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# USING MOLECULAR CLASSIFIER SYSTEMS

SIXTH FRAMEWORK PROGRAMME

ESIGNET

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### **GOAL OF WORK PRESENTED IN THIS POSTER**

A novel approach to study Complex Adaptive System is presented. This work refines John Holland's proposal to allow the study of Molecular Complex Adaptive Systems such as Cell Signaling Networks. We present preliminary experiments focusing on the self-replication ability of these complex systems. Counter intuitive results were encountered, which suggest the importance of molecular specificity and necessity of a theoretical framework for the study of Artificial Chemistries.

### LIMITATIONS OF HOLLAND'S APPROACH

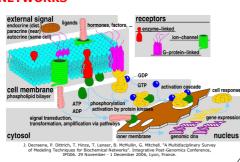
To study Complex Adaptive Systems, Holland proposed to employ an agent-based system in which Learning Classifier Systems (LCS) were used to determine the agents behavior and adaptivity. We argue that LCS are limited for the study of CAS: the rule-discovery mechanism is prespecified and may limit the evolvability of CAS. Secondly, LCS distinguish a demarcation between messages and rules, however operations are reflexive in CAS, e.g., in a cell, an agent (a molecule) may both act as a message (substrate) and as a catalyst (rule).

### **OUR APPROACH**

To address these issues, we proposed the **Molecular Classifier Systems** (MCS.b), a string-based Artificial Chemistry based on Holland's broadcast language. In the MCS.b, no explicit fitness function or rulediscovery mechanism is specified, moreover  $\ensuremath{\textbf{no}}$  distinction is made between messages and rules.

### **CELL SIGNALING NETWORKS**

In the context of the ESIGNET project, we employ the MCS.b to study a subclass of CAS: Cell Signaling Networks (CSNs) which are complex biochemical networks responsible for coordinating cellular activities.



### PRELIMINARY EXPERIMENTS ON SELF-REPLICATION

Closely relates to other studies: Tierra, Alchemy, Alpha Universe, Amoeba, etc, which were implemented differently but exhibited similar behavior.

### 1st experiment: domination of the self-replicators

#### $A + A + X \rightarrow 3A$

Common intuition: if a selfreplicase molecule A is present in the reactor (filled with random molecues and no mutation occur), then A should quickly dominate the reactor.

However our results differ: Fig 1

### Why is this happenning?

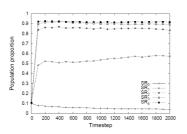
• In this experiment, replicase A was in fact not only a selfreplicase but а universal replicase.

• A universal replicator is said to have « zero specificity » replicate and would any molecules.

· Therefore a universal replicator does not possess any advantages over the other molecules

So, to dominate the reaction space, a replicator needs to be specific enough to prevent parasitic effects, Fig 2.

# 0.07 3000 4000 5000 6000 Timesten 7000 8000



Population growth of replicators  $SR_0, SR_1, SR_2, SR_3$  and  $SR_4$ ries is averaged over 30 simulation runs.

### So « specificity » matters!

### **MOLECULAR CLASSIFIER SYSTEMS - MCS.b**

- « Strong Artificial Life » approach
- String based artificial chemistry, loosely based on John Holland Broadcast Language
- Chemical processes as condition-action rules (IF condition x THEN action v)
- A molecule (string) can function both as substrate (message) and as catalyst (classifier)
- Formally an Abstract Term Rewriting System

### SELF-REPLICATION IN CELL SIGNALING NETWORKS

As CSNs occur in cells, these networks must replicate themselves prior to cell division. So that offspring cells obtain the necessary molecules to be functional. Errors may occur during this replication process, e.g., an offspring cell may inherit only a partial CSN. Thus resulting in a potentially defective cell which would lead to a variety of undesired effects (e.g., premature cell death). As a result, the **fitness** of a cell is implicitly represented by the survival and performance of a cell in achieving celllevel replication.

# Cell growth

Molecules (forming the CSN) are replicated

Cell division molecules are randomly distributed to offspring cells



· An ancestor molecule with high specificity was inserted « a la tierra ».

Mutation may now occur.

Expectetion: Domination of this ancestor over the population and emergence of organizations, such as collectively replicating sets of molecules, as observed in Tierra systems

results However, are again counter intuitive, see Fig 3.

### Why is this happenning?

- Fitter mutants emerged.
- Leading to an increase in molecule length.
- This molecular length growth leads to an elongation catastrophy, i.e., system extinction.

### CONCLUSION

These simple experiments exhibited **unexpected** results as opposed to those inferred from the literature. We demonstrated that molecular **specificity** plays an important role and may significantly influence the system dynamics. This work highlights the current deficit of a theoretical framework for the study of Artificial Chemistries.

### AKNOWLEGDMENTS

This work was supported by the ESIGNET project (Evolving Cell Signaling Networks in Silico). ESIGNET is a Specific Targeted Research Project funded by the European Commission under the Sixth Framework Programme (contract no. 12789).

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800 700 600 500 400 300 200 100 1500 2000 2500 Timestep

ules length growth upon overall system reactii m ancestor ( $SR_4 = \nabla 0101 : \nabla 0101$ ) is inser-ntration [ $SR_4$ ] = 0.1) in addition to randor eover mutation per molecule and per symbol nt, an an

