the limits of dynamical systems

DS represent agents only in terms of the interactions that exist and are assumed relevant in a given context; they do *not* represent agents in terms of *potential* interactions that are not causally effective now, but would become so if context was changing.

To describe agents in these terms requires a representation of their structure, which holds the key to possible interactions that can become actualized in other contexts.

It is an open question to what extent *physical* structure and modes of interaction can be lifted into a formal logical representation that permits such reasoning in the molecular realm.

The bottleneck in biology is not quantification, but description. (ouch!)

Turing Gas or Church Soup or AlChemy



fixed-points: self-maintenance



Self-maintenance is the consequence of a constructive feed-back loop: it occurs when the construction processes induced by the constituents of a system permit the continuous regeneration of these same constituents.

Immanuel Kant (Kritik der Urteilskraft, 1790): "[...] an organized product of nature is one in which all is end and, reciprocally, means too."

independent level of description



constrained extension





chemistry and proof theory



A more formal approach to agent-based systems is the π -calculus. Its application to biology was pioneered by Shapiro, Regev, and Priami. Its potential for biology was considerably deepened by Danos and Laneve.

The π -calculus (Milner, Walker and Parrow 1989)

- a program specifies a network of interacting processes
- processes are defined by their potential communication activities
- communication occurs on complementary channels, identified by names
- message content: channel name

Processes and channels

	P, Q, \cdots	process names (2.1)
\mathbf{Events}	x, y, \cdots	channel names (2.2)
	$\overline{x}, \overline{y}, \cdots$	channel co-names (2.3)
	$\pi ::= x \operatorname{comm}$	nunication on channel name x (2.4)
	\overline{x} comm	nunication on channel co-name x (2.5)
	x(y) receiv	$x \in y \text{ along } x \ (2.6)$
Process syntax $\overline{x}\langle y \rangle$ send y along x (2.7)		
P ::=	$P_1 \mid P_2$	parallel processes (2.8)
	π . P_1	sequential prefixing by communication (2.9)
	$\pi_1 . P_1 + \pi_2 . P_2$	mutually exclusive communications (2.10)
	$(new\;x)P$	new communication scope (2.11)
Structural con	0	inert process (2.12)

Structural congruence

$$P \mid Q \equiv Q \mid P$$

$$(P \mid Q) \mid R \equiv P \mid (Q \mid R)$$

$$P + Q \equiv Q + P$$

$$(P + Q) + R \equiv P + (Q + R)$$

$$(\text{new } x)0 \equiv 0$$

$$(\text{new } x)(\text{new } y)P \equiv (\text{new } y)(\text{new } x)P$$

$$((\text{new } x)P) \mid Q) \equiv (\text{new } x)(P \mid Q) \text{ if } x \notin FN(Q)$$

$$A(\vec{y}) \equiv \{\vec{y}/\vec{x}\}Q_A$$

$$x(y).P = x(z).(\{z/y\}P) \text{ if } z \notin FN(P)$$

$$(\text{new } y).P = (\text{new } z).(\{z/y\}P) \text{ if } z \notin FN(P)$$

associativity of PAR (2.14)commutativity of summation (2.15)associativity of summation (2.16)scope of inert processes (2.17)multiple communication scopes (2.18)scope extrusion (2.19)recursive parametric definition (2.20)renaming of input channel y (2.21)renaming of restricted channel y (2.22)

commutativity of PAR (2.13)

Reaction rules

$$\begin{array}{ll} (\cdots + \overline{x}\langle z \rangle.Q) | (\cdots + x(y).P) \rightarrow Q | P \{z/y\} & \text{communication (COMM)(2.23)} \\ & \text{if } P \rightarrow P' \text{ then } P | Q \rightarrow P' | Q & \text{reaction under parallel composition (2.24)} \\ & \text{if } P \rightarrow P' \text{ then (new } x)P \rightarrow (\text{new } x)P' & \text{reaction within restricted scope (2.25)} \\ & \text{if } Q \equiv P, P \rightarrow P', \text{ and } P' \equiv Q' \text{ then } Q \rightarrow Q' & \text{reaction up to structural congruence (2.26)} \end{array}$$

$E + S \rightleftharpoons ES \longrightarrow E + P$

- Says nothing about internal structure of E, S, P, ES
- We want to encode the reaction scheme... but according to certain principles

$\llbracket \mathsf{E} + \mathsf{S} \iff \mathsf{ES} \longrightarrow \mathsf{E} + \mathsf{P} \rrbracket_{\pi}$

•
$$\llbracket - \rightarrow_{\text{CHEM}} - \rrbracket_{\pi} = - \rightarrow_{\pi}^{*} -$$

•
$$\llbracket - +_{CHEM} - \rrbracket_{\pi} = - \downarrow -$$

L.Greg Meredith (2005)

$E + S \rightleftharpoons ES \longrightarrow E + P$

from these we deduce

•
$$\llbracket E + S \rrbracket_{\pi} = \llbracket E \rrbracket_{\pi} \mid \llbracket S \rrbracket_{\pi} \rightarrow_{\pi}^{*} \llbracket ES \rrbracket_{\pi}$$

• $\llbracket ES \rrbracket_{\pi} \rightarrow_{\pi}^{*} (\llbracket E \rrbracket_{\pi} \mid \llbracket S \rrbracket_{\pi}) + (\llbracket E \rrbracket_{\pi} \mid \llbracket P \rrbracket_{\pi})$

from these we deduce

- $\exists x_0 \cdot (\llbracket E \rrbracket_{\pi} \approx (\upsilon \ e)(x_0 [e] \cdot \llbracket E \rrbracket_{\pi'} + X_E)) \& (\llbracket S \rrbracket_{\pi} \approx x_0(y) \cdot \llbracket S \rrbracket_{\pi'} + X_S)$
- $\llbracket ES \rrbracket_{\pi} \approx (\upsilon \ e)(\llbracket E \rrbracket_{\pi}' \mid \llbracket S \rrbracket_{\pi}' \{ e/y \})$

therefore

• $(\upsilon e)(\llbracket E \rrbracket_{\pi}' | \llbracket S \rrbracket_{\pi}' \{e/y\}) \rightarrow_{\pi}^{*} (\llbracket E \rrbracket_{\pi} | \llbracket S \rrbracket_{\pi}) + (\llbracket E \rrbracket_{\pi} | \llbracket P \rrbracket_{\pi})$

L.Greg Meredith (2005)

$E + S \rightleftharpoons ES \longrightarrow E + P$

since *E* is an enzyme, $\llbracket E \rrbracket_{\pi}$ is the future of $\llbracket E \rrbracket_{\pi}'$, and $\llbracket S \rrbracket_{\pi}$ and $\llbracket P \rrbracket_{\pi}$ are the futures of $\llbracket S \rrbracket_{\pi}' \{ e/y \}$

• $(\upsilon e)(\llbracket E \rrbracket_{\pi}' | \llbracket S \rrbracket_{\pi}' \{e/y\}) \rightarrow_{\pi}^{*} (\llbracket E \rrbracket_{\pi} | \llbracket S \rrbracket_{\pi}) + (\llbracket E \rrbracket_{\pi} | \llbracket P \rrbracket_{\pi})$

implies

• $\exists x_1 x_2 \cdot (\llbracket E \rrbracket_{\pi}' \approx x_1(y) \cdot \llbracket E \rrbracket_{\pi} + x_2(y) \cdot \llbracket E \rrbracket_{\pi} + X_{E'}) & (\llbracket S \rrbracket_{\pi}' \approx x_1[e] \cdot \llbracket S \rrbracket_{\pi} + x_2[e] \cdot \llbracket P \rrbracket_{\pi} + X_{S'})$

setting X's to θ and minimizing the number of \rightarrow_{π} steps we arrive at

• $\llbracket E \rrbracket_{\pi} = (\upsilon \ e)(x_0[e].(x_1(y).\llbracket E \rrbracket_{\pi} + x_2(y).\llbracket E \rrbracket_{\pi}))$ • $\llbracket S \rrbracket_{\pi} = x_0(y).(x_1[e].\llbracket S \rrbracket_{\pi} + x_2[e].\llbracket P \rrbracket_{\pi})$

L.Greg Meredith (2005)

the concept of reaction or network type

$E + S \rightleftharpoons ES \longrightarrow E + P$

is really a reaction (network) type.

- use spatial logic (L.Caires) to capture the logical content (the characteristic formula F) of the process corresponding to this reaction
- translate biological networks into pi-processes xi
- model-check F against the xi
- thus identify networks with a (possibly dynamic) communication structure that behave like F (have that type)

The logic formula, "the largest process X that behaves in some way and eventually becomes X", describes the type "catalyst", which picks out the following red processes:



The spatial logic formula,

"the largest process X that behaves in some way and eventually becomes X|X", describes the type "autocatalyst", which picks out the following red processes:



an autocatalytic network at the dawn of life ?



Eric Smith & Harold Morowitz, PNAS, 101, 13168-13173 (2004)

The search for networks that inhabit certain types is important, because it extends current efforts at detecting network motifs.

Such efforts focus on syntactical motifs, but network types are behavioral motifs!

- detect whether, in a network, certain subgraphs occur more frequently than expected (expectation means a suitably randomized control)
- those that do are presumably solutions to some problem(s)
- figure out the problem(s)



the motives of motifs: "feed forward loop"

a delay mechanism...





...implementing a pulse-filter





from a physics of information to a biology of information

got guts?

i'm looking for a postdoc at the concurrency/biology interface of type:

must survive in a lab atmosphere & \diamond talk to biologists & have some physics intuition. (is this type inhabited?)