

Experiments in *Alchemy*

The Language

- Variation of Pure Lisp
- Expression
 - Atom
 - Operator: +, -, ', <, >, *, (,)
 - Variable: a, b, c,
 - List: (Expr1, Expr2, Expr3.....)

Semantics

(The “Meaning” of Expressions)

- The meaning of an expression is also an expression
 - Thus expressions have dual-identities as functions and objects
- The meaning of an expression is the result of “evaluating” the expression
 - Thus “meaning” is synonymous with expression evaluation

Molecule = Expression

- A Molecule in the system is an expression
 - Thus a molecule has a syntactic sense and a semantic sense

Interactions Between Molecules

Exp1 + Exp2 =

The meaning of ((' Exp1) (' Exp2))

Expression Evaluation

$$((+a)(*(>a)(>(+a))))$$
$$+$$
$$((*(+a)(*(-a)(<a)))a)$$

= meaning of [(('((+a)(*(>a)(>(+a)))))(('*(+a)(*(-a)(<a)))a))]

$$= (a(*(-a)(<a)))$$

See Trace of Expression Evaluation

```
• Evaluating Defined function (must first evaluate all expressions in the list):(((+a)(*>a)(>+a))))(((*+a)(*(-a)(<a)))a)))
• Evaluating '(((+a)(*>a)(>+a))))' with args:a:=a
• Evaluating Primitive function (must first evaluate all argument Expressions) :('((+a)(*>a)(>+a))))
• Value returned = ((+a)(*>a)(>+a)))
• Value returned = ((+a)(*>a)(>+a)))
• Evaluating '((*+a)(*(-a)(<a)))a)' with args:a:=a
• Evaluating Primitive function (must first evaluate all argument Expressions) :('((*+a)(*(-a)(<a)))a)
• Value returned = ((*+a)(*(-a)(<a)))a)
• Value returned = ((*+a)(*(-a)(<a)))a)
• ----Expression Evaluation Completed. We now Sequentially apply the resultant values to each other
• Evaluating ((+a)(*>a)(>+a))) with args:a:=(((*+a)(*(-a)(<a)))a)
• Evaluating Defined function (must first evaluate all expressions in the list):((+a)(*>a)(>+a)))
• Evaluating (+a) with args:a:=(((*+a)(*(-a)(<a)))a)
• Evaluating Primitive function (must first evaluate all argument Expressions) :(+a)
• Evaluating a with args:a:=(((*+a)(*(-a)(<a)))a)
• ----Argument evaluation Completed. We now apply the primitive to the resultant values
• Evaluating + with args:a:=(((*+a)(*(-a)(<a)))a)
• Value returned = ((*+a)(*(-a)(<a)))
• Value returned = ((*+a)(*(-a)(<a)))
• Evaluating (*>a)(>+a))) with args:a:=(((*+a)(*(-a)(<a)))a)
• Evaluating Primitive function (must first evaluate all argument Expressions) :(*>a)(>+a)))
• Evaluating >a) with args:a:=(((*+a)(*(-a)(<a)))a)
• Evaluating Primitive function (must first evaluate all argument Expressions) :(>a)
• Evaluating a with args:a:=(((*+a)(*(-a)(<a)))a)
• ----Argument evaluation Completed. We now apply the primitive to the resultant values
• Evaluating > with args:a:=(((*+a)(*(-a)(<a)))a)
• Value returned = (a)
• Value returned = a
• Evaluating (>+a)) with args:a:=(((*+a)(*(-a)(<a)))a)
• Evaluating Primitive function (must first evaluate all argument Expressions) :(>+a))
• Evaluating (+a) with args:a:=(((*+a)(*(-a)(<a)))a)
• Evaluating Primitive function (must first evaluate all argument Expressions) :(+a)
• Evaluating a with args:a:=(((*+a)(*(-a)(<a)))a)
• ----Argument evaluation Completed. We now apply the primitive to the resultant values
• Evaluating + with args:a:=(((*+a)(*(-a)(<a)))a)
• Value returned = ((*+a)(*(-a)(<a)))
• Value returned = ((*+a)(*(-a)(<a)))
• ----Argument evaluation Completed. We now apply the primitive to the resultant values
• Evaluating > with args:a:=(((*+a)(*(-a)(<a)))a)
• Value returned = ((*(-a)(<a)))
• Value returned = ((*(-a)(<a)))
• ----Argument evaluation Completed. We now apply the primitive to the resultant values
• Evaluating * with args:a:=a b:=((*(-a)(<a)))
```

Interaction Graph

- What do the lines mean?
- Notion of Elastic collisions

Iterated Interaction Graph

- See handout
- Autocatalytic Subgraphs
 - All nodes in the subgraph must have two incoming edges from other nodes in the subgraph
- Seeding Sets of an Autocatalytic subgraph
 - What happens if you remove a subset of the nodes in a subgraph
- Minimal Seeding Set of an Autocatalytic subgraph

Iterated Interaction Graph (cntd...)

- Closure

Turing Gas

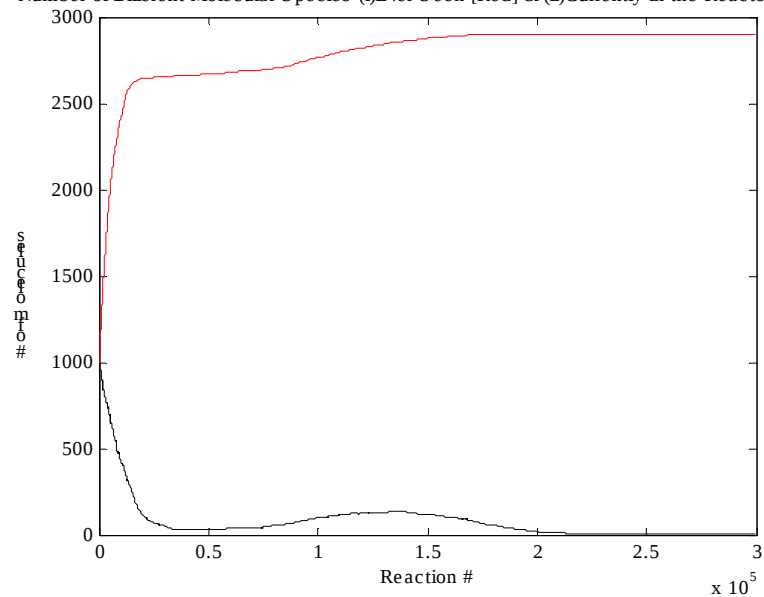
- Dynamics
 - Fixed size reactor
 - Reactions
 - Random collisions between molecules
 - Dilution
 - Random removal of a molecule in the reactor

Innovation and Absolute Innovation

- Innovative reaction: the species of the product molecule is not currently in the reactor
- Absolutely innovative reaction: the species of the product molecule has never been seen by the reactor during the run

One Sample Run

Number of Different Molecular Species (i)Ever Seen [Red] & (ii)Currently in the Reactor [Black]



(i) Innovation Rate [blue] & (ii) Absolute Innovation Rate [magenta]

