# Framsticks genetic representations comparison experiments

Maciej Komosinski and Adam Rotaru-Varga. Comparison of dierent genotype encodings for simulated 3D agents. *Articial Life Journal*, 7(4):395-418, Fall 2001. [view pdf]

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#### Goals

- co-evolve bodies and brains
- design various methods of description of body and brain
- study and compare the effectiveness of evolution using these methods
- <u>in a single system.</u>

#### Why to co-evolve brains and bodies?

- because it yields better results than with body separated from brain
- because it is natural
- embodiment: physical interactions (between body parts, signal processing) perform computations, a part of overall behavior
- brain and body strongly connected
  - evolution of body changes the cognitive space of the brain (e.g.: an eye placed on a limb, new senses)
  - evolution of brain changes usage of the body
- co-evolution: can cause change even in the absence of environmental change

# What is the trouble?

- the 'matching' problem
  - parts of brain (neurons, nodes) must be connected to parts of body (sensors, actuators)
  - if matching is explicit, it can be disrupted by the change of either side, which can be catastrophic
- both are variable size
- crossover on complex representations



# General problems in optimization of realistic autonomous agents

- Infinite search space
- Discrete-continuous space
- Hard to define neighborhood
- Solutions contain varying amount of information
- Hard to choose representation
- Very strong dependencies and connections between parts of a solution
- Evaluation function with many local optima
- Many non-feasible solutions and diverse constraints
- Non-determinism and complexity of evaluation
- Multi-criteria evaluation, complex definition of criteria, evaluation delayed to action
- Hard to estimate the time needed for evaluation and optimization

...the big problem is size and nature of the search space

#### Genetics is important because each representation and its operators

- establishes different structure and order in the search space
- defines important information and 'building blocks' in another way
- is scalable in a different degree
- introduces different bias which leads to finding qualitatively different solutions
- imposes diverse local optima and displays various levels of robustness against being trapped into them
- can limit the space of valid solutions in a particular way
- has a specific degree of coherency, redundancy, easiness of interpretation, etc.

# Current artificial genomes – very diverse



No comparison possible!

What is the best representation?

- •Fitness values
- •Nature of solutions
- •Simplicity
- •Understandability

# Experiments

- 3 one-criterion tasks
  - Average height of agent center (maximize; NN turned off)
  - Average height of agent center (maximize; NN turned on)
  - Average velocity (maximize)
- 10 runs for each task and each genotype format (*simul*, *recur*, *devel*)
- 90 runs in total
- System main parameters
  - Steady-state
  - Population size: 200
  - Cloning probability: 20%
  - Crossing over prob.: 16%
  - Mutation prob.: 64%
  - Stabilization period



# Non-deterministic evaluation



#### Results (Quantitative)



#### Genetics experiment

#### Results (Qualitative; height passive)



# Conclusions

- *Simul* representation with full abilities of expressing agents was the worst one
- The limitation of the search space by higher level representations has not deteriorate results, but has improved them
- The most advanced *devel* encoding was not significantly better than *recur*
- Each higher-level representation introduces a specific bias and new quality (characteristics) into solutions
- For all representations, the best individuals were successful in terms of fitness value. It was difficult or impossible to construct better agents by hand, mainly because of high time costs
- It may be sometimes worthwhile to introduce advanced mechanisms into a representation, in order to obtain different nature of solutions, even when they are not improved in terms of fitness

#### Genetics experiment

# Conclusions, cont.

- punctuated equilibria
- convergence
- exploitation of simulator imperfections
- redundancy, randomness
- many strong (implicit) dependencies inside agents

