

pre-kappa expander for κ language

Héctor Urbina

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pre- κ
expander

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κ is a formal language for defining **agents** as sets of **sites**.

Sites hold an internal state as well as a binding state.

κ also enables the expression of rules of interaction between agents.

These rules are executable, inducing a stochastic dynamics on a mixture of agents.

A κ model is a collection of rules (with rate constants) and an initial mixture of agents on which such rules begin to act.

Krivine et. al. Programs as models: Kappa language basics. Unpublished work.

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Rule in English:

"Unphosphorilated Site1 of A binds to Site1 of B."

κ Rule:

$A(\text{Site1} \sim u), B(\text{Site1}) \rightarrow A(\text{Site1} \sim u!1), B(\text{Site1}!1)$

- Agent Names : an identifier.
- Agent Sites : an identifier.
- Internal States : $\sim \langle \text{value} \rangle$.
- Binding States : $! \langle n \rangle$, $!_-$ or $!?$.

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Kappa file structure

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```
1 ##### Signatures
2 %agent: A(x,c) # Declaration of agent A
3 %agent: B(x) # Declaration of B
4 %agent: C(x1~u~p,x2~u~p) # Declaration of C with 2 modifiable sites
5 ##### Rules
6 'a.b' A(x),B(x) -> A(x!1),B(x!1) @ 'on_rate' #A binds B
7 'a..b' A(x!1),B(x!1) -> A(x),B(x) @ 'off_rate' #AB dissociation
8 'ab.c' A(x!_,c),C(x1~u) ->A(x!_,c!2),C(x1~u!2) @ 'on_rate' #AB binds C
9 'mod x1' C(x1~u!1),A(c!1) ->C(x1~p),A(c) @ 'mod_rate' #AB modifies x1
10 'a.c' A(x,c),C(x1~p,x2~u) -> A(x,c!1),C(x1~p,x2~u!1) @ 'on_rate' #A binds C on x2
11 'mod x2' A(x,c!1),C(x1~p,x2~u!1) -> A(x,c),C(x1~p,x2~p) @ 'mod_rate' #A modifies x2
12 ##### Variables
13 %var: 'on_rate' 1.0E-4 # per molecule per second
14 %var: 'off_rate' 0.1 # per second
15 %var: 'mod_rate' 1 # per second
16 %obs: 'AB' A(x!x.B)
17 %obs: 'Cuu' C(x1~u,x2~u)
18 %obs: 'Cpu' C(x1~p,x2~u)
19 %obs: 'Cpp' C(x1~p,x2~p)
20 ##### Initial conditions
21 %init: 1000 A,B
22 %init: 10000 C
```

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DLab members study complex dynamical systems.

Currently, Cesar Ravello is modeling muscle contraction and Felipe Nuñez is simulating massive responses to zombie attacks on human populations, whereas Ricardo Honorato is adapting Model Checking techniques to be used with systems expressed in κ language.

Without intervening the κ language, we have reached some interesting levels of abstraction!

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- Space-related simulations.
 - Compartmentalization.
 - Diffusion events.
- Timing control.
 - Polymer-driven rules to manipulate latency.

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#Signatures

%agent: A(x,c,loc~i~j~k)

%agent: B(x,loc~i~j~k)

#Rules

#A binds B

A(x,loc~i),B(x,loc~i) \rightarrow A(x!1,loc~i),B(x!1,loc~i) @ 'on_rate'

A(x,loc~j),B(x,loc~j) \rightarrow A(x!1,loc~j),B(x!1,loc~j) @ 'on_rate'

A(x,loc~k),B(x,loc~k) \rightarrow A(x!1,loc~k),B(x!1,loc~k) @ 'on_rate'

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#Signatures

%agent: A(x,c,loc~i~j~k)

%agent: B(x,loc~i~j~k)

#Rules

#A binds B

$A(x, \text{loc} \sim i), B(x, \text{loc} \sim i) \rightarrow A(x!1, \text{loc} \sim i), B(x!1, \text{loc} \sim i) @ \text{'on_rate'}$

$A(x, \text{loc} \sim j), B(x, \text{loc} \sim j) \rightarrow A(x!1, \text{loc} \sim j), B(x!1, \text{loc} \sim j) @ \text{'on_rate'}$

$A(x, \text{loc} \sim k), B(x, \text{loc} \sim k) \rightarrow A(x!1, \text{loc} \sim k), B(x!1, \text{loc} \sim k) @ \text{'on_rate'}$

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#Locations i, j and k have different volumen/area!

#Signatures

%agent: A(x,c,loc~i~j~k)

%agent: B(x,loc~i~j~k)

#Rules

#A binds B

A(x,loc~i),B(x,loc~i) → A(x!1,loc~i),B(x!1,loc~i) @ 'on_rate_loc(i)'

A(x,loc~j),B(x,loc~j) → A(x!1,loc~j),B(X!1,loc~j) @ 'on_rate_loc(j)'

A(x,loc~k),B(x,loc~k) → A(x!1,loc~k),B(X!1,loc~k) @ 'on_rate_loc(k)'

#AB dissociation

A(x!1,loc~i),B(x!1,loc~i) → A(x,loc~i),B(x,loc~i) @ 'off_rate'

A(x!1,loc~j),B(x!1,loc~j) → A(x,loc~j),B(x,loc~j) @ 'off_rate'

A(x!1,loc~k),B(x!1,loc~k) → A(x,loc~k),B(x,loc~k) @ 'off_rate'

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#Locations i, j and k have different volumen/area!

#Signatures

%agent: A(x,c,loc~i~j~k)

%agent: B(x,loc~i~j~k)

#Rules

#A binds B

A(x,loc~i),B(x,loc~i) \rightarrow A(x!1,loc~i),B(x!1,loc~i) @ 'on_rate_loc(i)'

A(x,loc~j),B(x,loc~j) \rightarrow A(x!1,loc~j),B(X!1,loc~j) @ 'on_rate_loc(j)'

A(x,loc~k),B(x,loc~k) \rightarrow A(x!1,loc~k),B(X!1,loc~k) @ 'on_rate_loc(k)'

#AB dissociation

A(x!1,loc~i),B(x!1,loc~i) \rightarrow A(x,loc~i),B(x,loc~i) @ 'off_rate'

A(x!1,loc~j),B(x!1,loc~j) \rightarrow A(x,loc~j),B(x,loc~j) @ 'off_rate'

A(x!1,loc~k),B(x!1,loc~k) \rightarrow A(x,loc~k),B(x,loc~k) @ 'off_rate'

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%agent: A(x,c,loc~i~j~k)

%agent: B(x,loc~i~j~k)

#Rules

#A binds B

A(x,loc~i),B(x,loc~i) \rightarrow A(x!1,loc~i),B(x!1,loc~i) @ 'on_rate_loc(i)'

A(x,loc~j),B(x,loc~j) \rightarrow A(x!1,loc~j),B(X!1,loc~j) @ 'on_rate_loc(j)'

A(x,loc~k),B(x,loc~k) \rightarrow A(x!1,loc~k),B(X!1,loc~k) @ 'on_rate_loc(k)'

#AB dissociation

A(x!1,loc~i),B(x!1,loc~i) \rightarrow A(x,loc~i),B(x,loc~i) @ 'off_rate'

A(x!1,loc~j),B(x!1,loc~j) \rightarrow A(x,loc~j),B(x,loc~j) @ 'off_rate'

A(x!1,loc~k),B(x!1,loc~k) \rightarrow A(x,loc~k),B(x,loc~k) @ 'off_rate'

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#Signatures

%agent: A(x,c,loc~i~j~k)

%agent: B(x,loc~i~j~k)

%agent: T(s,org~i~j~k,dst~i~j~k)

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#Rules

#A diffusions

$A(\text{loc} \sim i, x, c), T(\text{org} \sim i, \text{dst} \sim j) \rightarrow A(\text{loc} \sim j, x, c), T(\text{org} \sim i, \text{dst} \sim j) @ 'Adiff_ij'$

$A(\text{loc} \sim i, x, c), T(\text{org} \sim i, \text{dst} \sim k) \rightarrow A(\text{loc} \sim k, x, c), T(\text{org} \sim i, \text{dst} \sim k) @ 'Adiff_ik'$

$A(\text{loc} \sim j, x, c), T(\text{org} \sim j, \text{dst} \sim i) \rightarrow A(\text{loc} \sim i, x, c), T(\text{org} \sim j, \text{dst} \sim i) @ 'Adiff_ji'$

$A(\text{loc} \sim j, x, c), T(\text{org} \sim j, \text{dst} \sim k) \rightarrow A(\text{loc} \sim k, x, c), T(\text{org} \sim j, \text{dst} \sim k) @ 'Adiff_jk'$

$A(\text{loc} \sim k, x, c), T(\text{org} \sim k, \text{dst} \sim i) \rightarrow A(\text{loc} \sim i, x, c), T(\text{org} \sim k, \text{dst} \sim i) @ 'Adiff_ki'$

$A(\text{loc} \sim k, x, c), T(\text{org} \sim k, \text{dst} \sim j) \rightarrow A(\text{loc} \sim j, x, c), T(\text{org} \sim k, \text{dst} \sim j) @ 'Adiff_kj'$

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#Signatures

%agent: $S(x)$

%agent: $Z()$

%agent: $V(p,n)$

#Rules

'Infection' $Z(), S(x) \rightarrow Z(), S(x!1), V(p!1,n) @ \text{'infection_rate'}$

'Polymerization' $V(n) \rightarrow V(n!1), V(p!1,n) @ \text{'polymer_rate'}$

'Expression' $S(x!1), V(p!1,n!2), V(p!2,n!3), V(p!3,n!4), \backslash$
 $V(p!4,n!5), V(p!5,n!6), V(p!6,n!7), V(p!7,n!8), V(p!8,n!9), \backslash$
 $V(p!9,n!10), V(p!10,n) \rightarrow Z() @ [\text{inf}]$

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 $V(p!4,n!5), V(p!5,n!6), V(p!6,n!7), V(p!7,n!8), V(p!8,n!9), \backslash$
 $V(p!9,n!10), V(p!10,n) \rightarrow Z() @ [\text{inf}]$

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'Expression' $S(x!1), V(p!1,n!2), V(p!2,n!3), V(p!3,n!4), \backslash$
 $V(p!4,n!5), V(p!5,n!6), V(p!6,n!7), V(p!7,n!8), V(p!8,n!9), \backslash$
 $V(p!9,n!10), V(p!10,n) \rightarrow Z() @ [\text{inf}]$

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A Python (V2) script that takes as input a (built in-house) **pre- κ** file and outputs a kappa file which can subsequently be used with KaSim.

This is done using Lexer & Parser techniques, available in Python through ply library.

It facilitates κ abstraction while reducing error-proneness.

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#Locations

%loc: i 100 10000 1201

%loc: j 1000 20000 3902

%loc: k 500 30000 2890

#Location list

%loc!: all i j k

#Signatures

%expand-agent: all A(x,c)

%expand-agent: all B(x)

gives:

%agent: A(x,c,loc~i~j~k)

%agent: B(x,loc~i~j~k)

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%loc: j 1000 20000 3902

%loc: k 500 30000 2890

#Location list

%locl: all i j k

#Signatures

%expand-agent: all A(x,c)

%expand-agent: all B(x)

gives:

%agent: A(x,c,loc~i~j~k)

%agent: B(x,loc~i~j~k)

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#Locations

%loc: i 100 10000 1201

%loc: j 1000 20000 3902

%loc: k 500 30000 2890

#Location list

%locl: all i j k

#Signatures

%expand-agent: all A(x,c)

%expand-agent: all B(x)

gives:

%agent: A(x,c,loc~i~j~k)

%agent: B(x,loc~i~j~k)

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#Locations

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%loc: j 1000 20000 3902

%loc: k 500 30000 2890

#Location list

%locl: all i j k

#Signatures

%expand-agent: all A(x,c)

%expand-agent: all B(x)

gives:

%agent: A(x,c,loc~i~j~k)

%agent: B(x,loc~i~j~k)

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#Locations

```
%loc: i 100 10000 1201
```

```
%loc: j 1000 20000 3902
```

```
%loc: k 500 30000 2890
```

#Location list

```
%locl: all i j k
```

#Initializations

```
%expand-init: all %loc[0] A(x,c)
```

```
%expand-init: all %loc[1] B(x)
```

gives:

```
%init: 100 A(x,c,loc~i)
```

```
%init: 1000 A(x,c,loc~j)
```

```
%init: 500 A(x,c,loc~k)
```

```
%init: 10000 B(x,loc~i)
```

```
%init: 20000 B(x,loc~j)
```

```
%init: 30000 B(x,loc~k)
```

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A bimolecular stochastic rate constant γ , expressed in $s^{-1} molecule^{-1}$, is related to its deterministic counterpart k , expressed in $s^{-1} M^{-1}$ as

$$\gamma = \frac{k}{AV}, \quad (1)$$

where A is Avogadro's number.

Krivine et. al. Programs as models: Execution. Unpublished work.

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#Locations

%loc: i 100 10000 1.201

%loc: j 1000 20000 3.902

%loc: k 500 30000 2.89

#Location list

%locl: all i j k

#A binds B

%expand-rule: all $A(x), B(x) \rightarrow A(x!1), B(x!1) @ \%loc[2]$

gives:

$A(x, loc\sim i), B(x, loc\sim i) \rightarrow A(x!1, loc\sim i), B(x!1, loc\sim i) @ 1.201$

$A(x, loc\sim j), B(x, loc\sim j) \rightarrow A(x!1, loc\sim j), B(x!1, loc\sim j) @ 3.902$

$A(x, loc\sim k), B(x, loc\sim k) \rightarrow A(x!1, loc\sim k), B(x!1, loc\sim k) @ 2.89$

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gives:

$A(x, loc \sim i), B(x, loc \sim i) \rightarrow A(x!1, loc \sim i), B(x!1, loc \sim i) @ 1.201$

$A(x, loc \sim j), B(x, loc \sim j) \rightarrow A(x!1, loc \sim j), B(x!1, loc \sim j) @ 3.902$

$A(x, loc \sim k), B(x, loc \sim k) \rightarrow A(x!1, loc \sim k), B(x!1, loc \sim k) @ 2.89$

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#Location matrices

%locm:

TM	i	j	k
i	0	0.5	1.5
j	2.0	0	1.8
k	1.0	1.1	0

#A diffusions

%expand-rule: TM A(x,c),T() \rightarrow A(%x,c),T() @ %cell

gives:

A(loc~i,x,c),T(org~i,dst~j) \rightarrow A(loc~j,x,c),T(org~i,dst~j) @ 0.5

A(loc~i,x,c),T(org~i,dst~k) \rightarrow A(loc~k,x,c),T(org~i,dst~k) @ 1.5

A(loc~j,x,c),T(org~j,dst~i) \rightarrow A(loc~i,x,c),T(org~j,dst~i) @ 2.0

A(loc~j,x,c),T(org~j,dst~k) \rightarrow A(loc~k,x,c),T(org~j,dst~k) @ 1.8

A(loc~k,x,c),T(org~k,dst~i) \rightarrow A(loc~i,x,c),T(org~k,dst~i) @ 1.0

A(loc~k,x,c),T(org~k,dst~j) \rightarrow A(loc~j,x,c),T(org~k,dst~j) @ 1.1

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A(loc~k,x,c),T(org~k,dst~i) \rightarrow A(loc~i,x,c),T(org~k,dst~i) @ 1.0

A(loc~k,x,c),T(org~k,dst~j) \rightarrow A(loc~j,x,c),T(org~k,dst~j) @ 1.1

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gives:

$A(\text{loc}^{\sim}i,x,c),T(\text{org}^{\sim}i,\text{dst}^{\sim}j) \rightarrow A(\text{loc}^{\sim}j,x,c),T(\text{org}^{\sim}i,\text{dst}^{\sim}j) @ 0.5$

$A(\text{loc}^{\sim}i,x,c),T(\text{org}^{\sim}i,\text{dst}^{\sim}k) \rightarrow A(\text{loc}^{\sim}k,x,c),T(\text{org}^{\sim}i,\text{dst}^{\sim}k) @ 1.5$

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$A(\text{loc}^{\sim}k,x,c),T(\text{org}^{\sim}k,\text{dst}^{\sim}i) \rightarrow A(\text{loc}^{\sim}i,x,c),T(\text{org}^{\sim}k,\text{dst}^{\sim}i) @ 1.0$

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#Location matrices

%locm:

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i	0	0.5	1.5
j	2.0	0	1.8
k	1.0	1.1	0

#Observing transporters

%expand-obs: TM 'Transporter(%org,%dst)' T()

gives:

%obs: 'Transporter(i,j)' T(org~i,dst~j)

%obs: 'Transporter(i,k)' T(org~i,dst~k)

%obs: 'Transporter(j,i)' T(org~j,dst~i)

%obs: 'Transporter(j,k)' T(org~j,dst~k)

%obs: 'Transporter(k,i)' T(org~k,dst~i)

%obs: 'Transporter(k,j)' T(org~k,dst~j)

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%obs: 'Transporter(i,j)' T(org~i,dst~j)

%obs: 'Transporter(i,k)' T(org~i,dst~k)

%obs: 'Transporter(j,i)' T(org~j,dst~i)

%obs: 'Transporter(j,k)' T(org~j,dst~k)

%obs: 'Transporter(k,i)' T(org~k,dst~i)

%obs: 'Transporter(k,j)' T(org~k,dst~j)

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#Rules

'Expression' $S(x!1), V(p!1, n!2), V(p!2, n!3), \dots, V(p!10, n) \rightarrow Z() @ [inf]$

gives:

'Expression' $S(x!1), V(p!1, n!2), V(p!2, n!3), V(p!3, n!4), \backslash$
 $V(p!4, n!5), V(p!5, n!6), V(p!6, n!7), V(p!7, n!8), V(p!8, n!9), \backslash$
 $V(p!9, n!10), V(p!10, n) \rightarrow Z() @ [inf]$

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 $V(p!4, n!5), V(p!5, n!6), V(p!6, n!7), V(p!7, n!8), V(p!8, n!9), \backslash$
 $V(p!9, n!10), V(p!10, n) \rightarrow Z() @ [inf]$