

Evolutionary design for the behaviour of cellular automaton-based complex systems

Peter Siepmann, Germán Terrazas, Natalio Krasnogor

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Adaptive Computing in Design and Manufacture
25th April 2006
Bristol

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Motivation

- automated design and optimisation of complex systems
 - cellular automata models
- how can we verify that our fitness function is robust?
- present a methodology for answering this question
 - supported by experimental illustration

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Introduction

- Self-organising processes
- Modelled using **cellular automata**

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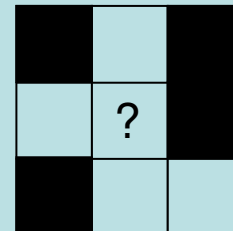
Introduction

- Self-organising processes
- Modelled using **cellular automata**

- infinite, regular grid of cells
- each cell in one of a finite number of states
- at a given time, t , the state of a cell is a function of the states of its *neighbourhood* at time $t-1$.

Example

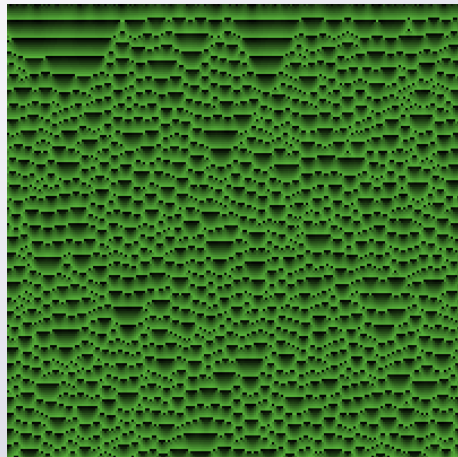
- infinite sheet of graph paper
- each square is either black or white
- in this case, neighbours of a cell are the eight squares touching it
- for each of the 2^8 possible patterns, a rules table would state whether the center cell will be black or white on the next time step.



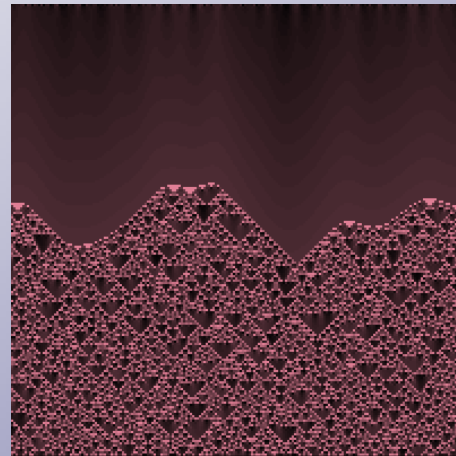
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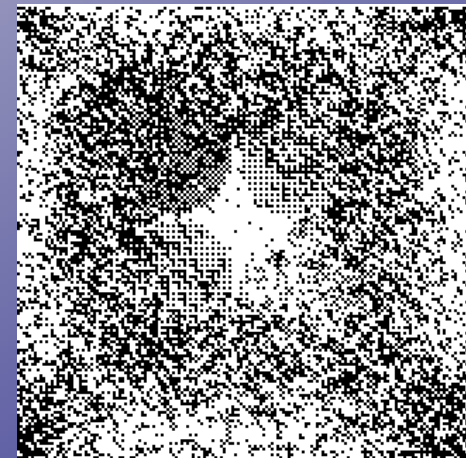
Introduction



CA continuous



Turbulence



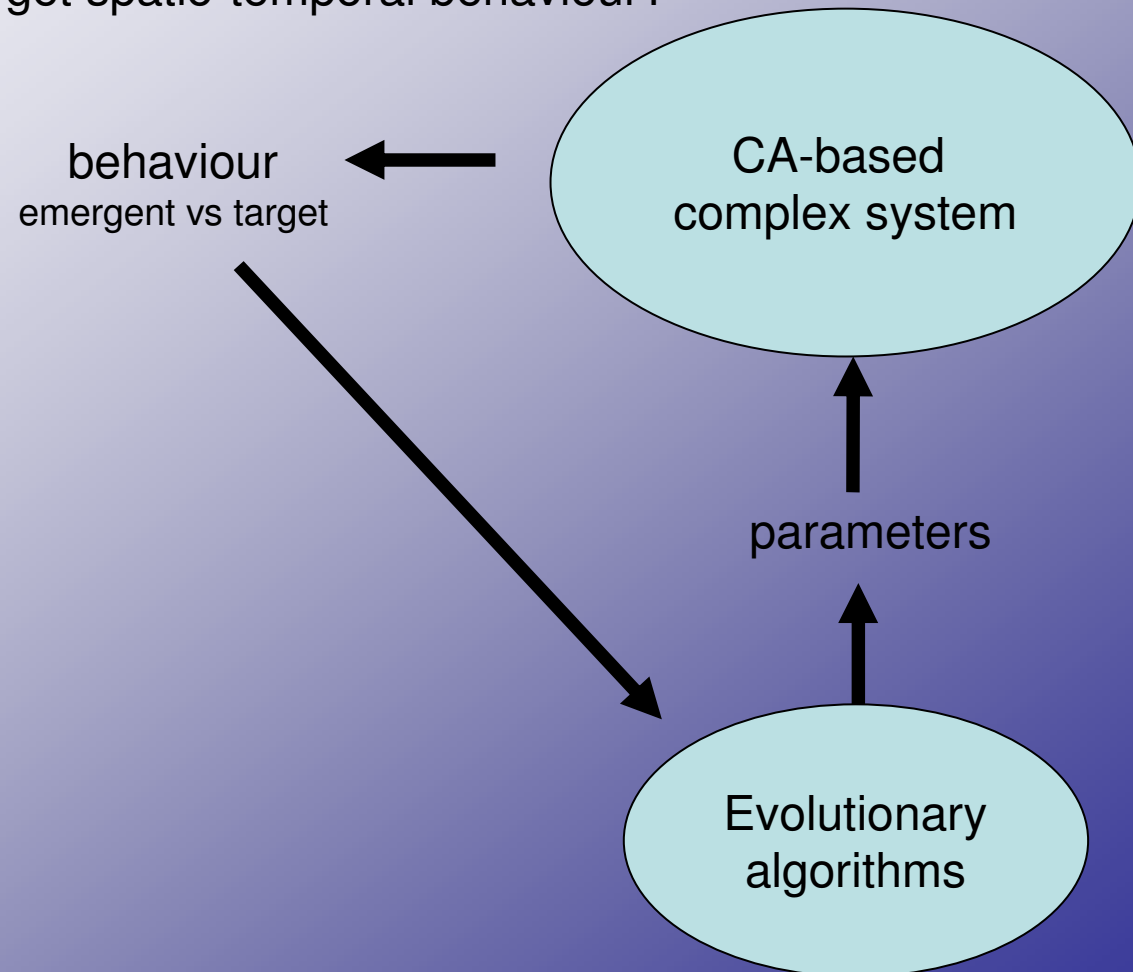
Gas Lattice

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Introduction

- Can we generate a target spatio-temporal behaviour?

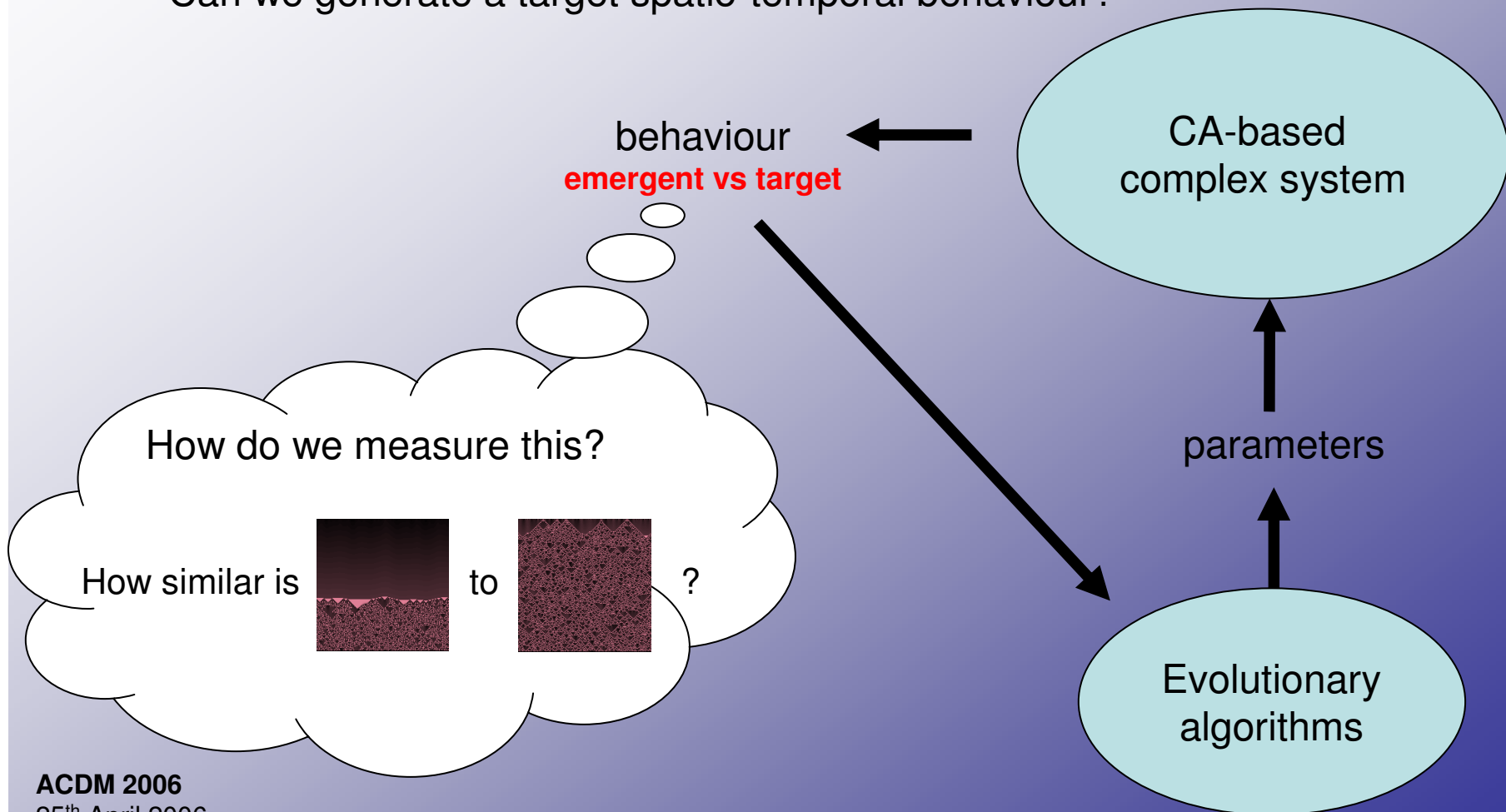


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The Universal Similarity Metric (USM)

is a measure of similarity between two given objects in terms of information distance:

$$d(o_1, o_2) = \frac{\max\{K(o_1 | o_2), K(o_2 | o_1)\}}{\max\{K(o_1), K(o_2)\}}$$

where $K(o)$ is the Kolmogorov complexity

Prior Kolmogorov complexity $K(o)$: The length of the shortest program for computing o by a Turing machine

Conditional Kolmogorov complexity $K(o_1|o_2)$:

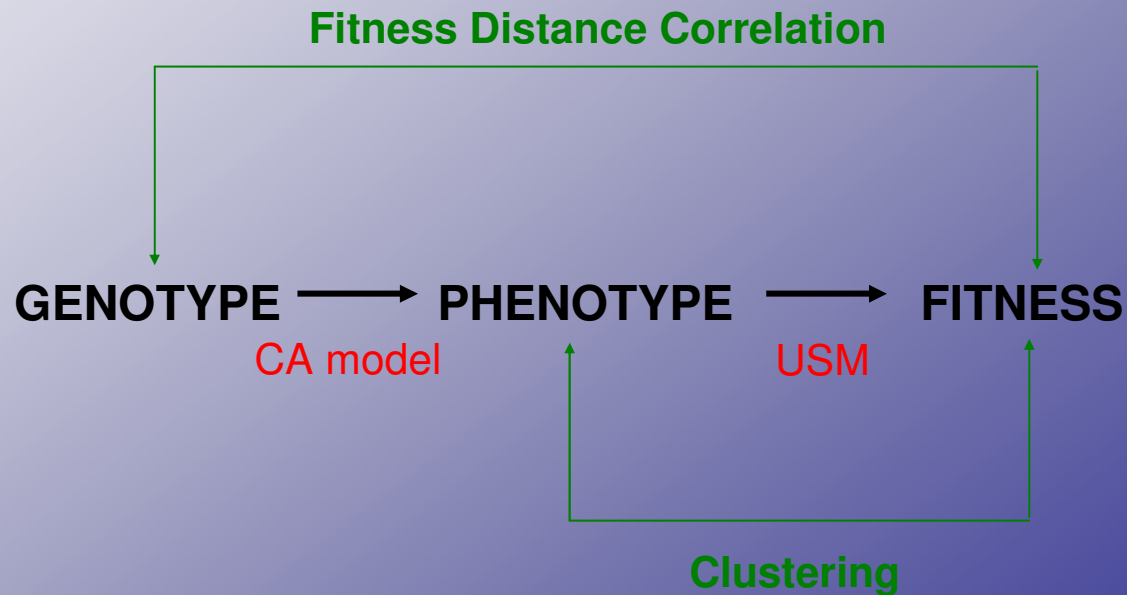
How much (more) information is needed to produce object o_1 if one already knows object o_2 (as input)

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The Universal Similarity Metric (USM)

- Is the USM a good objective function for evolving target spacio-temporal behaviour in a CA system?
- methodology for answering this question
- experimental results



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Data set

For each CA system:

- Keep all but one parameter the same
- Produce 10 behaviour patterns through the variable parameter
- Repeat for other parameters

EXAMPLE

`turb_c4` refers to the spacio-temporal pattern produced by the fourth variation in parameter *c* of a *Turbulence CA* system

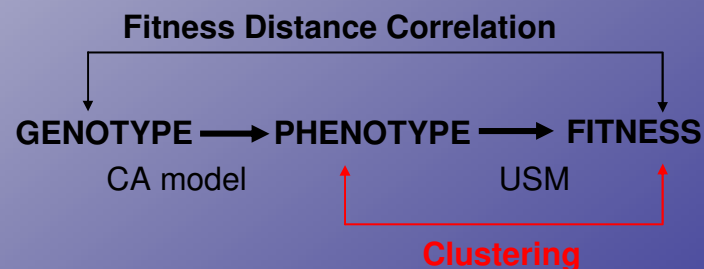
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Clustering

- does the USM detect similarity of phenotype?
- calculate a similarity matrix filled with the results of the application of the USM to a set of objects
- during the clustering process, similar objects should be grouped together





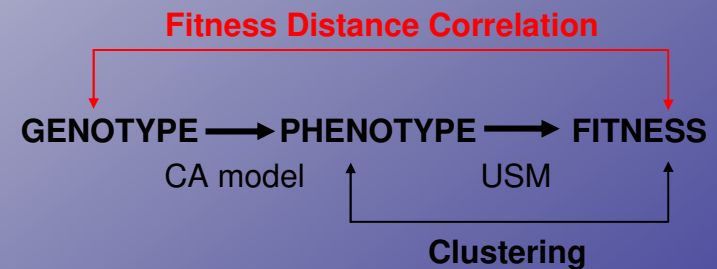
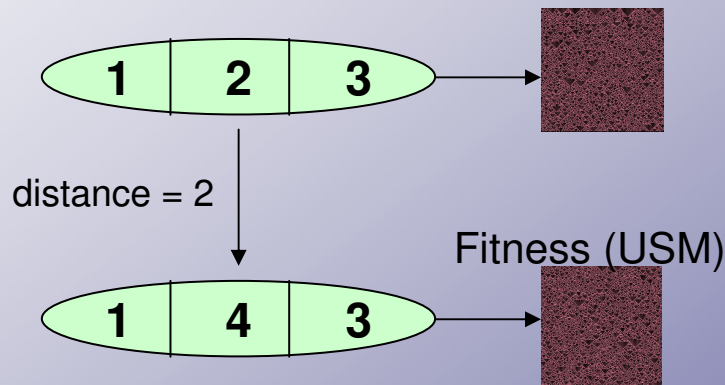


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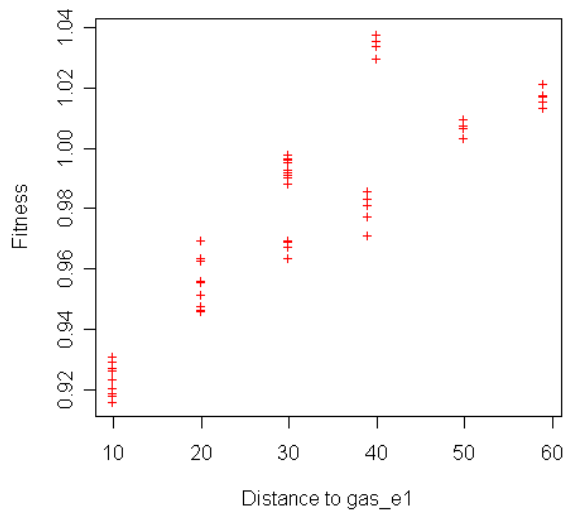
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Fitness Distance Correlation

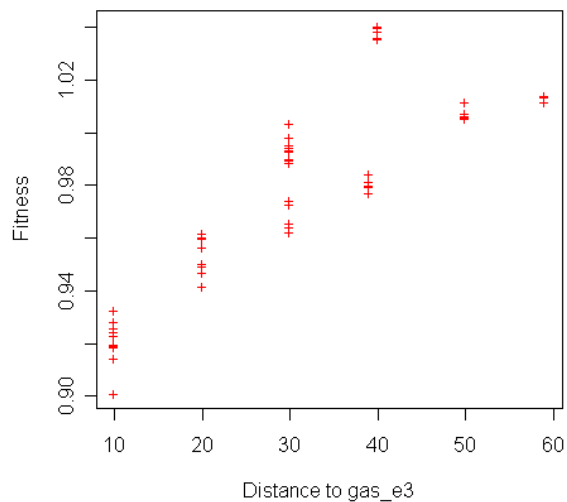
- correlation analyses of a given fitness function versus parametric (genotype) distance.
- larger numbers indicate the problem could be optimised by a GA
- numbers around zero $[-0.15, 0.15]$ indicate bad correlation
- scatter plots are helpful



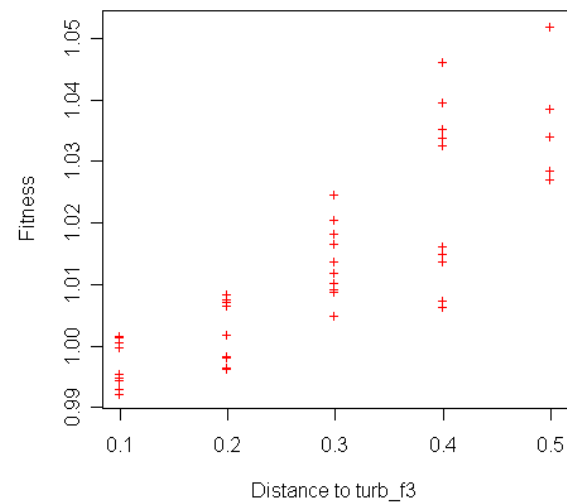
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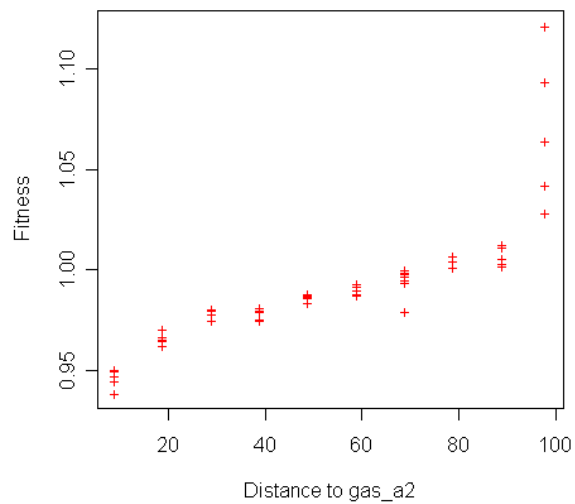
FDC = 0.473056717160691



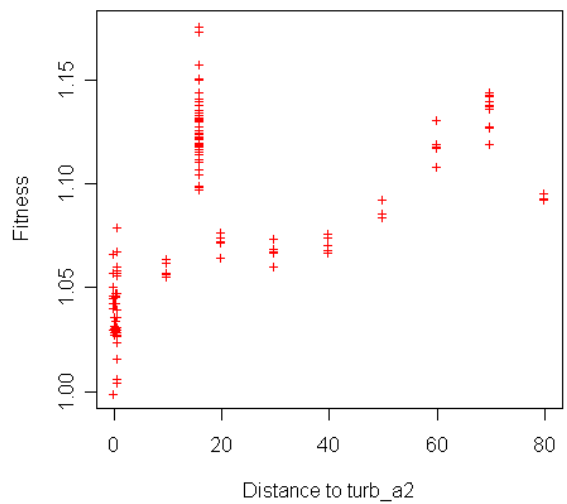
FDC = 0.330265456807397



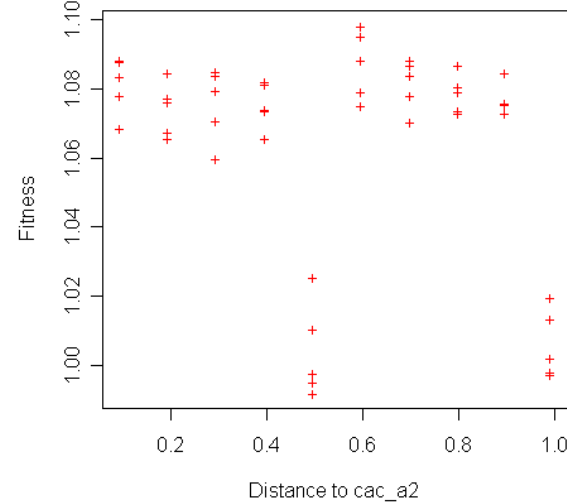
FDC = 0.49942809674373



FDC = 0.297666084271867



FDC = 0.264936105455606



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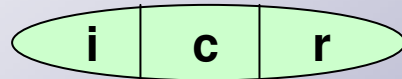
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The Genetic Algorithm

Aim: to evolve a behaviour as similar as possible to some given target behaviour

Parameters

Chromosome:



Population size: 10

Offspring: 4

Selection: roulette

Recombination: uniform crossover (probability 1)

Mutation: BCG (probability 0.6)

selects a value in a fixed distribution
either side of current value

Stopping condition: 200 generations

$i \in \mathcal{R} . 0 \leq i \leq 100$

initial turbulence

$c \in \mathcal{R} . 0 \leq c \leq 1$

coupling strength

$r \in \mathcal{R} . 0 \leq r \leq 0.025$

roughness

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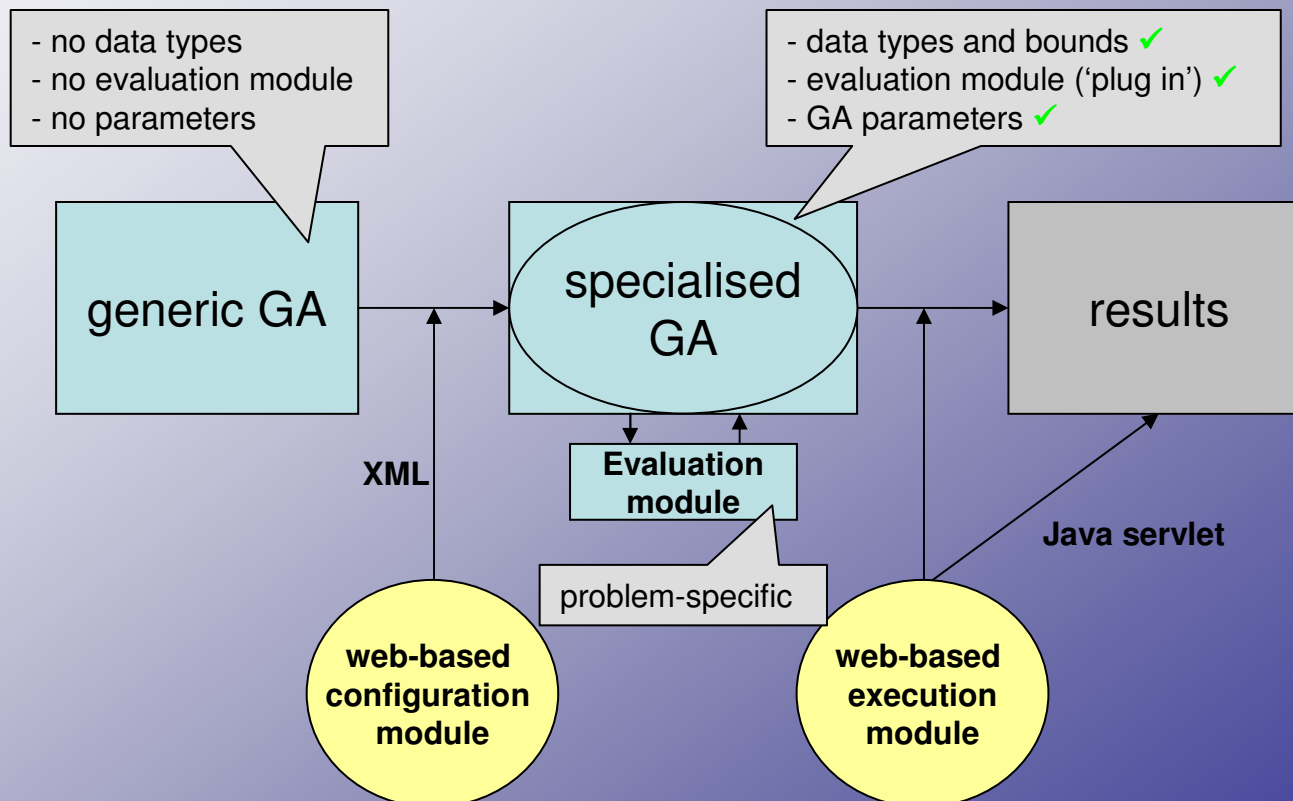
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The Genetic Algorithm

Our GA platform

“we will implement an object-oriented, platform-independent, evolutionary engine (EE). The EE will have a user-friendly interface that will allow the various platform users to specify the platform with which the EE will interact”

Evolvable CHELLware grant application



The Genetic Algorithm

Measuring success

- Visual comparison
- USM
- Genotypic error

We define the error for each gene, $e(g) = \frac{abs(g_T - g_F)}{g_T}$

and the *average error* for a given individual, $E(F) = \frac{e(i_F) + e(c_F) + e(r_F)}{3}$

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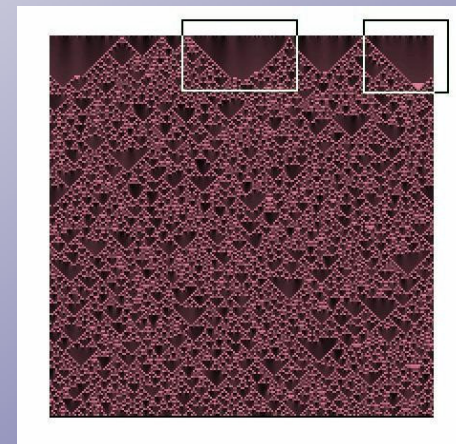
The Genetic Algorithm

Results

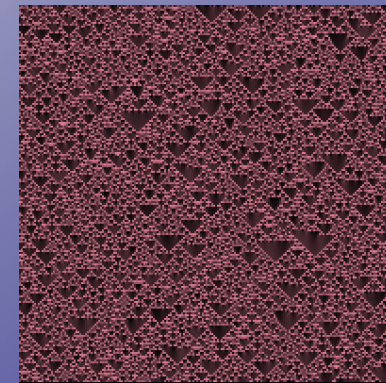
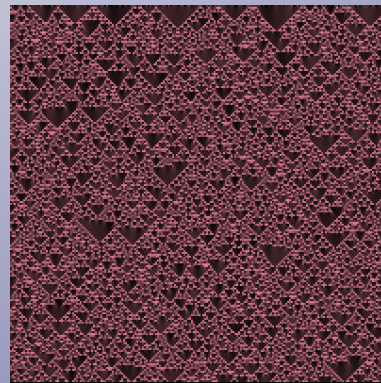
Target

Designoid

e5



f3



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Results

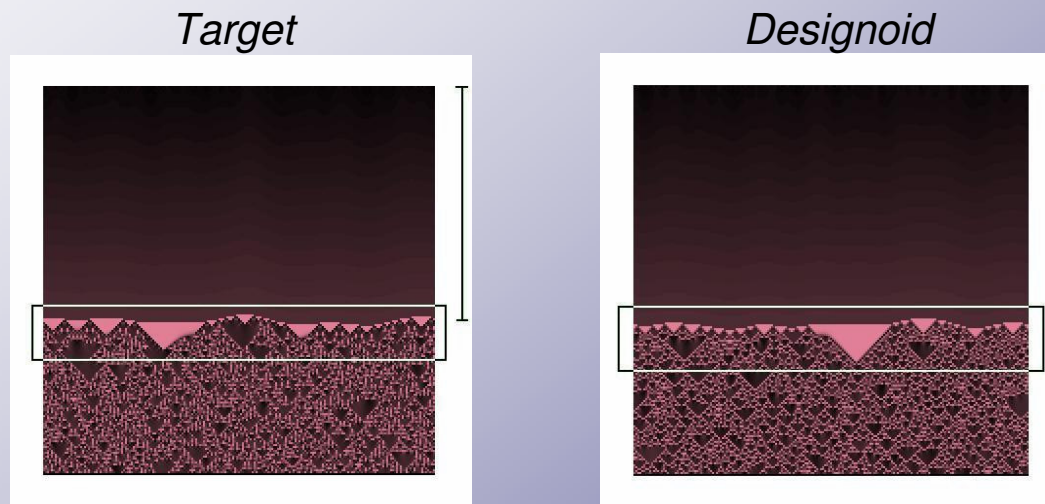
Target	Genotypic error				
	$usm(F,T)$	$e(i)$	$e(c)$	$e(r)$	E
$f2$	0.95657	0.30659	0.01885	12.77470	4.36672
$f3$	0.95608	0.35404	0.00426	10.40107	3.58646

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Target	$usm(F,T)$	$e(i)$	$e(c)$	$e(r)$	E
p	0.91980	0.26843	0.35314	0.05552	0.22569

Conclusions

- using Fitness Distance Correlation and Clustering, we can show the USM to be an appropriate objective function in this domain.
- can we generate a target spatio-temporal behaviour in a CA system?

YES

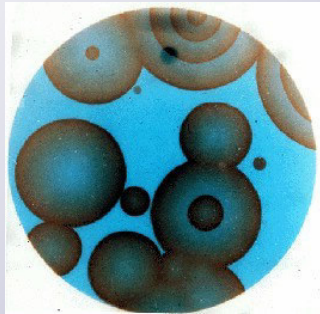
- GA generates very convincing designoid patterns

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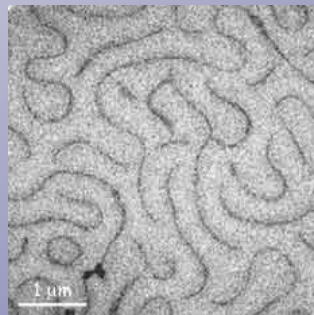
Applications (in design and manufacture) and further work

- Many, many systems can be modelled using CAs
- Research into chemical 'design'



e.g. designoid patterns in the BZ reaction

and self-organising nanostructured systems



We are actively working towards these practical goals in the context of the EPSRC grant *CHELLnet* (EP/D023343/1), which comprises

Evolvable CHELLware (EP/D021847/1),
vesiCHELL (EP/D022304/1),
brainCHELL (EP/D023645/1) and
wellCHELL (EP/D023807/1).

CHELLNet

<http://www.chellnet.org>



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