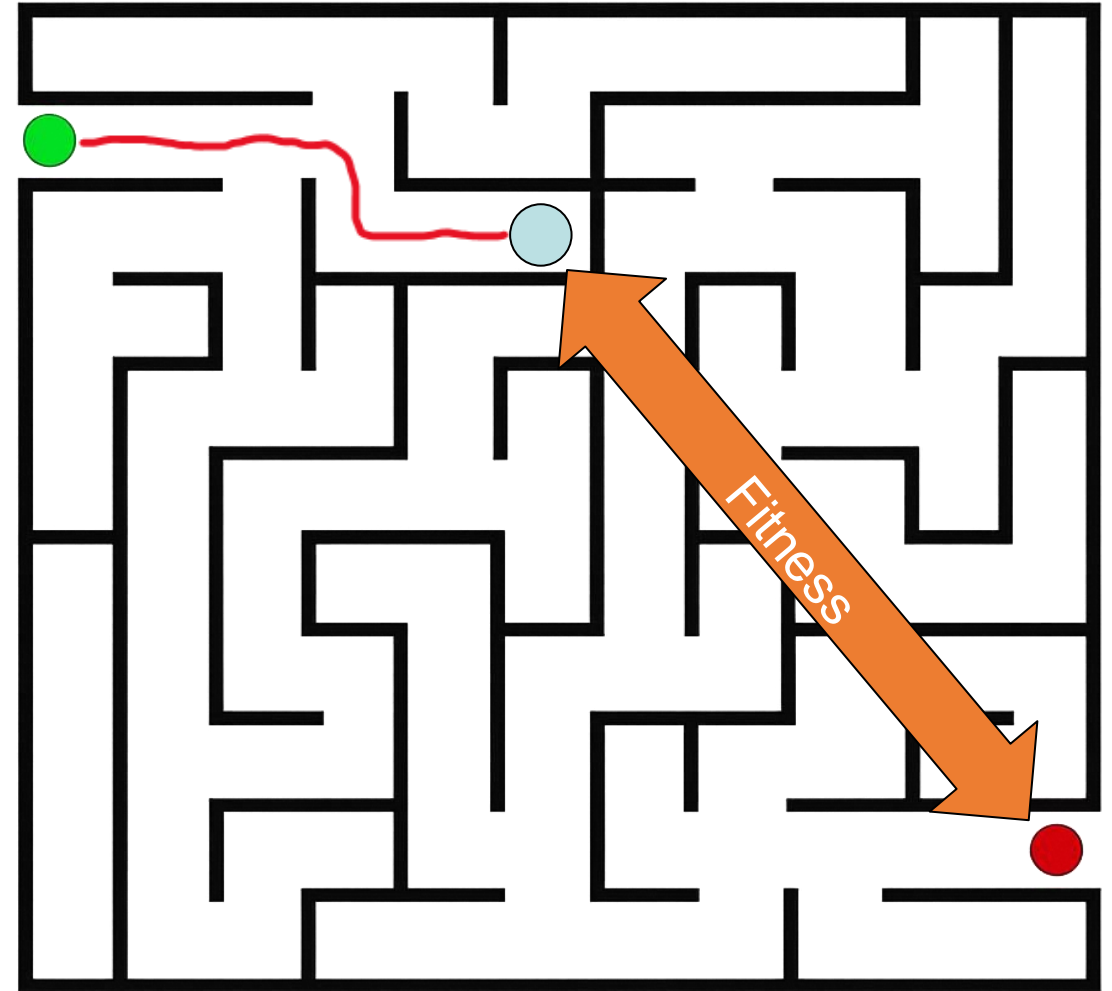
Featured Topics

scientists estimate we share about 50 percent of our DNA with the **banana**. This is the nature of DNA in evolution: that all organisms currently alive developed over the eons from common earlier organisms.

Maze problem

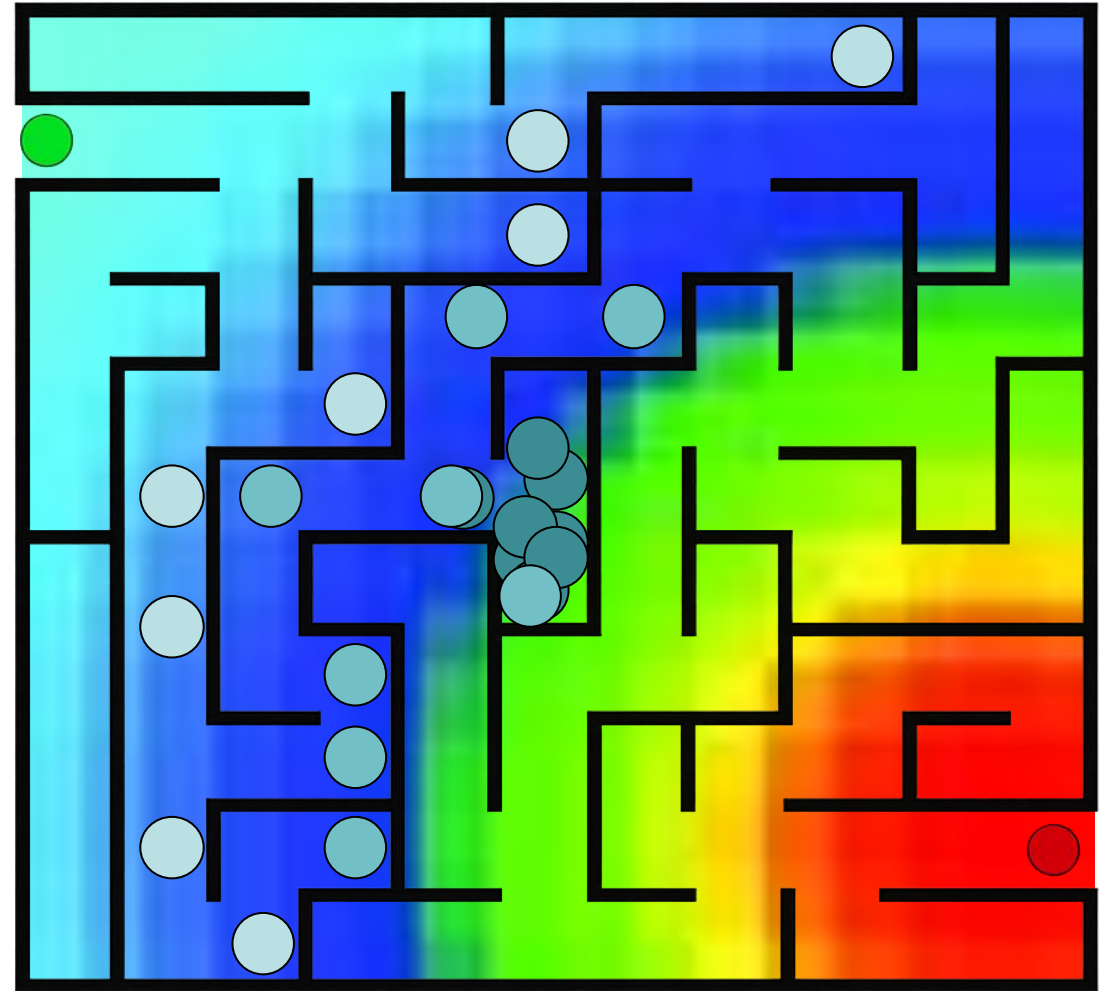
Maze problem:

- Robot actions $\{<, ^, v, >\}$
- Genotype = T actions (e.g. $[>, v, >]$)
- Fitness = $\text{dist}(\text{robot_pos}_T, \text{end_goal})$



The deceptive trap

Optimizing the maze problem leads to a deceptive trap



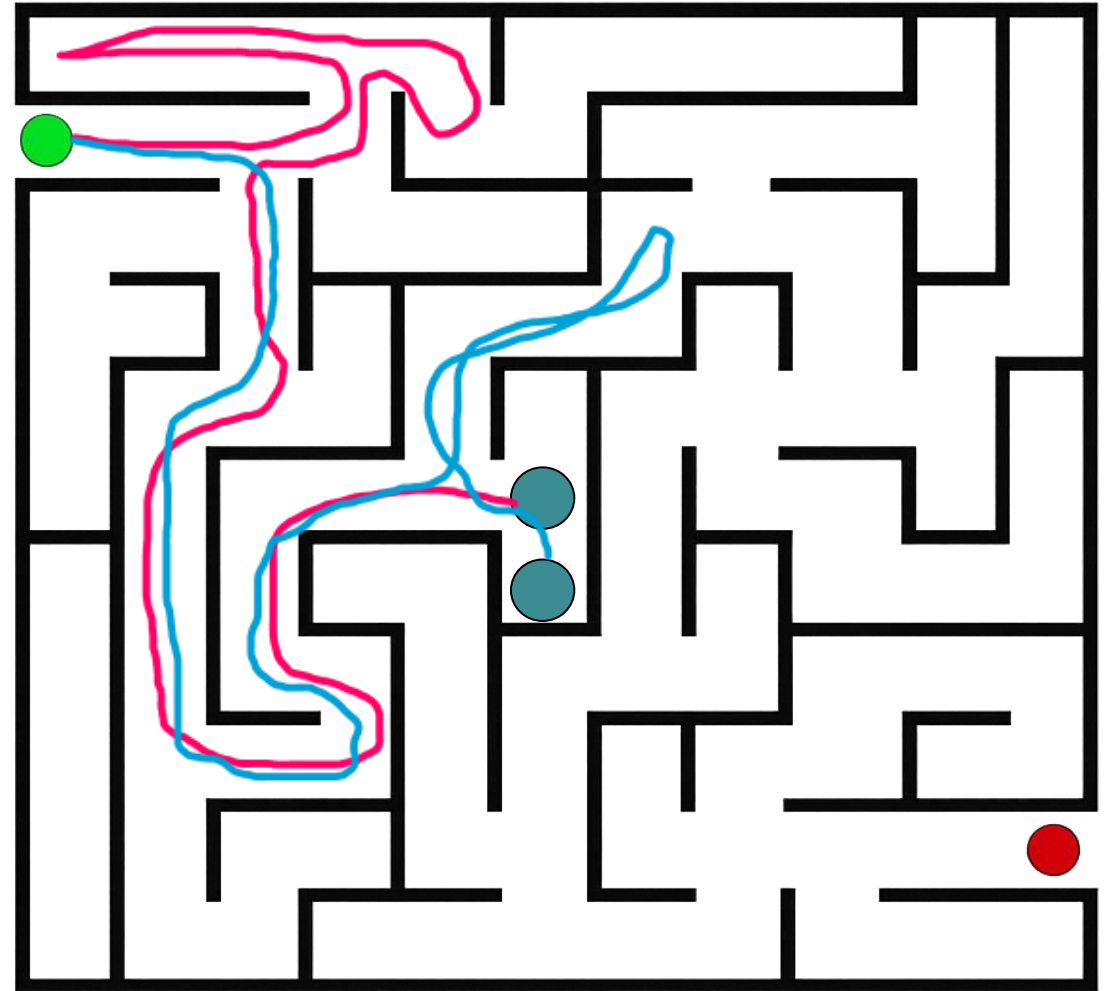
Genotype variation is not enough

Maze problem:

- Robot actions $\{<, ^, v, >\}$

T1 = $\{>, ^, <, >, v, ^, <, v, <, v, <, v, >, ^, <, ^, >\}$

T2 = $\{>, v, <, v, >, ^, <, ^, >, ^, >, ^, v, <, v, >, v\}$



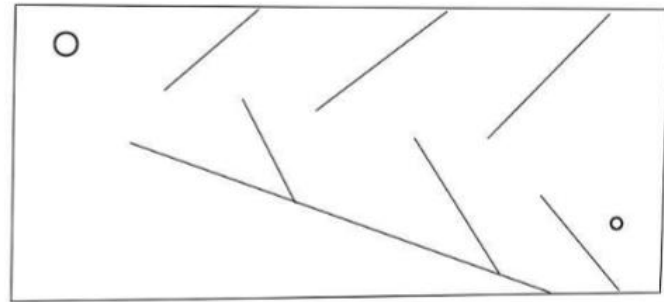
Genotype vs. Phenotype

Premature convergence is the situation where a population loses diversity too quickly and converges to a suboptimal solution (local optimum) before adequately exploring the search space.

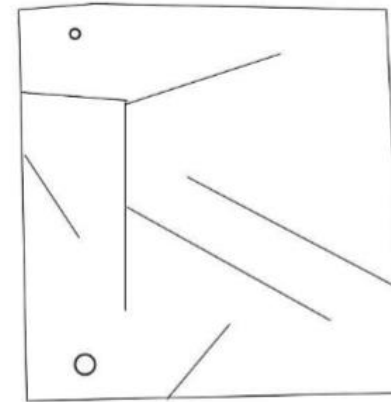
- Reduced spread in fitness (sign of convergence)
- Reduced spread in genotype
 - Adaptive step size (e.g. CMAES)
 - Co-evolve mutation rate
- Reduced spread in phenotype

Deception in Objective-Based Search

- When Fitness is calculated as the distance between the robot's final position and the end goal.
- The more complex the maze, the more deceptive the fitness landscape becomes, trapping standard algorithms in local optima



(a) Medium Map



(b) Hard Map

Lehman, J., & Stanley, K. O. (2008, August). Exploiting open-endedness to solve problems through the search for novelty. In *ALIFE* (Vol. 65, pp. 329-336).

Lehman, J., & Stanley, K. O. (2011). Abandoning objectives: Evolution through the search for novelty alone. *Evolutionary computation*, 19(2), 189-223.

Novelty search

The *novelty* of a newly generated individual is computed with respect to the *behaviors* (i.e. phenotype) of an *archive* of past individuals whose behaviors were highly novel when they originated.

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Novelty search

The *novelty* of a newly generated individual is computed with respect to the *behaviors* (i.e. phenotype) of an *archive* of past individuals whose behaviors were highly novel when they originated.

1. Archive N individuals
2. Measure phenotype distance of each new individual with respect to k-nearest neighbors in the archive
3. If the measure exceeds a threshold, add new individual to the archive

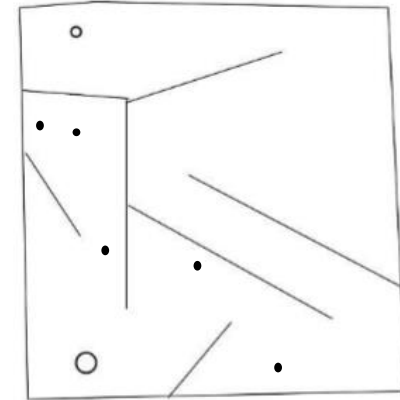
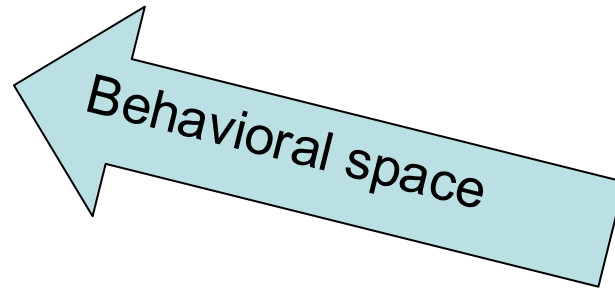
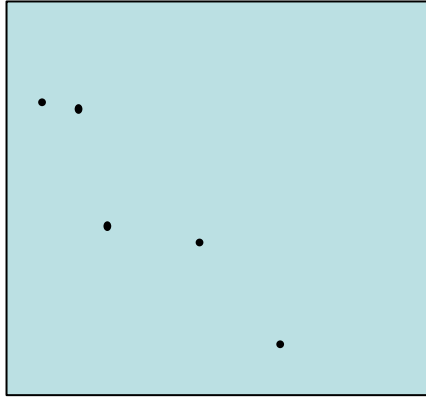
$$\rho(x) = \frac{1}{k} \sum_{i=0}^k \text{dist}(x, \mu_i)$$

Lehman, J., & Stanley, K. O. (2008, August). Exploiting open-endedness to solve problems through the search for novelty. In *ALIFE* (Vol. 65, pp. 329-336).

Lehman, J., & Stanley, K. O. (2011). Abandoning objectives: Evolution through the search for novelty alone. *Evolutionary computation*, 19(2), 189-223.

Novelty search: 1) Archive N individuals

Archive:



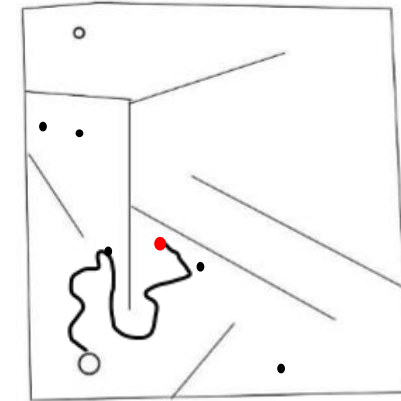
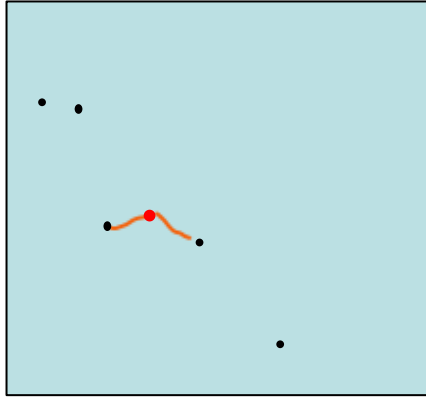
(b) Hard Map

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Lehman, J., & Stanley, K. O. (2011). Abandoning objectives: Evolution through the search for novelty alone. *Evolutionary computation*, 19(2), 189-223.

Novelty search: 2) Measure phenotype distance

Archive:



(b) Hard Map

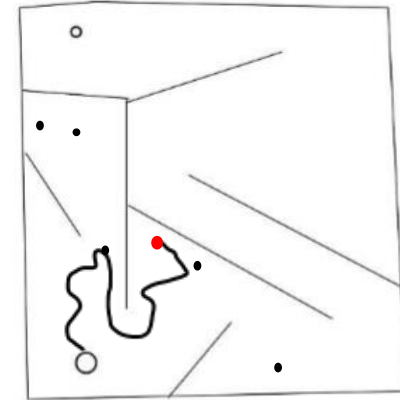
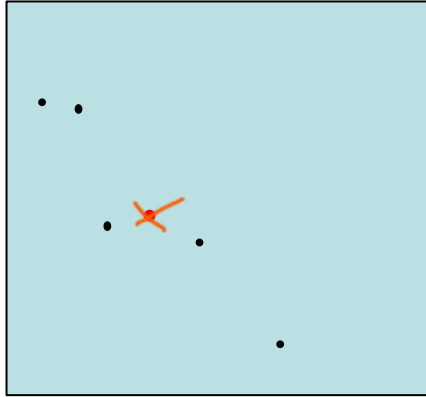
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Novelty search: 3) if measure exceeds threshold..

Archive:



(b) Hard Map

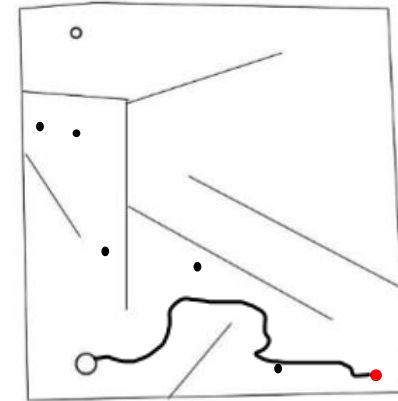
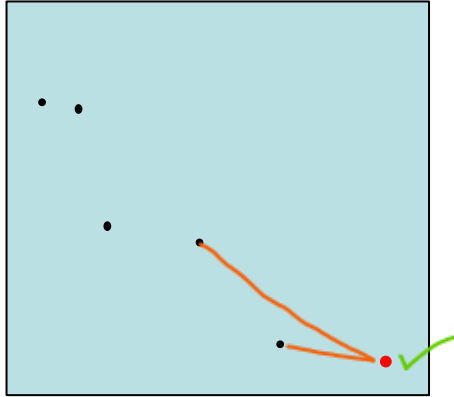
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Novelty search: 3) if measure exceeds threshold..

Archive:



(b) Hard Map

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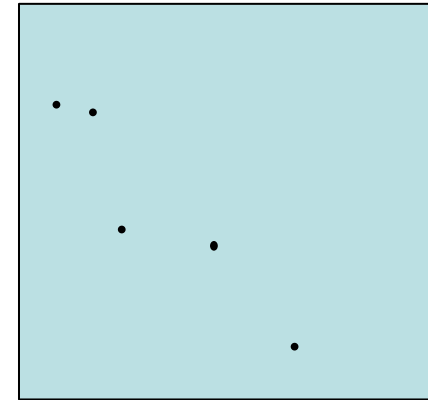
Lehman, J., & Stanley, K. O. (2011). Abandoning objectives: Evolution through the search for novelty alone. *Evolutionary computation*, 19(2), 189-223.

New objective: fill archive as much as possible

Instead of goal-based optimization that can get trapped

We fill the behavioral space by optimizing for novelty

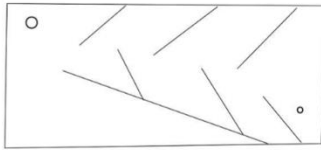
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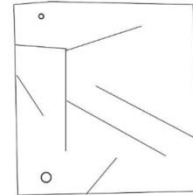
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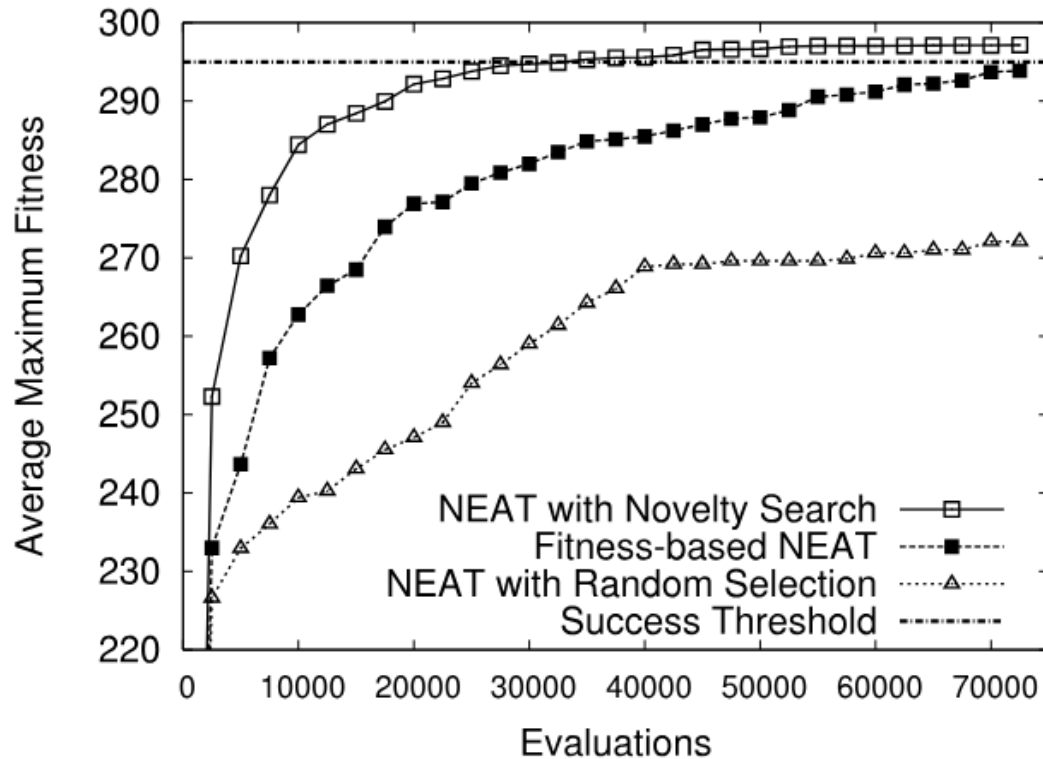
Novelty search outperforms distance-based fitness



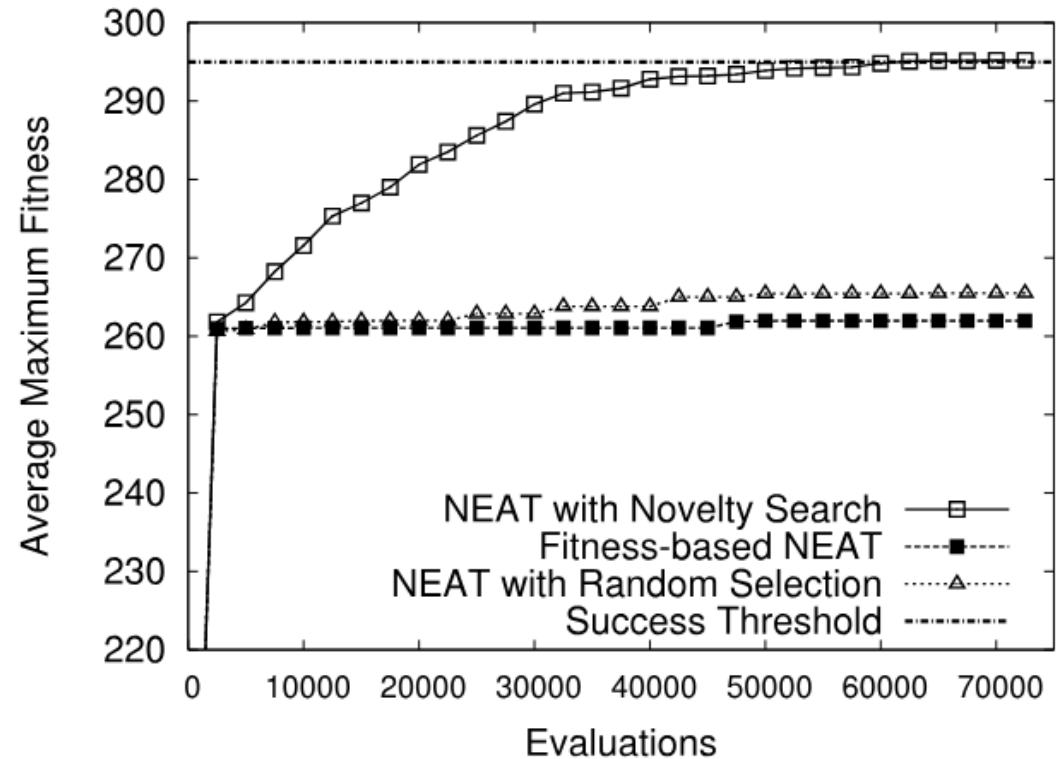
(a) Medium Map



(b) Hard Map



(a) Medium Map



(b) Hard Map

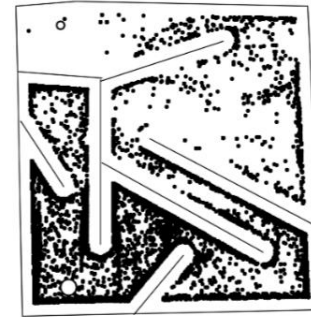
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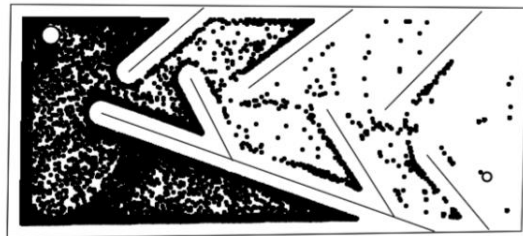
Novelty search completes hard maze



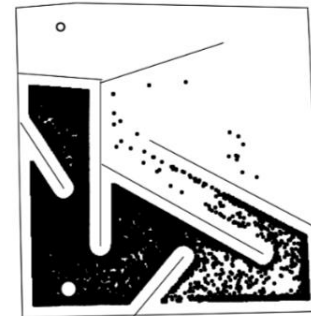
(a) Medium Map Novelty



(b) Hard Map Novelty



(c) Medium Map Fitness



(d) Hard Map Fitness

Lehman, J., & Stanley, K. O. (2008, August). Exploiting open-endedness to solve problems through the search for novelty. In *ALIFE* (Vol. 65, pp. 329-336).

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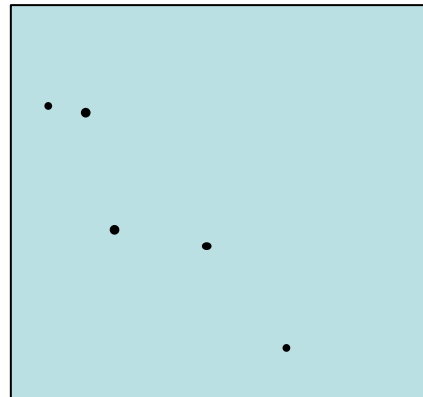
Novelty search: key takeaways

- Instead of optimizing for an objective, Novelty Search rewards unique behaviors.
- The algorithm bypasses deceptive local optima, by optimizing phenotypical variation.

Open-ended evolution

- Premature convergence is caused by focusing on an objective.

Open-ended evolution redefines optimization as a never-ending process, rather than settling into a stable equilibrium.

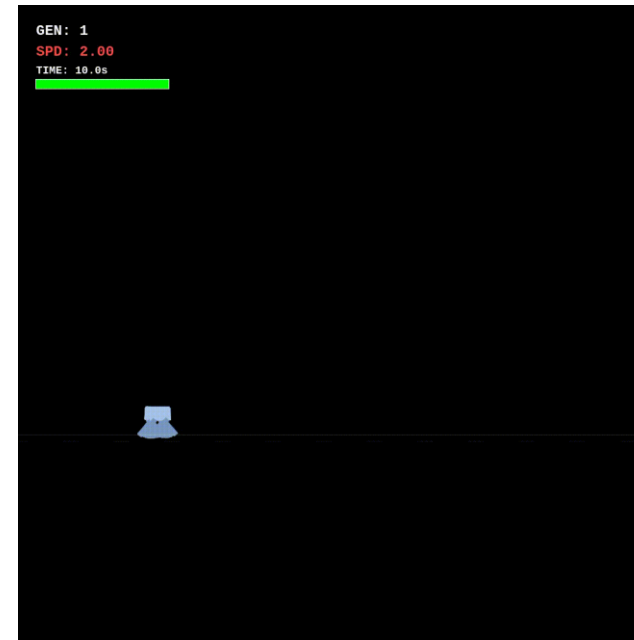


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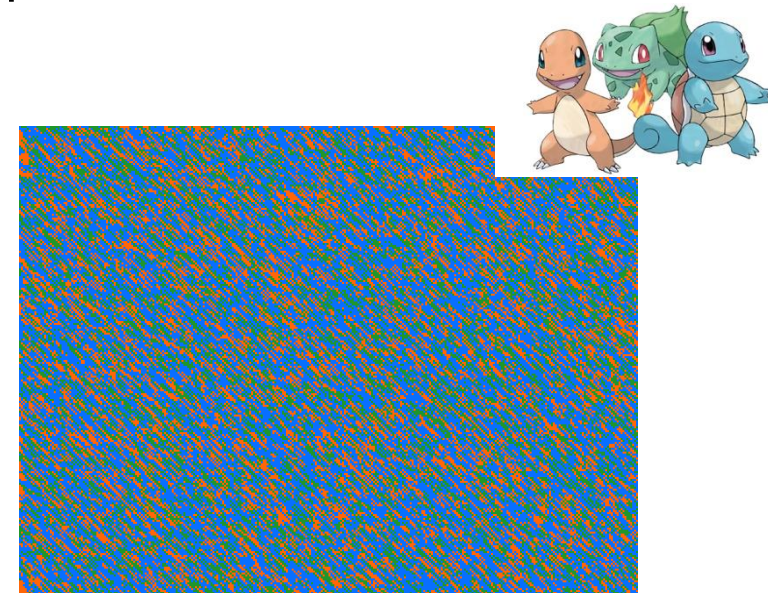
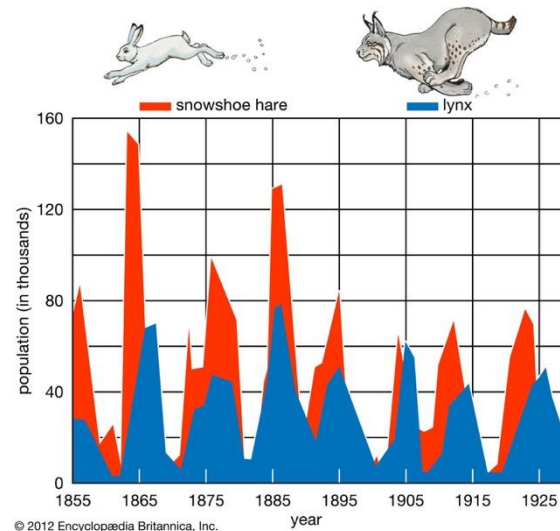


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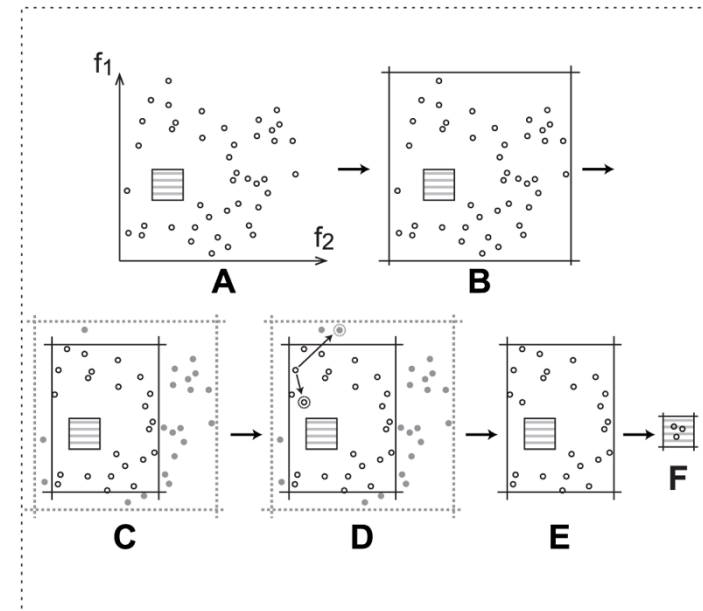


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- Viability Evolution



Open-ended evolution

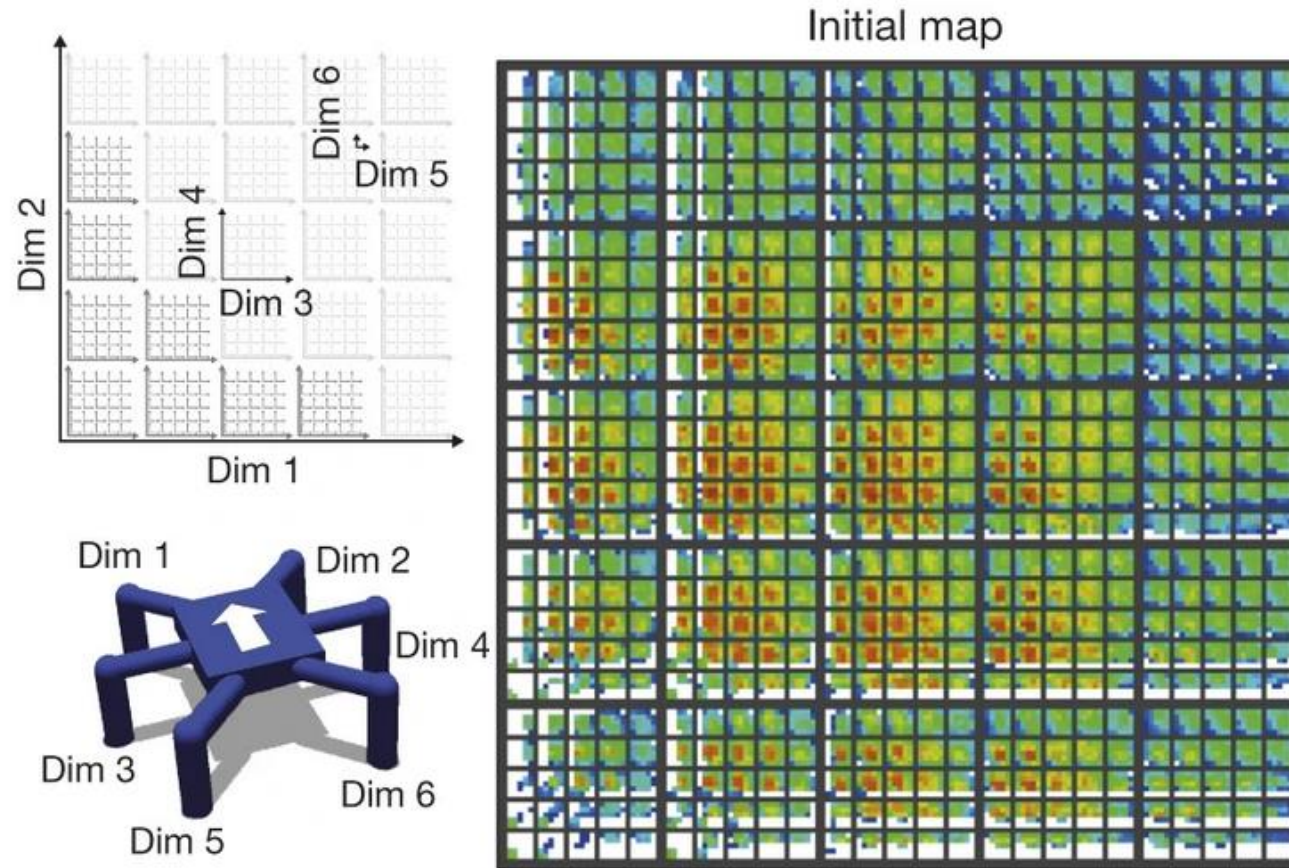
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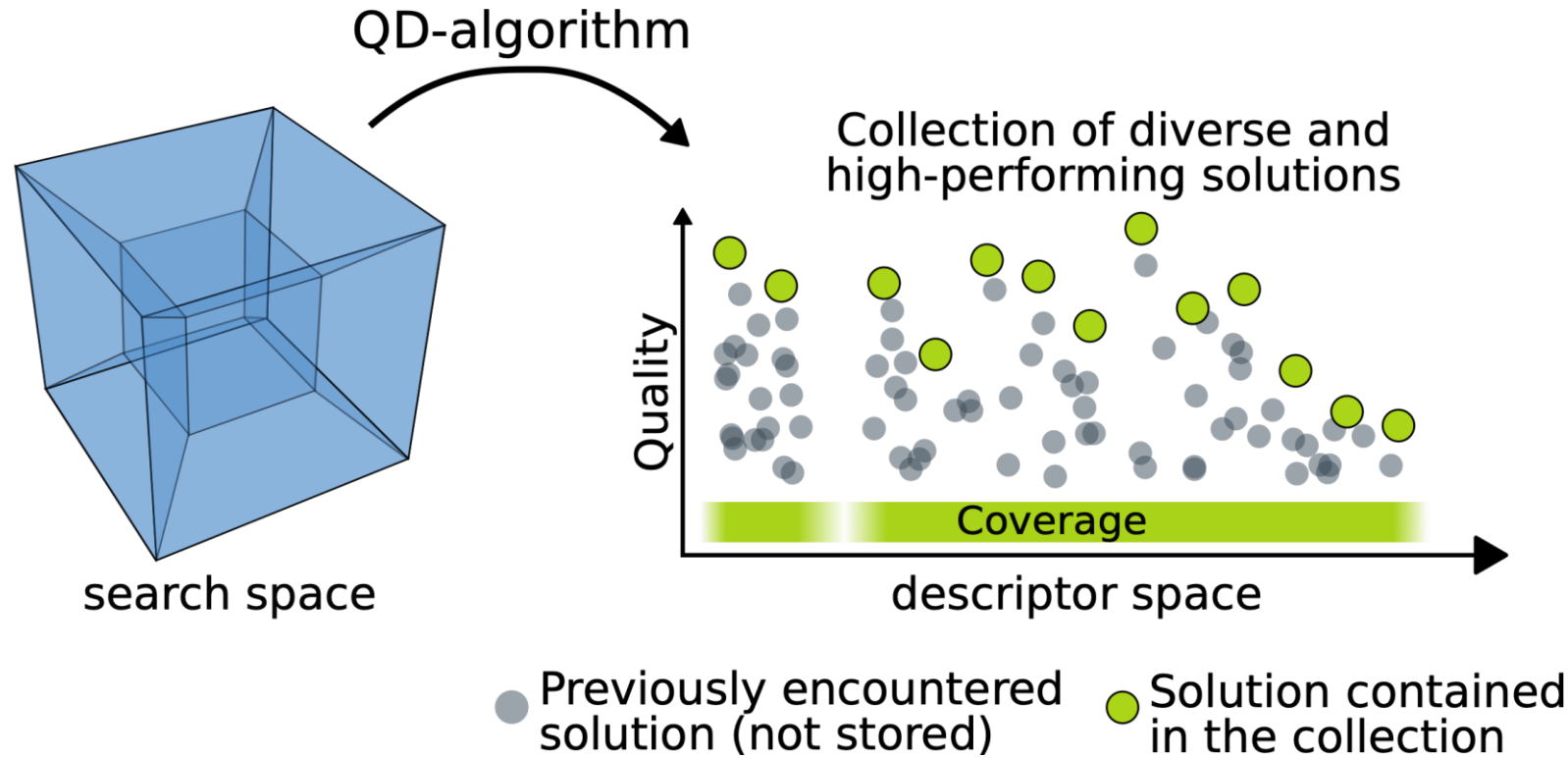
Diversity does not guarantee quality!

MAP Elites



Premature convergence in robot co-design

The co-design of control/body evolution results in different performances for similar designs, thereby worsening premature convergence

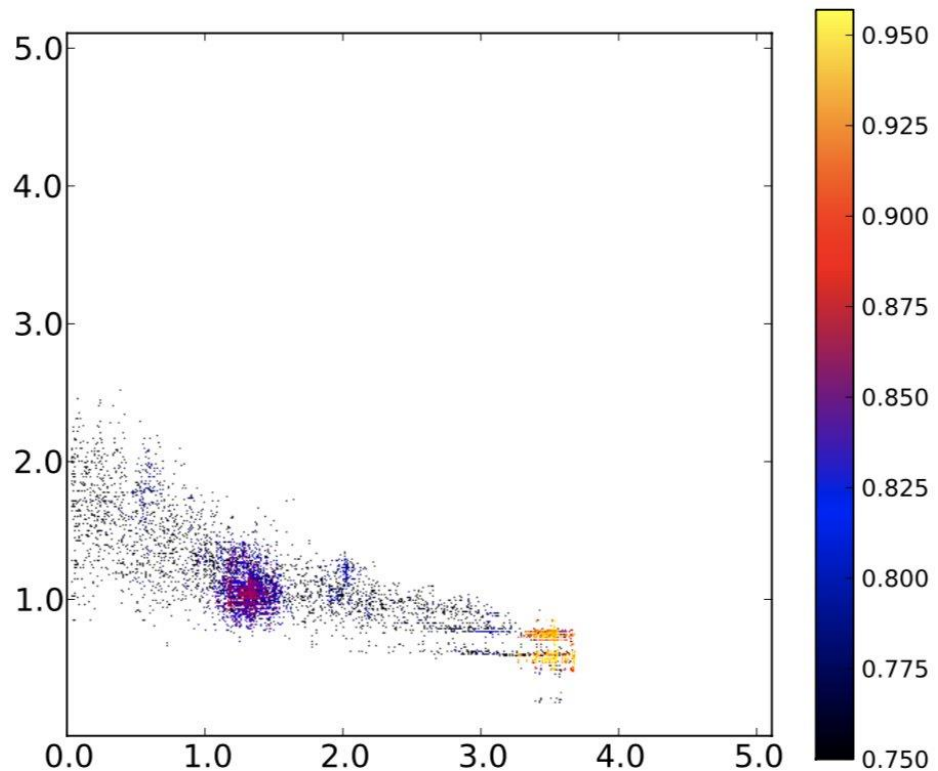


Quality Diversity algorithms

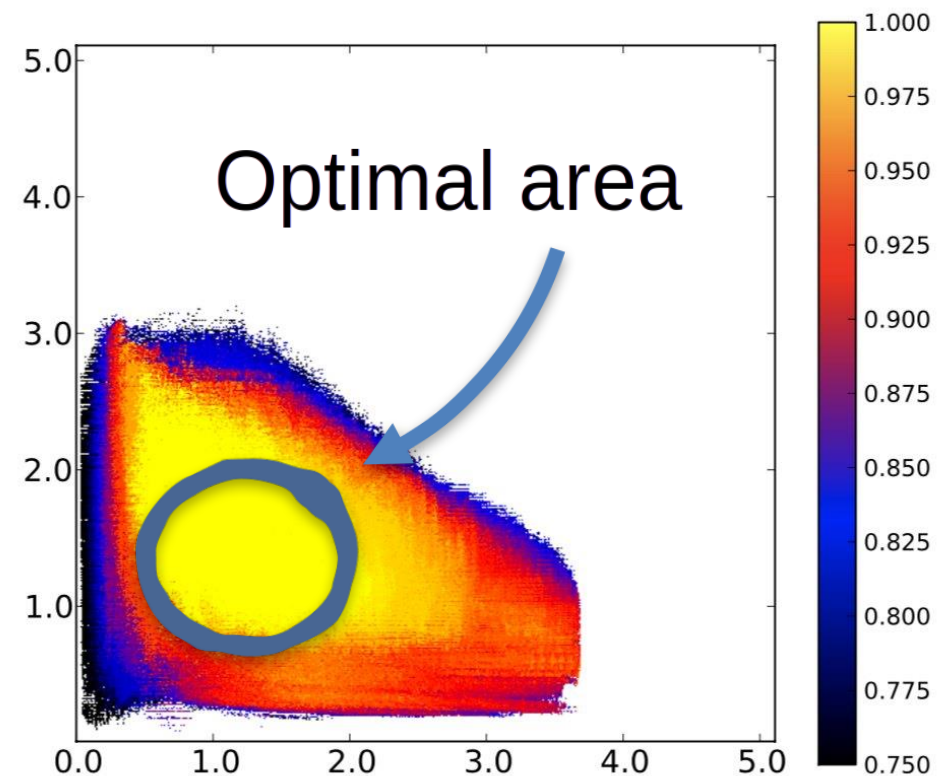
The co-design of control/body evolution results in different performances for similar designs, thereby worsening premature convergence

Quality-Diversity: Optimize both goal (quality) and phenotypical coverage (diversity)

Novelty search -> Maximize differences



MAP-Elites -> Optimize within niche



MAP Elites Archive

Define a phenotypical feature space (Archive)

Discretize feature space

Test individual fitness

Place individual in phenotype grid

- If individual fitness > current cell best; **replace**

procedure MAP-ELITES ALGORITHM (SIMPLE, DEFAULT VERSION)

$(\mathcal{P} \leftarrow \emptyset, \mathcal{X} \leftarrow \emptyset)$

for iter = 1 \rightarrow I **do**

if iter < G **then**

$\mathbf{x}' \leftarrow \text{random_solution}()$

else

$\mathbf{x} \leftarrow \text{random_selection}(\mathcal{X})$

$\mathbf{x}' \leftarrow \text{random_variation}(\mathbf{x})$

$\mathbf{b}' \leftarrow \text{feature_descriptor}(\mathbf{x}')$

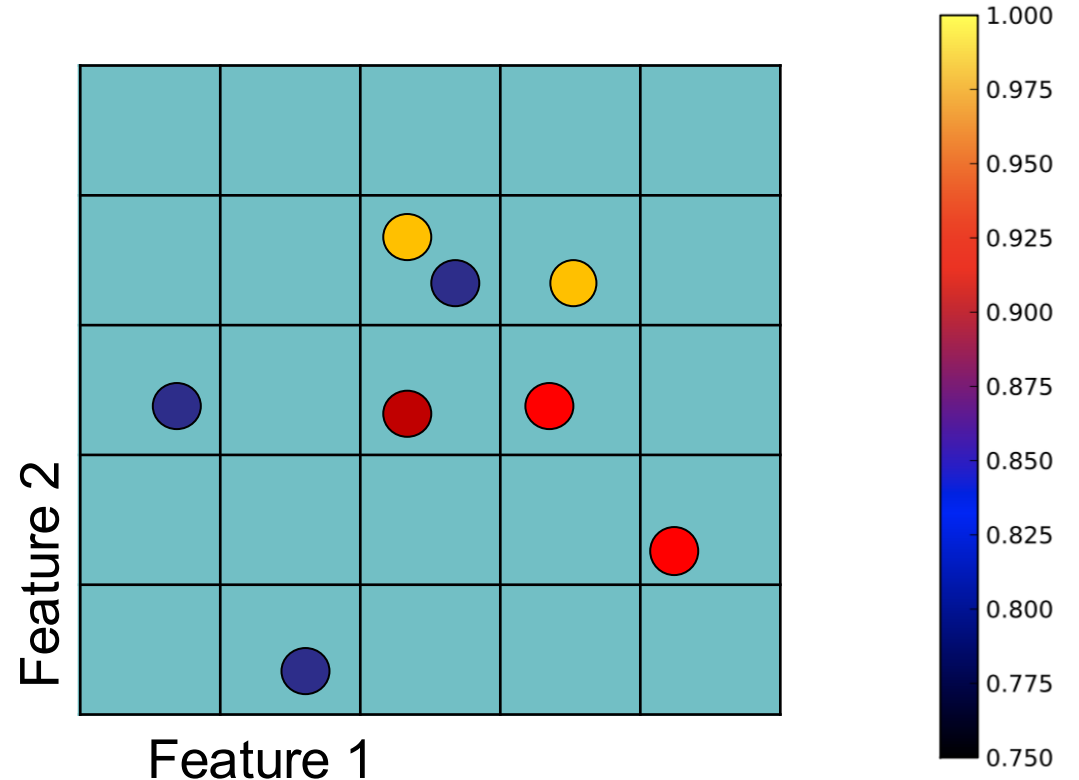
$p' \leftarrow \text{performance}(\mathbf{x}')$

if $\mathcal{P}(\mathbf{b}') = \emptyset$ or $\mathcal{P}(\mathbf{b}') < p'$ **then**

$\mathcal{P}(\mathbf{b}') \leftarrow p'$

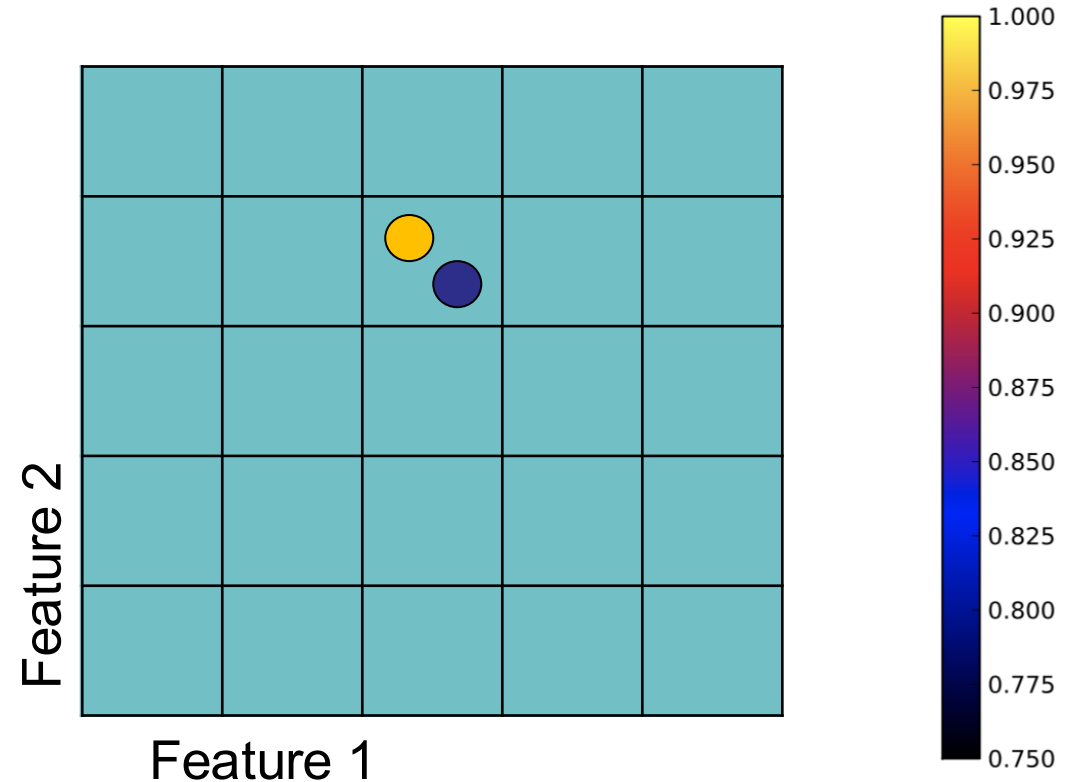
$\mathcal{X}(\mathbf{b}') \leftarrow \mathbf{x}'$

return feature-performance map (\mathcal{P} and \mathcal{X})



Niche construction in the archive

- MAP-Elites forces similar solutions to compete only with each other.
- Each map cell preserves a local elite, thus allowing for Phenotypical diversity



MAP Elites: soft-voxel robots

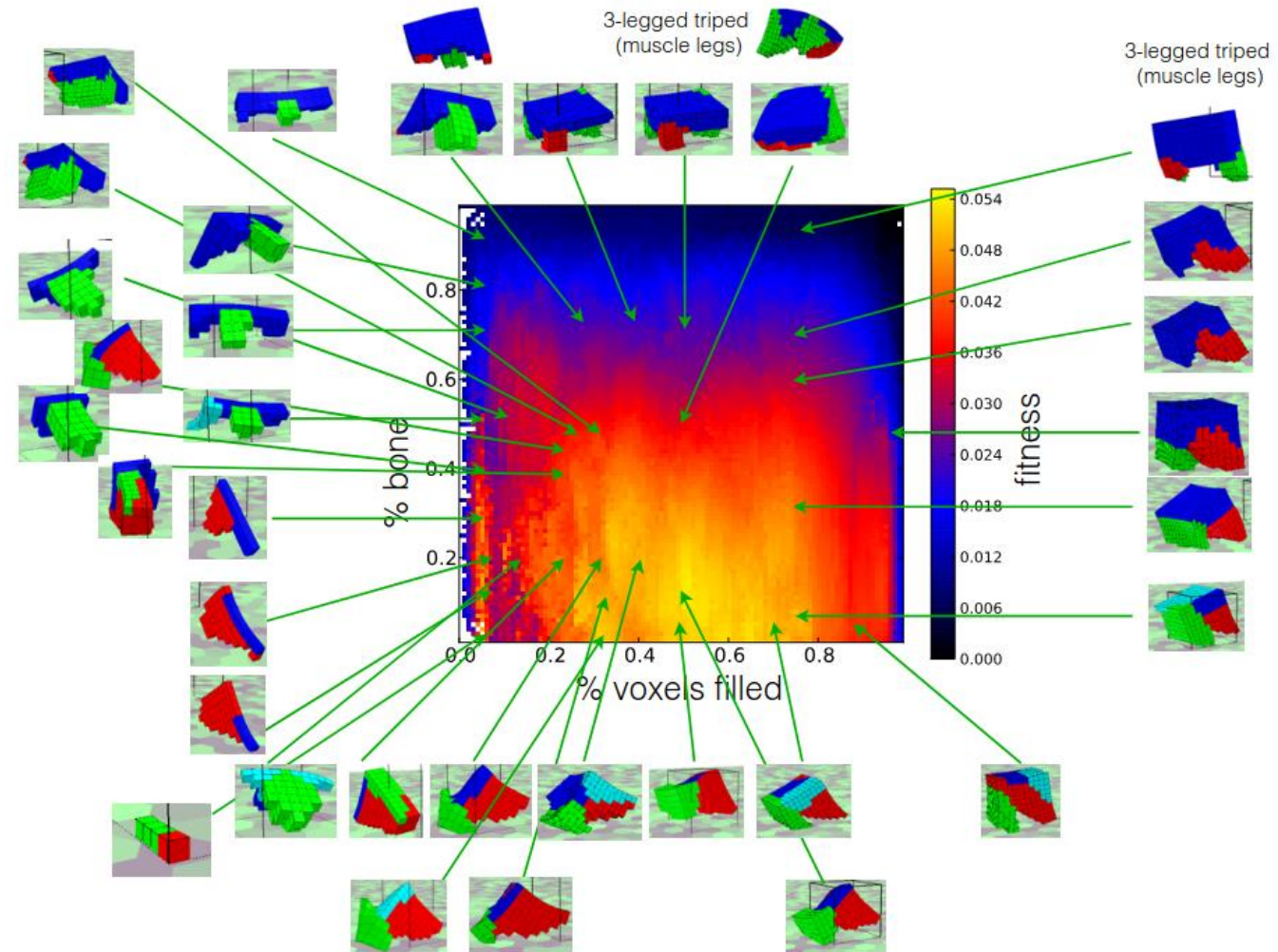
Example:

Evolve soft voxel robots walk as far as possible

Voxels: {empty, bone, soft, contract, expand}

Phenotype space:

- F1: Voxels not empty [%]
- F2: Voxels = bones [%]



MAP Elites: 6-legged robot

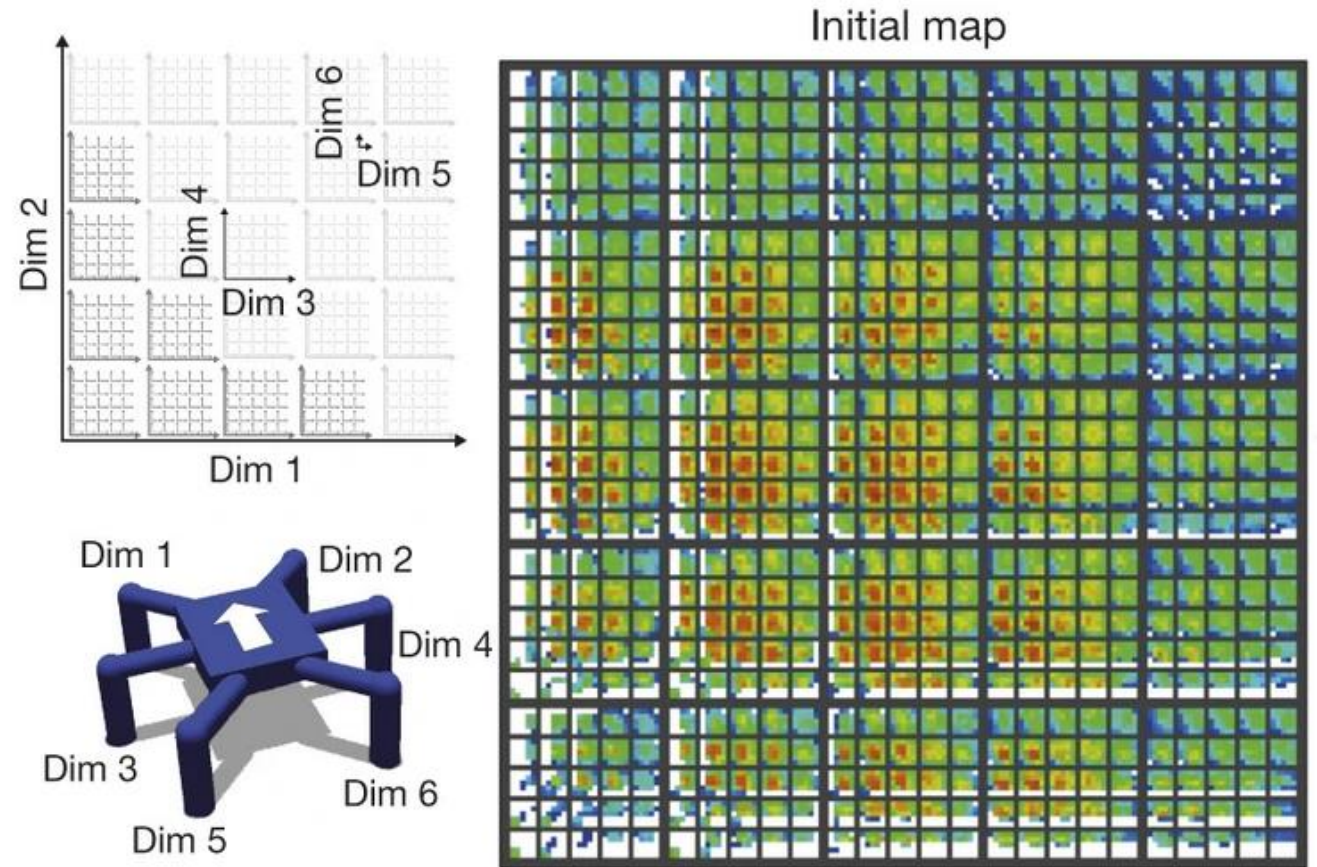
Example:

Evolve 6-legged robot walk as far as possible 5s

Parameterized controller

Phenotype space Contact time per leg:

- F1: Dim 1 (leg 1)
- F2: Dim 2
- ...
- F6: Dim 6

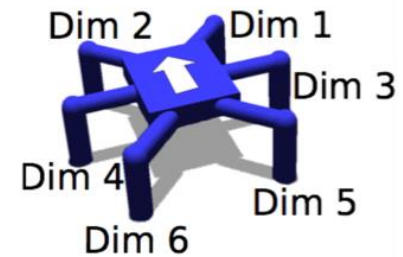
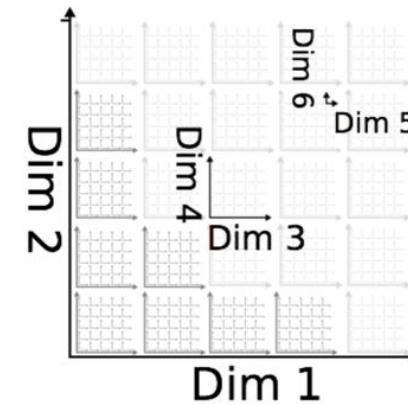
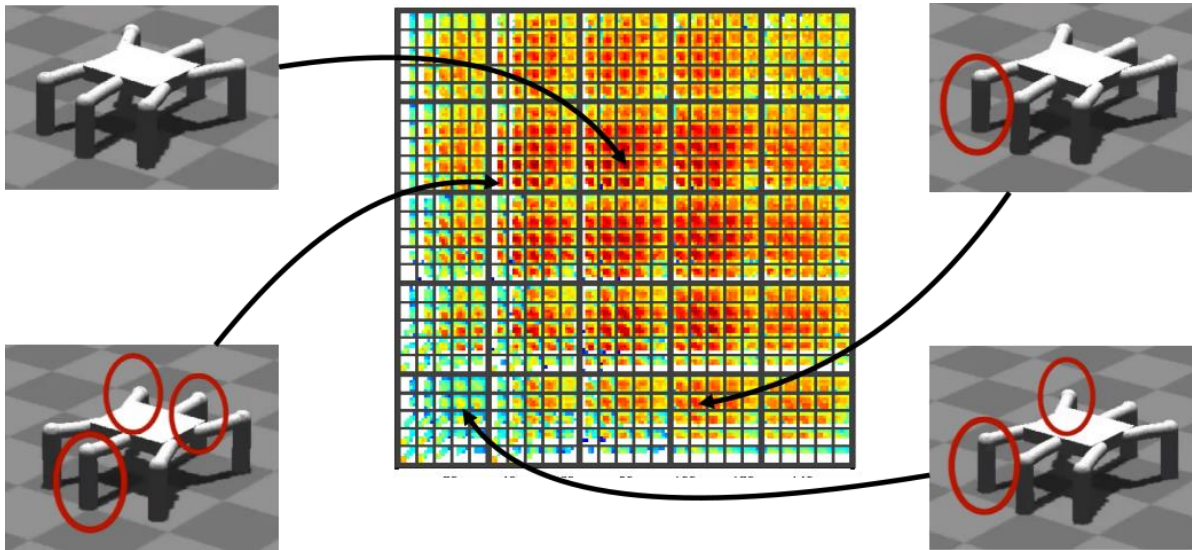


Understanding the feature space

Analyzing feature space (time of ground contact)

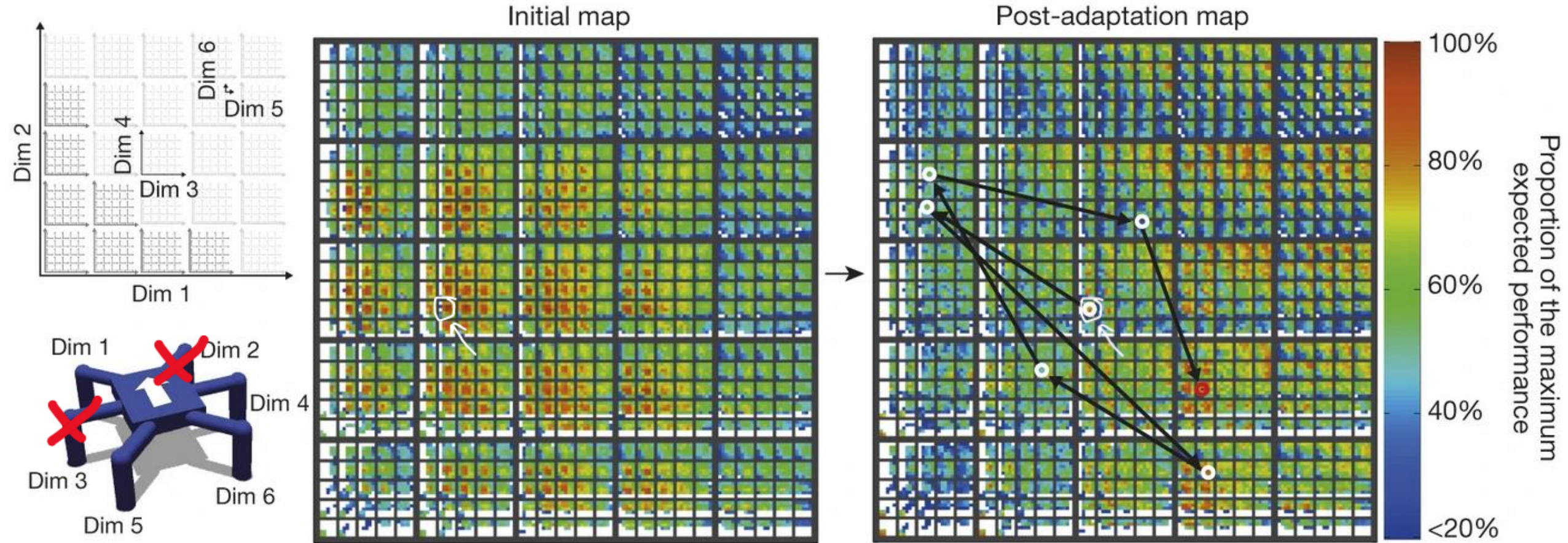
E.G. right bottom robot

Dim 1 & Dim 2: low



Adaptation through archival motor primitives

Learning a wide range of behaviors allows for quick adaptation



Cully, A., Clune, J., Tarapore, D., & Mouret, J. B. (2015). Robots that can adapt like animals. *Nature*, 521(7553), 503-507.

Real world test

Learning a wide range of behaviors allows for quick adaptation

Robots that can adapt like animals

Nature, 2015

which describes damage recovery via Intelligent Trial and Error



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University of Wyoming
(USA)



Danesh Tarapore
UPMC/CNRS
(France)



Jean-Baptiste Mouret
UPMC/CNRS/Inria/UL
(France)



Cully, A., Clune, J., Tarapore, D., & Mouret, J. B. (2015). Robots that can adapt like animals. *Nature*, 521(7553), 503-507.

MAP-Elites: key takeaways

- MAP-elites Optimize both goal (quality) and phenotypical coverage (diversity)
- Archive is a phenotype-performance space
- Unlike novelty-search MAP-elites do not reject similar phenotypes; but replaces individuals when performance improves (i.e. local competition/niches)

Checkpoints

- Describe premature convergence
- Describe the differences between genotype/phenotype diversity
- What is open-ended evolution and how does it differ from goal-based evolution
- Describe the following Quality-Diversity algorithms:
 - Novelty search
 - MAP-Elites

