

Dr. Miguel Nicolau

Artificial Regulatory Network Models Definition and Application

COMP30290 Natural Computing COMP41190 Natural Computing and Applications



Natural Computing and Optimisation

Outline

Gene Regulatory Networks

2 Artificial Model

- Representation
- Example Dynamics
- **3** I/O Extensions to Model
- Pole-Balancing Experiments
- **5** Index Trading Experiments
- **Observations**



Genetic Regulatory Networks

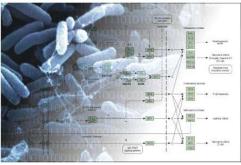


Gene Expression

- Cells express proteins when needed;
- Regulatory processes enhance/inhibit expression of genes;
- Responsible for differentiation and morphogenesis.



Genetic Regulatory Networks



Transcription Regulation

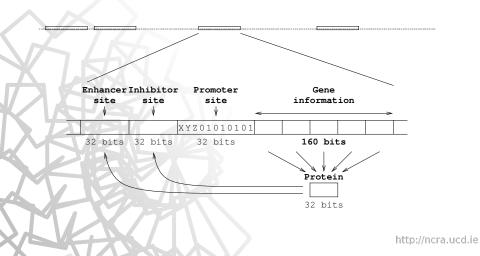
- Genes are transcribed at different rates;
- Transcription Factors (TFs) influence expression of genes;
- Genes can produce TFs, thus forming Regulatory Networks.



Wolfgang Banzhaf: Artificial Regulatory Networks and Genetic Programming. GPTP, 2003.

- Composition:
 - Genome (binary string);
 - Mobile Proteins (TFs).
- Promoter Sites identify genes;
- Genes composed of:
 - Enhancer and Inhibitor sites;
 - Gene information.







Dynamics

- Regulatory model is a closed world (no I/O);
- Production of proteins dictates their concentration: $\sum_{i} c_i = 1.0$
- Transcription factors bind to enhancer and inhibitor sites:
 - All TFs bind to all sites;
 - Strength of connection = # complementary bits (XOR):
 - Each gene has an **enhancing** and **inhibiting** signal:

$$e_i, h_i = rac{1}{N} \sum_{j=1}^N c_j \exp(eta(u_{i,j} - u_{max}))$$

 $u_{i,j} =$ match between site and protein j $u_{max} =$ maximum match $\beta =$ scaling factor



Dynamics

- Concentration of each protein changes per iteration:
 - Regulated by differential equation:

$$\frac{dc_i}{dt} = \delta(e_i - h_i)c_i - \Phi$$

$$\begin{split} \delta &= \text{scaling factor (time unit)} \\ \Phi &= \text{protein scaling factor } (\sum_i c_i = 1.0) \end{split}$$

- Concentrations change over time, depending on TFs matching sites;
- Rich dynamic regulatory network.



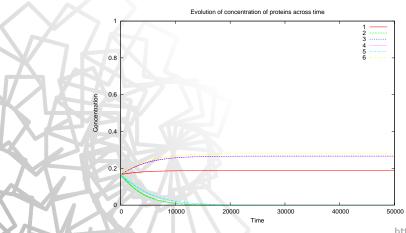
Dynamics

Expression level of genes change over time:

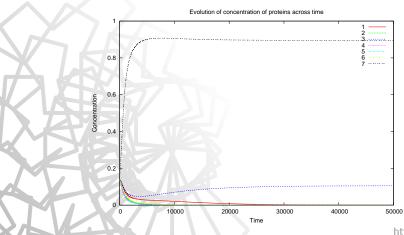
$$c_{t+1} = \text{signal} \times c_t$$

- Plot expression level of all genes across time;
- Different dynamics can be observed.

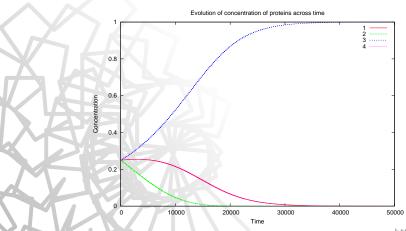






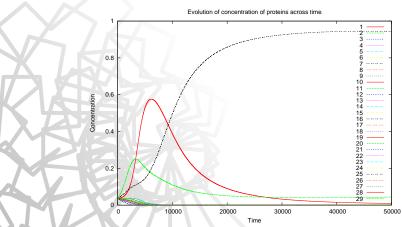




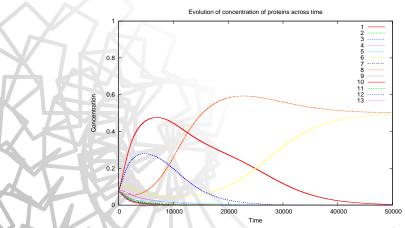


http://ncra.ucd.ie

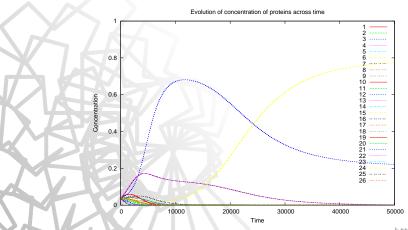




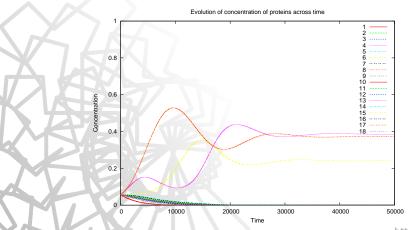




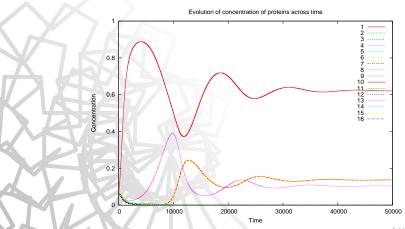




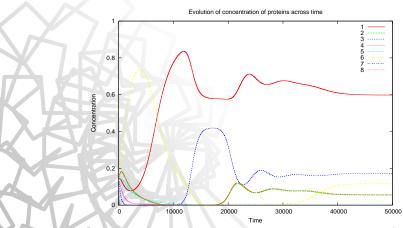














ARN: I/O Extensions

Modelling output

- Genes produce 2 types of genes:
 - TF-genes: produce TFs;
 - P-genes: produce product proteins (output signals).
- Different promoter site signatures identify different genes.

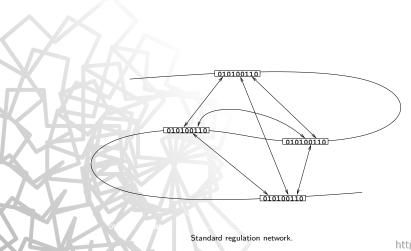
Modelling input

- Model input through extra regulatory (TF) proteins:
 - Not produced by any gene (external signals);
 - Co-regulate all genes with existing TFs.



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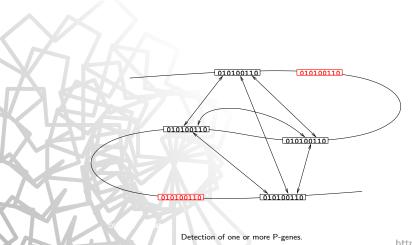
ARN: I/O Extensions





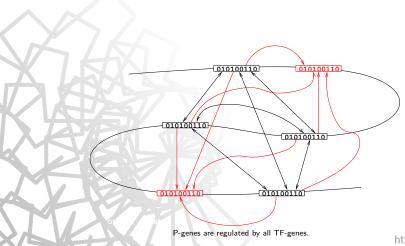
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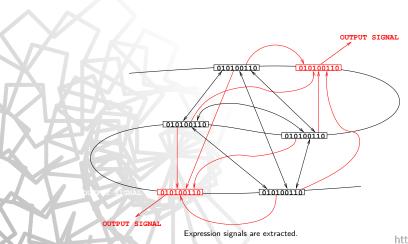


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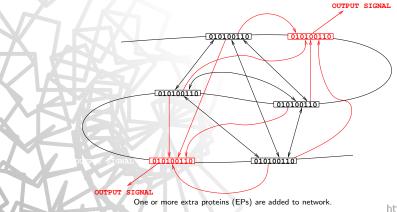




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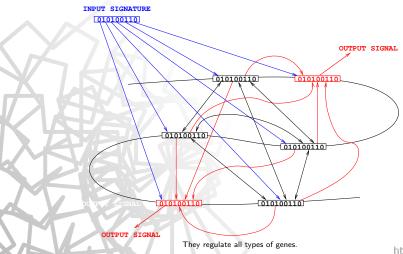
INPUT SIGNATURE





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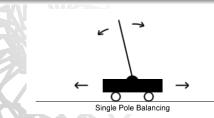




Pole-Balancing Experiments

Single-Pole Balancing (Inverted Pendulum)

- Famous dynamic control benchmark;
- Cart with rigid pole attached to it;
- Control cart along 1-dimensional track:
 - Apply constant force to either side of cart each 0.2s;
 - Keep pole within acceptable angle;
 - Keep cart within track boundaries.





Pole-Balancing Experiments

Pole-Balancing

- 4 inputs, 1 output:
 - $x \in [-2.4, 2.4]$ m = position of the cart relative to the centre;
 - $\theta \in [-12, 12]^{\circ} =$ angle of the pole with the vertical;
 - $\dot{x} \in [-1, 1]$ m/s = cart velocity;
 - $\dot{\theta} \in [-1.5, 1.5]^{\circ}/s =$ rate of change of the angle.
 - $F(t) = \pm 10$ N = bang-bang force applied to cart.

Experimental Setup

- (250+250)-ES (1% mutation rate, adapted);
- 0.2s = 2000 time steps in model;
- Measure concentration of chosen P-gene:
 - > .5 = push cart right, < .5 = push cart left.



Pole-Balancing Experiments

Results

	Best	Worst	Mean
random genomes	422	3	202.18
DM genomes	416	23	235.68

Analysis

- Results close to optimum (457 or less);
- Not far from best in literature (446).

UCD CASL Complex & Adaptive Systems Laboratory

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Index Trading Experiments

4000 TRAIN TEST1 TEST2 3500 D/N 3000 Montun 2500 Raw price -Trade signal 75 440 804 1170 1534 Day

FTSE Trader

GRN Trader

- Index Trading;
- Inputs: MA(n), SO(n), MC(n);
- Output is buy/sell/nothing;
- FTSE, DAX, Nikkei;

- Investment \$1000/\$10000, fixed term;
- Results equally good/better than B&H;
- Proof of concept.



Observations

Regulatory Networks as computational models

- Use of GRNs as new representation for GP;
- Investigation of more complex biological models for evolutionary computation.
 - Clear computational potential with current model;
 - Adaptation of such complex models often not obvious.

Future Work

- Calibration of parameters not easy;
 - Signature and concentration of extra proteins;
 - Calibration of biological and physical model.
- Dynamic control problems seem adequate.



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Next Classes

Discussion group (Thursday) 15h - 16h:

Toribash paper (blood and guts).