

Artificial Biochemistry

Combining Stochastic Collectives

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Microsoft Research

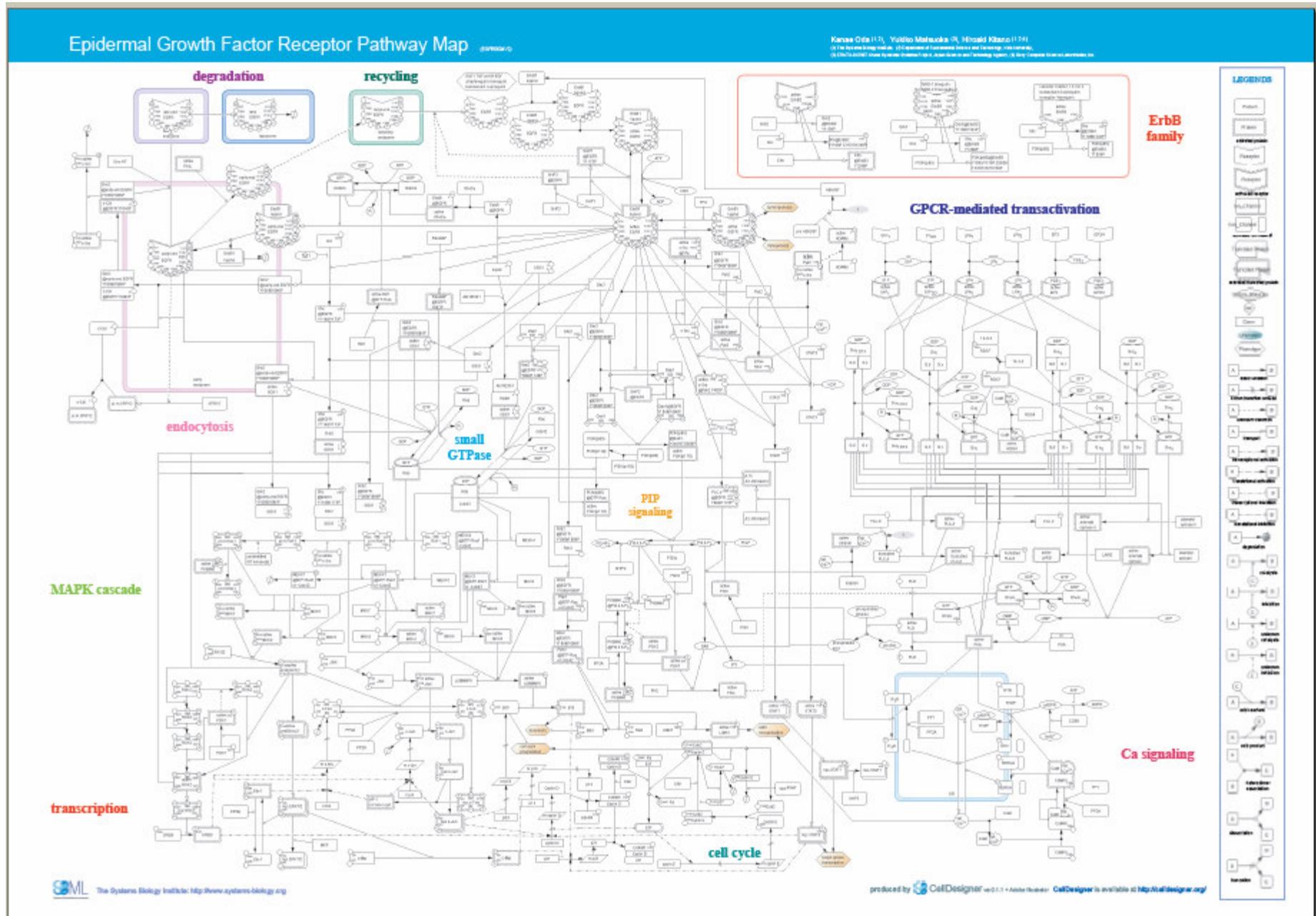
Dagstuhl 2006-04-20

www.luca.demon.co.uk

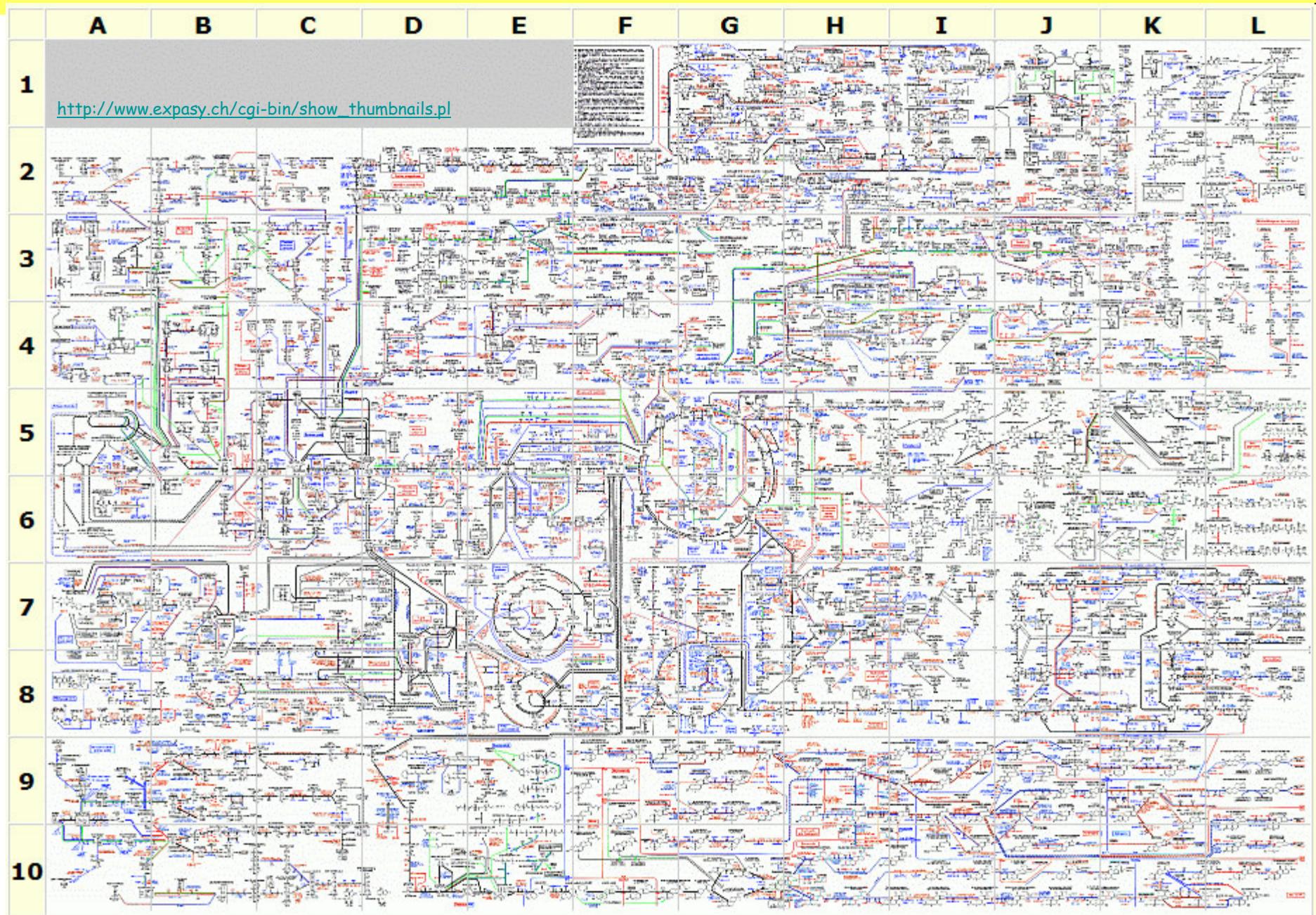
Stochastic Collectives

- “Collective”:
 - A large set of interacting finite state automata:
 - Not quite language automata (“large set”)
 - Not quite cellular automata (“interacting” but not on a grid)
 - Not quite process algebra (“finite state” and “collective”)
 - Not quite calculus (rate of change of “automata”??)
 - Cf. “multi-agent systems” and “swarm intelligence”
- “Stochastic”:
 - Interactions have *rates*
- Very much like biochemistry
 - Which is a large set of stochastically interacting molecules/proteins
 - Are proteins **finite state** and subject to automata-like **transitions**?
 - Let's say they are, at least because:
 - Much of the knowledge being accumulated in Systems Biology is described as state transition diagrams [Kitano].

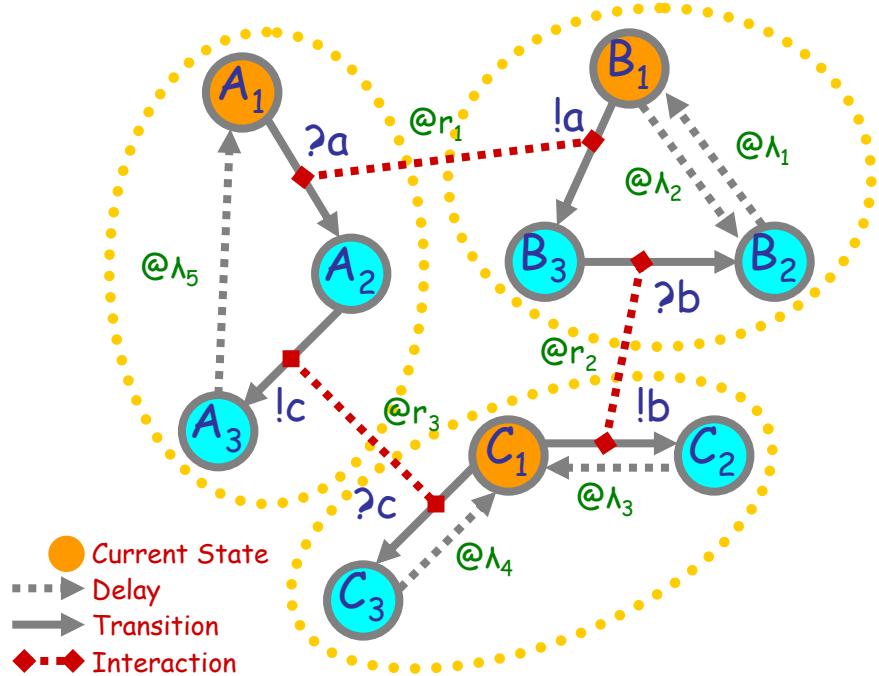
State Transitions



Even More State Transitions



Interacting Automata

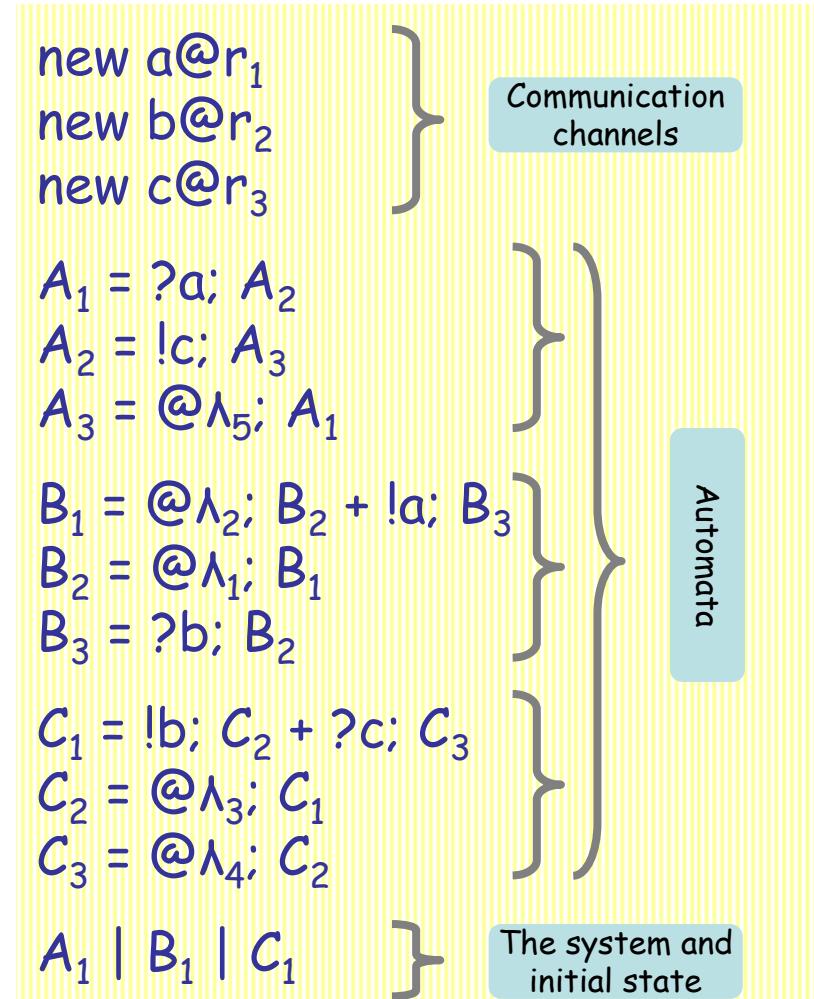


Communicating automata: a graphical FSA-like notation for “finite state restriction-free π -calculus processes”. *Interacting automata* do not even exchange values on communication.

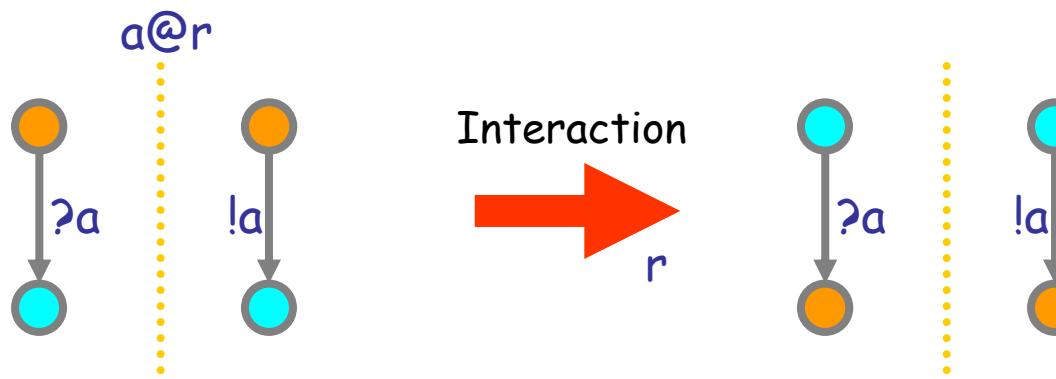
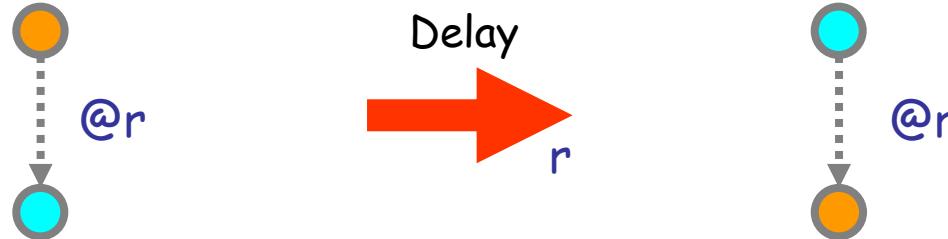
The stochastic version has *rates* on communications, and delays.

“Finite state” means: no composition or restriction inside recursion.

Analyzable by standard Markovian techniques, by first computing the “product automata” to obtain the underlying finite Markov transition system. [Buchholz]

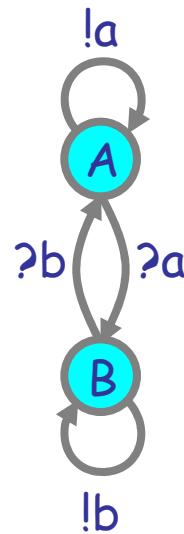


Interacting Automata Transition Rules



- Current State
- Delay
- Transition

Groupies and Celebrities



Celebrity

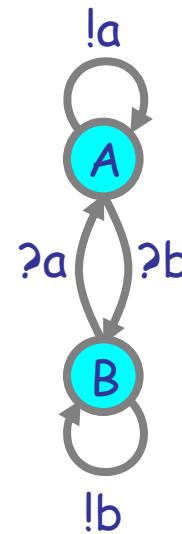
(does not want to be like somebody else)

directive sample 0.1 1000
directive plot A(); B()

new a@1.0:chan()
new b@1.0:chan()

let A() = do !a; A() or ?a; B()
and B() = do !b; B() or ?b; A()

run 100 of (A() | B())



Groupie

(wants to be like somebody different)

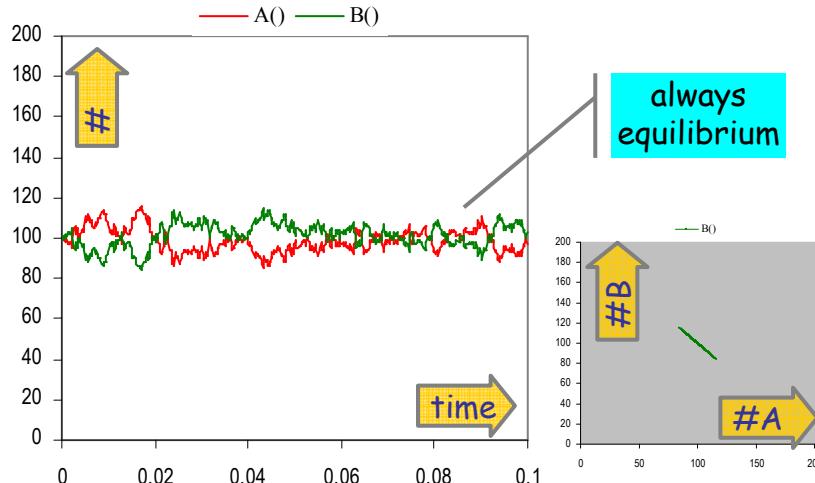
directive sample 5.0 1000
directive plot A(); B()

new a@1.0:chan()
new b@1.0:chan()

let A() = do !a; A() or ?b; B()
and B() = do !b; B() or ?a; A()

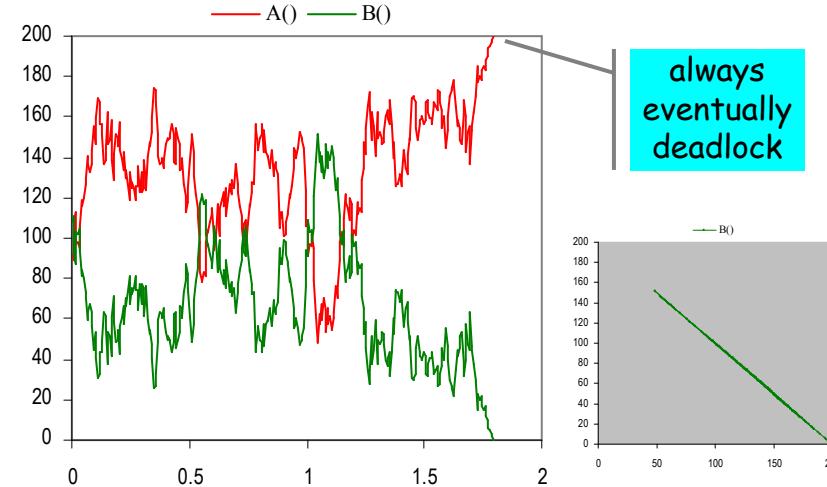
run 100 of (A() | B())

A stochastic collective of celebrities:



Stable because as soon as a A finds itself in the majority, it is more likely to find somebody in the same state, and hence change, so the majority is weakened.

A stochastic collective of groupies:

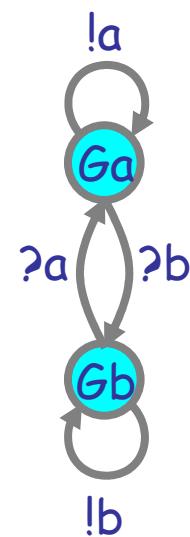


Unstable because within an A majority, an A has difficulty finding a B to emulate, but the few B's have plenty of A's to emulate, so the majority may switch to B. Leads to deadlock when everybody is in the same state and there is nobody different to emulate.

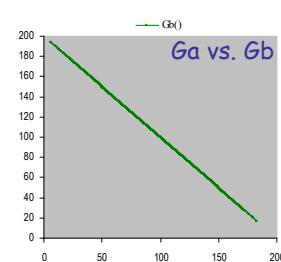
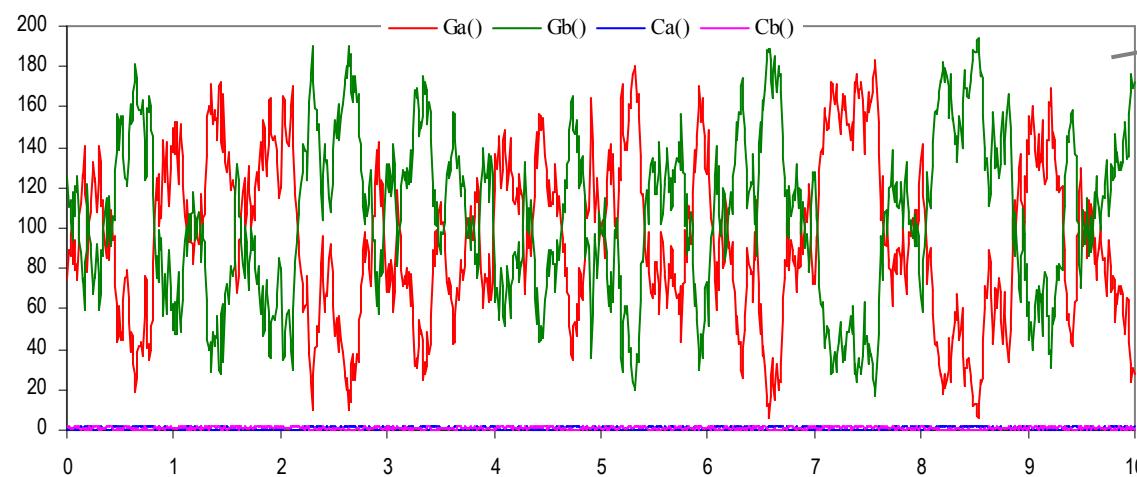
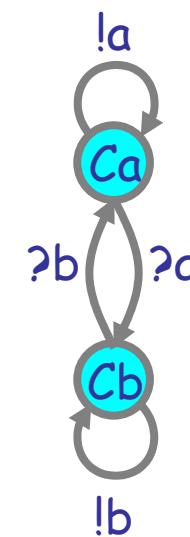
Both Together

A tiny bit of
"noise" can make a
huge difference

Many
Groupies



A few
Celebrities



```

directive sample 10.0 1000
directive plot Ga(); Gb(); Ca(); Cb()

new a@1.0:chan()
new b@1.0:chan()

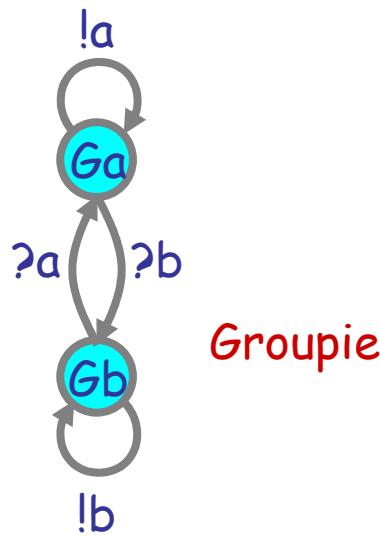
let Ca() = do !a; Ca() or ?a; Cb()
and Cb() = do !b; Cb() or ?b; Ca()

let Ga() = do !a; Ga() or ?b; Gb()
and Gb() = do !b; Gb() or ?a; Ga()

run 1 of (Ca() | Cb())
run 100 of (Ga() | Gb())
  
```

Doped Groupies

A similar way to break the deadlocks: destabilize the groupies by a small perturbation.



```

directive sample 10.0 1000
directive plot Ga(); Gb(); Da(); Db()

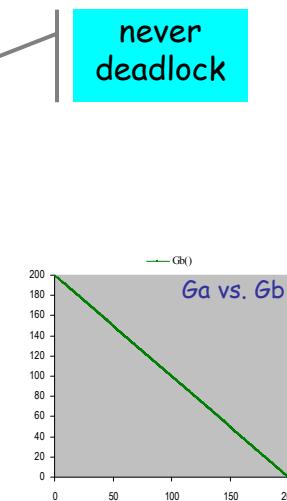
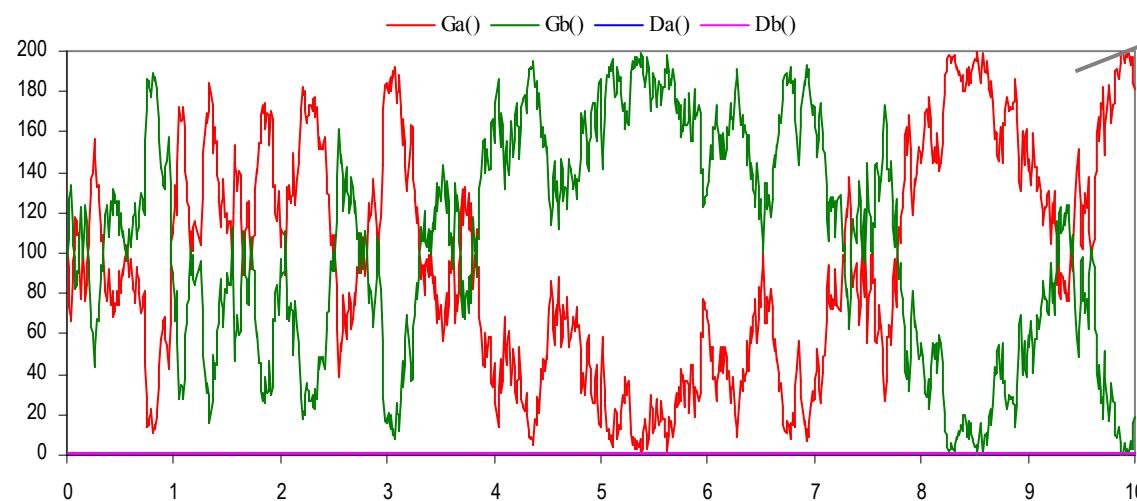
new a@1.0:chan()
new b@1.0:chan()

let Ga() = do !a; Ga() or ?b; Gb()
and Gb() = do !b; Gb() or ?a; Ga()

let Da() = !a; Da()
and Db() = !b; Db()

run 1 of (Da() | Db())
run 100 of (Ga() | Gb())

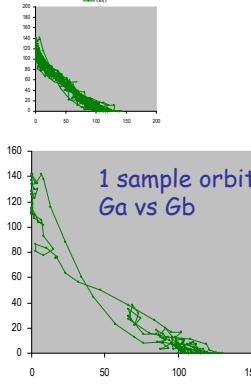
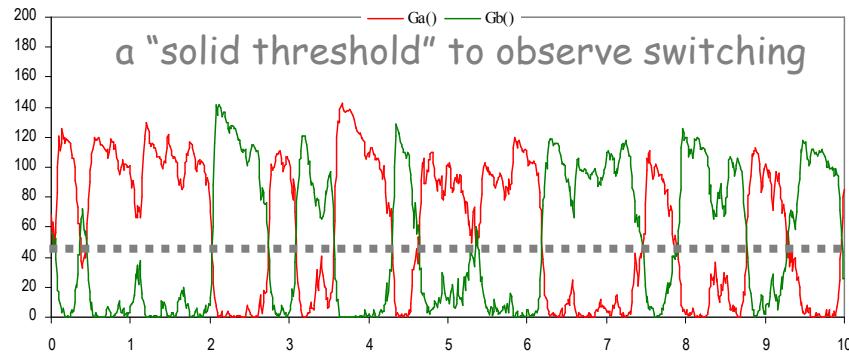
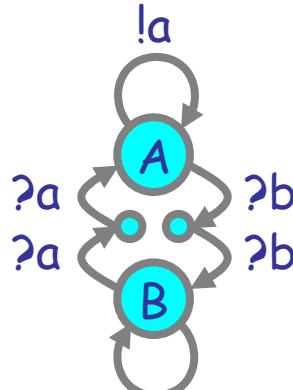
```



⁽¹⁾A technical term in microelectronics

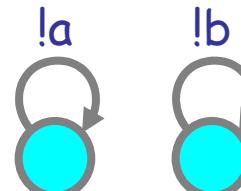
Hysteric Groupies

We can get more regular behavior from groupies if they "need more convincing", or "**hysteresis**" (history-dependence), to switch states.



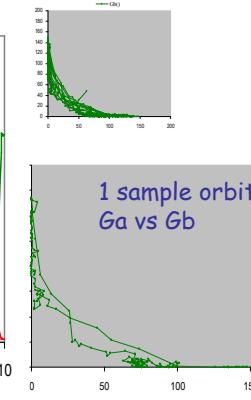
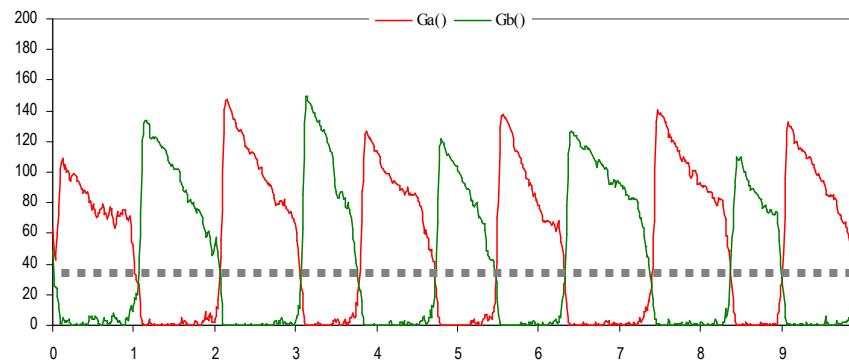
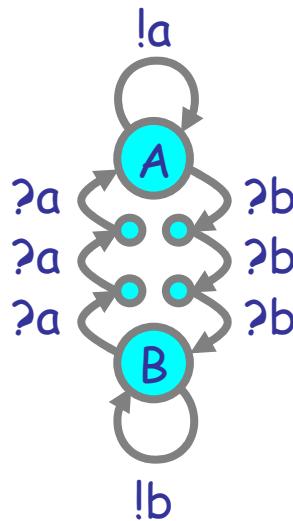
```
directive sample 10.0 1000
directive plot  $\text{Ga}(); \text{Gb}()$ 
new a@1.0:chan()
new b@1.0:chan()
let  $\text{Ga}()$  = do !a;  $\text{Ga}()$  or ?b; ?b;  $\text{Gb}()$ 
and  $\text{Gb}()$  = do !b;  $\text{Gb}()$  or ?a; ?a;  $\text{Ga}()$ 
let  $\text{Da}()$  = !a;  $\text{Da}()$ 
and  $\text{Db}()$  = !b;  $\text{Db}()$ 
run 100 of ( $\text{Ga}()$  |  $\text{Gb}()$ )
run 1 of ( $\text{Da}()$  |  $\text{Db}()$ )
```

$!b$



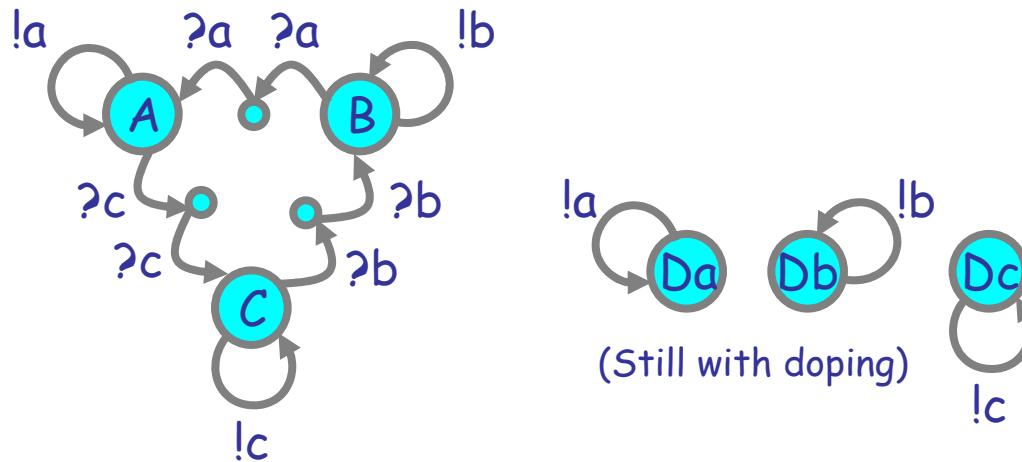
(Still with doping)

N.B.: It will not oscillate without doping (noise)

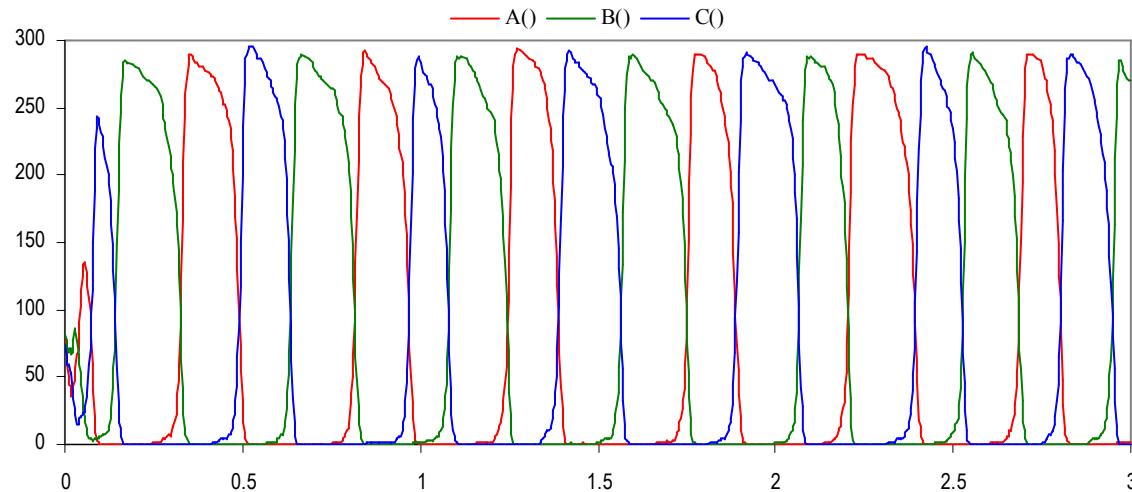


```
directive sample 10.0 1000
directive plot  $\text{Ga}(); \text{Gb}()$ 
new a@1.0:chan()
new b@1.0:chan()
let  $\text{Ga}()$  = do !a;  $\text{Ga}()$  or ?b; ?b;  $\text{Gb}()$ 
and  $\text{Gb}()$  = do !b;  $\text{Gb}()$  or ?a; ?a;  $\text{Ga}()$ 
let  $\text{Da}()$  = !a;  $\text{Da}()$ 
and  $\text{Db}()$  = !b;  $\text{Db}()$ 
run 100 of ( $\text{Ga}()$  |  $\text{Gb}()$ )
run 1 of ( $\text{Da}()$  |  $\text{Db}()$ )
```

Hysteric 3-Way Groupies



N.B.: It will not oscillate without doping (noise)



```

directive sample 3.0 1000
directive plot A(); B(); C()

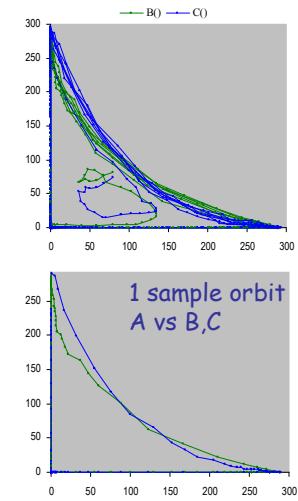
new a@1.0:chan()
new b@1.0:chan()
new c@1.0:chan()

let A() = do !a; A() or ?c; ?c; C()
and B() = do !b; B() or ?a; ?a; A()
and C() = do !c; C() or ?b; ?b; B()

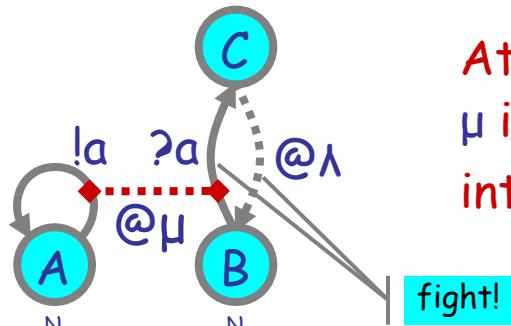
let Da() = !a; Da()
and Db() = !b; Db()
and Dc() = !c; Dc()

run 100 of (A() | B() | C())
run 1 of (Da() | Db() | Dc())

```



The Strength of Populations



At size $2N$, on a shared channel,
 μ is N times stronger than λ :
 interaction easily wins over delay.

```
directive sample 0.01 1000
directive plot B()

val lam = 1000.0
val mu = 1.0

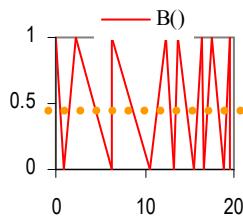
new a@mu:chan
let A() = !a; A()
and B() = ?a; C()
and C() = delay@lam; B()

run 1000 of (A() | B())
```

$$N=1$$

$$\lambda=1$$

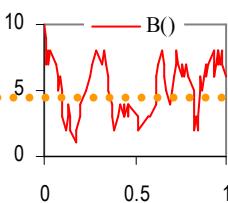
$$\mu=1$$



$$N=10$$

$$\lambda=10$$

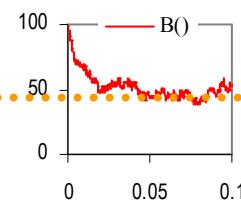
$$\mu=1$$



$$N=100$$

$$\lambda=100$$

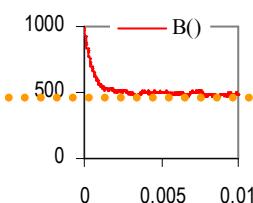
$$\mu=1$$



$$N=1000$$

$$\lambda=1000$$

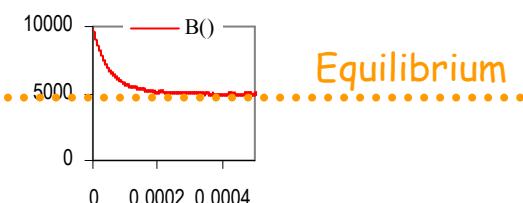
$$\mu=1$$



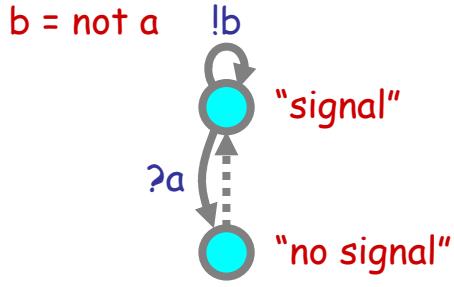
$$N=10000$$

$$\lambda=10000$$

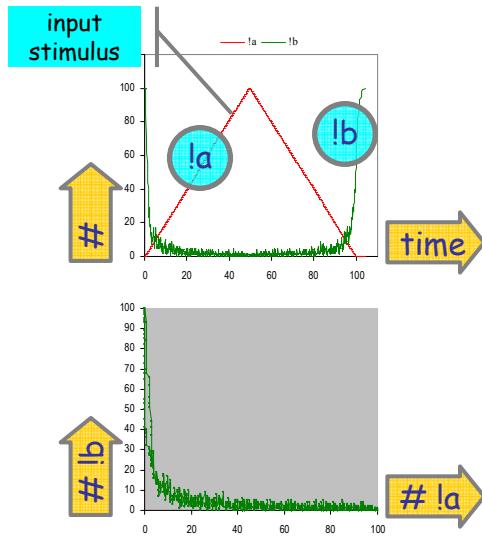
$$\mu=1$$



Boolean Inverter Collectives



in presence of a , b goes low
in absence of a , b goes high

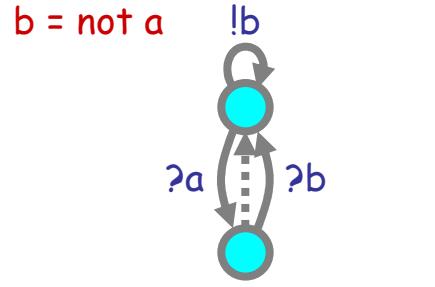


```
directive sample 110.0 1000
directive plot !a; !b

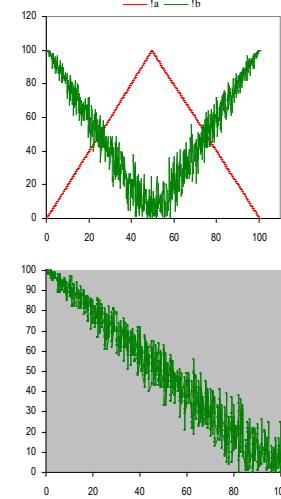
new a@1.0 chan new b@1.0 chan

new a@1.0 chan new b@1.0 chan
new b@1.0 chan new a@1.0 chan

let Inv_h(a,chan,b,chan)=
do !b: Inv_h(a,b)
or !a: Inv_j(a,b)
and Inv_j(a,chan,b,chan)=
delay@1.0: Inv_h(a,b)
```



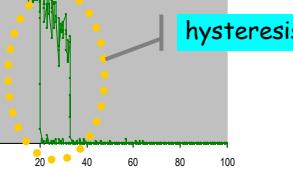
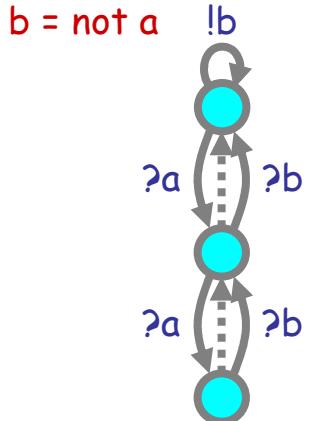
the high b state reinforces itself (as a population)



```
directive sample 110.0 1000
directive plot !a; !b

new a@1.0 chan new b@1.0 chan
new b@1.0 chan new a@1.0 chan

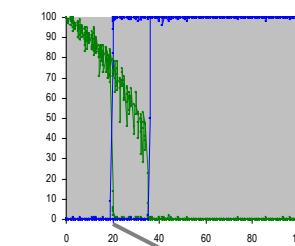
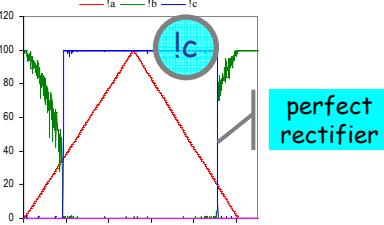
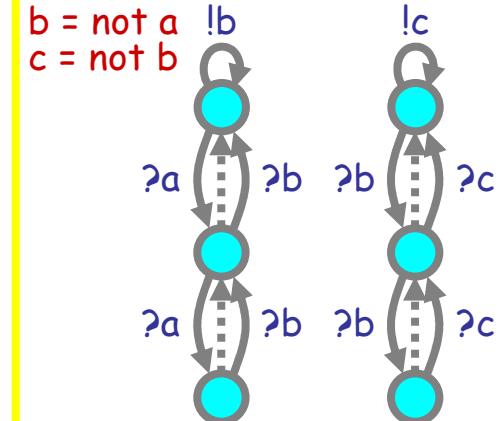
let Inv_h(a,chan,b,chan)=
do !b: Inv_h(a,b)
or !a: Inv_j(a,b)
and Inv_j(a,chan,b,chan)=
delay@1.0: Inv_h(a,b)
```



```
directive sample 110.0 1000
directive plot !a; !b

new a@1.0 chan new b@1.0 chan
new b@1.0 chan new a@1.0 chan

let Inv_h(a,chan,b,chan)=
do !b: Inv_h(a,b)
or !a: Inv_j(a,b)
and Inv_j(a,chan,b,chan)=
delay@1.0: Inv_h(a,b)
```



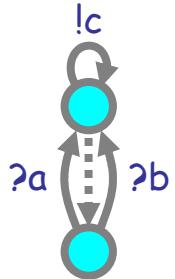
```
directive sample 110.0 1000
directive plot !a; !b; !c

new a@1.0 chan new b@1.0 chan new c@1.0 chan

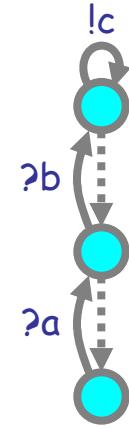
let Inv2_h(a,chan,b,chan)=
do !b: Inv2_h(a,b)
or !c: Inv2_m(a,b)
and Inv2_m(a,chan,b,chan)=
do !b: Inv2_j(a,b) or delay@1.0: Inv2_h(a,b)
or !c: Inv2_m(a,b)
and Inv2_j(a,chan,b,chan)=
do !b: Inv2_j(a,b) or delay@1.0: Inv2_m(a,b)
```

Boolean Gate Collectives

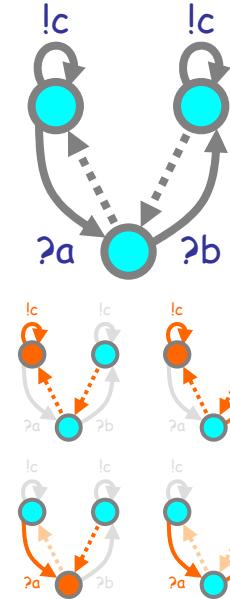
$c = a \text{ or } b$



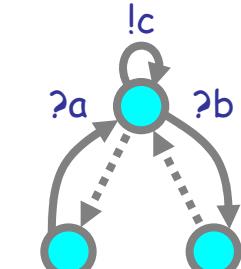
$c = a \text{ and } b$



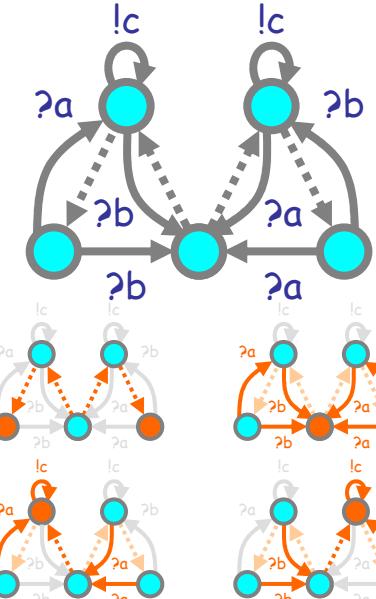
$c = a \text{ imply } b$



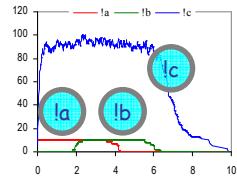
$c = a \text{ unless } b$



$c = a \text{ xor } b$



Inputs:
10 la for 4t
2t; 10 !b for 4t



directive sample 10.0 1000
directive plot la; lc

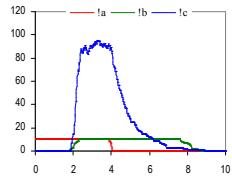
new a@1.0:chan new b@1.0:chan new c@1.0:chan
val del = 1.0

```
let Or_hi(a:chan, b:chan, c:chan) =  
do lc: Or_hi(b,c) or delay@del: Or_lo(a,b,c)  
and Or_lo(a:chan, b:chan, c:chan) =  
do hc: Or_hi(a,b,c) or !hc: Or_lo(a,b,c)
```

run 100 of Or_lo(a,b,c)

```
let clock@(float, tickchan) = (* sends a tick every t time *)  
(val t = 1/200.0 val d = 1.0/t)  
let step(n)=  
if n>0 then tick; clock(t, tick) else delay@d: step(n-1)  
run step(200)
```

```
let S_a(tickchan) = do la: S_a(tick) or !hc: ()  
let S_b(tickchan) = !hc & S_b(tick)  
and S_b(tickchan) = do lb: S_b(tick) or !hc: S_b2(tick)  
end S_b2(tickchan) = do lb: S_b2(tick) or !hc: S_b2(tick)  
and S_b2(tickchan) = do lb: S_b3(tick) or !hc: ()  
run 10 of (new tickchan run (clock(4.0,tick) | S_a(tick)))  
run 10 of (new tickchan run (clock(2.0,tick) | S_b(tick)))
```



directive sample 10.0 1000
directive plot la; lc

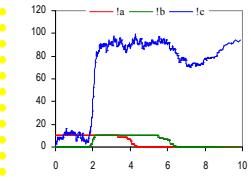
new a@1.0:chan new b@1.0:chan new c@1.0:chan
val del = 1.0

```
let And_hi(a:chan, b:chan, c:chan) =  
do lc: And_hi(b,c) or delay@del: And_lo(a,b,c)  
and And_lo(a:chan, b:chan, c:chan) =  
do hc: And_hi(a,b,c) or delay@del: And_lo(a,b,c)  
do hb: And_lo(b,a,c) or delay@del: And_lo(a,b,c)
```

run 100 of And_lo(a,b,c)

```
let clock@(float, tickchan) = (* sends a tick every t time *)  
(val t = 1/200.0 val d = 1.0/t)  
let step(n)=  
if n>0 then tick; clock(t, tick) else delay@d: step(n-1)  
run step(200)
```

```
let S_a(tickchan) = do la: S_a(tick) or !hc: ()  
let S_b(tickchan) = !hc & S_b(tick)  
and S_b(tickchan) = do lb: S_b(tick) or !hc: S_b2(tick)  
end S_b2(tickchan) = do lb: S_b2(tick) or !hc: S_b2(tick)  
and S_b2(tickchan) = do lb: S_b3(tick) or !hc: ()  
run 10 of (new tickchan run (clock(4.0,tick) | S_a(tick)))  
run 10 of (new tickchan run (clock(2.0,tick) | S_b(tick)))
```



directive sample 10.0 1000
directive plot la; lc

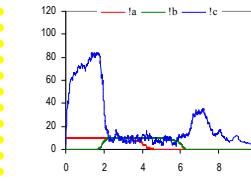
new a@1.0:chan new b@1.0:chan new c@1.0:chan
val del = 1.0

```
let Imply_hi(a:chan, b:chan, c:chan) =  
do lc: Imply_hi(b,c) or !hc: Imply_lo(a,b,c)  
and Imply_lo(a:chan, b:chan, c:chan) =  
do hc: Imply_hi(b,c) or delay@del: Imply_lo(a,b,c)  
do hb: Imply_lo(b,a,c) or delay@del: Imply_hi(a,b,c)
```

run 100 of Imply_lo(a,b,c)

```
let clock@(float, tickchan) = (* sends a tick every t time *)  
(val t = 1/200.0 val d = 1.0/t)  
let step(n)=  
if n>0 then tick; clock(t, tick) else delay@d: step(n-1)  
run step(200)
```

```
let S_a(tickchan) = do la: S_a(tick) or !hc: ()  
let S_b(tickchan) = !hc & S_b(tick)  
and S_b(tickchan) = do lb: S_b(tick) or !hc: S_b2(tick)  
end S_b2(tickchan) = do lb: S_b2(tick) or !hc: S_b2(tick)  
and S_b2(tickchan) = do lb: S_b3(tick) or !hc: ()  
run 10 of (new tickchan run (clock(4.0,tick) | S_a(tick)))  
run 10 of (new tickchan run (clock(2.0,tick) | S_b(tick)))
```



directive sample 10.0 1000
directive plot la; lc

new a@1.0:chan new b@1.0:chan new c@1.0:chan
val del = 1.0

```
let OOO_hi(a:chan, b:chan, c:chan) =  
do lc: OOO_hi(b,c) or !hc: OOO_lo(a,b,c) or !hb: OOO_lo(b,a,c)  
and OOO_lo(a:chan, b:chan, c:chan) =  
do hc: OOO_hi(b,c) or delay@del: OOO_lo(a,b,c)  
do hb: OOO_hi(b,a,c) or delay@del: OOO_lo(a,b,c)  
do hc: OOO_lo(b,a,c) or delay@del: OOO_hi(a,b,c)
```

run 50 of (OOO_lo(a,b,c) | OOO_hi(b,a,c))

```
let clock@(float, tickchan) = (* sends a tick every t time *)  
(val t = 1/200.0 val d = 1.0/t)  
let step(n)=  
if n>0 then tick; clock(t, tick) else delay@d: step(n-1)  
run step(200)
```

```
let S_a(tickchan) = do la: S_a(tick) or !hc: ()  
let S_b(tickchan) = !hc & S_b(tick)  
and S_b(tickchan) = do lb: S_b(tick) or !hc: S_b2(tick)  
end S_b2(tickchan) = do lb: S_b2(tick) or !hc: S_b2(tick)  
and S_b2(tickchan) = do lb: S_b3(tick) or !hc: ()  
run 10 of (new tickchan run (clock(4.0,tick) | S_a(tick)))  
run 10 of (new tickchan run (clock(2.0,tick) | S_b(tick)))
```

directive sample 10.0 1000
directive plot la; lc

new a@1.0:chan new b@1.0:chan new c@1.0:chan

```
let Xor_hi(a:chan, b:chan, c:chan) =  
do lc: Xor_hi(a,b,c) or !hb: Xor_lo_ab(a,b,c) or delay@1.0: Xor_lo_b(a,b,c)
```

```
and Xor_lo(a:chan, b:chan, c:chan) =  
do hc: Xor_hi(b,c) or !hc: Xor_lo_ab(b,c) or delay@1.0: Xor_lo_b(b,c)
```

```
and Xor_lo(a:chan, b:chan, c:chan) =  
do hc: Xor_hi(b,c) or !hb: Xor_lo_ab(b,c) or delay@1.0: Xor_lo_b(b,c)
```

```
and Xor_lo(a:chan, b:chan, c:chan) =  
do hb: Xor_hi(b,c) or !hc: Xor_lo_ab(c,b) or delay@1.0: Xor_lo_b(c,b)
```

```
do delay@1.0: Xor_hi_ab(c,b) or !hc: Xor_lo_hi(b,a,c)
```

run 50 of (Xor_lo_ab(b,c) | Xor_lo_hi(b,a,c))

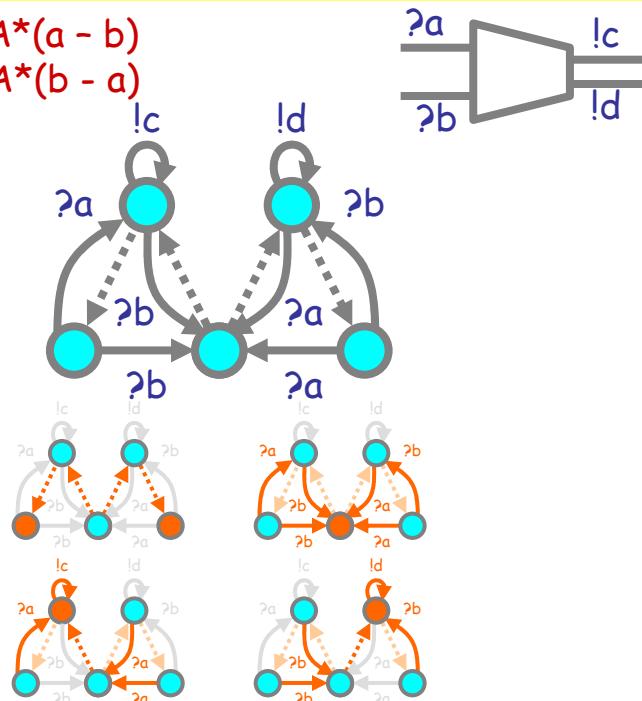
```
let clock@(float, tickchan) = (* sends a tick every t time *)  
(val t = 1/200.0 val d = 1.0/t)  
let step(n)=  
if n>0 then tick; clock(t, tick) else delay@d: step(n-1)  
run step(200)
```

```
let S_a(tickchan) = do la: S_a(tick) or !hc: ()  
let S_b(tickchan) = !hc & S_b(tick)  
and S_b(tickchan) = do lb: S_b(tick) or !hc: S_b2(tick)  
end S_b2(tickchan) = do lb: S_b2(tick) or !hc: S_b2(tick)  
and S_b2(tickchan) = do lb: S_b3(tick) or !hc: ()  
run 10 of (new tickchan run (clock(4.0,tick) | S_a(tick)))  
run 10 of (new tickchan run (clock(2.0,tick) | S_b(tick)))
```

Xor as an Op Amp

$$c = A^*(a - b)$$

$$d = A^*(b - a)$$



```

directive sample 20.0 1000
directive plot ta tb tc ld

new a@0:chan new b@1:chan new c@1:chan new d@1:chan

let Xor_hi_a(chan, b chan, c chan, d chan) =
  do l: Xor_hi_a(chan, b, chan, c, chan) or pb: Xor_lo_ab(a,b,c,d) or delay@1:0; Xor_lo_a(a,b,c,d)
  and Xor_hi_b(chan, b chan, c chan, d chan) =
  do l: Xor_hi_b(chan, b, chan, c, chan) or pb: Xor_lo_ab(a,b,c,d) or delay@1:0; Xor_lo_b(a,b,c,d)
  and Xor_lo_ab(chan, b chan, c chan, d chan) =
  do pb: Xor_hi_ab(b,c,d) or pb: Xor_lo_ab(a,b,c,d)
  and Xor_lo_ab(chan, b chan, c chan, d chan) =
  do pb: Xor_hi_ab(b,c,d) or pb: Xor_lo_ab(a,b,c,d)
  and Xor_lo_ab(chan, b chan, c chan, d chan) =
  do delay@1:0; Xor_hi_ab(a,b,c,d) or delay@1:0; Xor_hi_b(a,b,c,d)

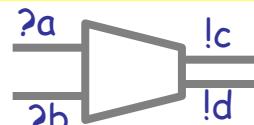
run 50 of (Xor_hi_a(b,c,d) | Xor_hi_b(b,c,d))

let clock(t:float, tickchan) = (* sends a tick every t time *)
  (val ti := t/200.0 val d := 1.0/ti
  let step(n:int) =
    if n<0 then tickchan.clock(t, tickchan) else delay@d; step(n-1)
  run step(200))

let S_a(tickchan) = do l: S_a(tick) or Htick: () and S_b1(tickchan) = Htick: S_b1(tick)
and S_b1(tickchan) = do l: S_b1(tick) or Htick: S_b2(tick)
and S_b2(tickchan) = do l: S_b2(tick) or Htick: S_b3(tick)
and S_b3(tickchan) = Htick: S_b4(tick)
and S_b4(tickchan) = l: S_b4(tick)

run 100 of (new tickchan run (clock@0.0 tick) | S_a(tick)))
run 100 of (new tickchan run (clock@0.0 tick) | S_b(tick)))

```

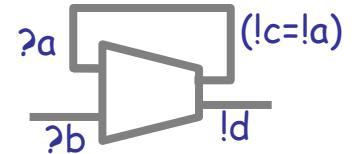


Follower (a standard OpAmp trick)

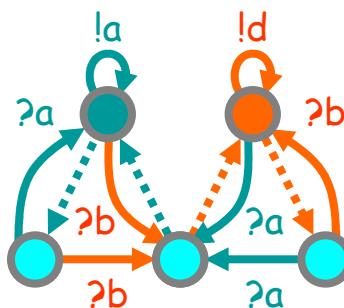
$$\begin{aligned} a=0 \ b=0 &\Rightarrow d=b-a=0 \ a=c=a-b=0 \\ a=0 \ b=1 &\Rightarrow d=b-a=1 \ a=c=a-b=0 \\ a=1 \ b=0 &\Rightarrow d=b-a=0 \ a=c=a-b=1 \\ a=1 \ b=1 &\Rightarrow d=b-a=0 \ a=c=a-b=0 \end{aligned}$$

hence $d=1$ at next step

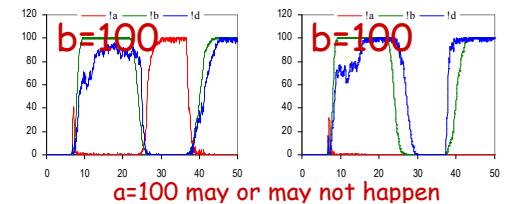
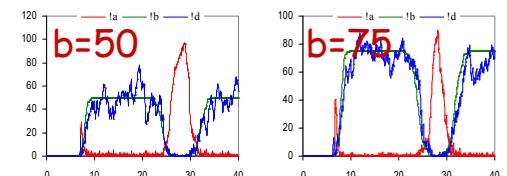
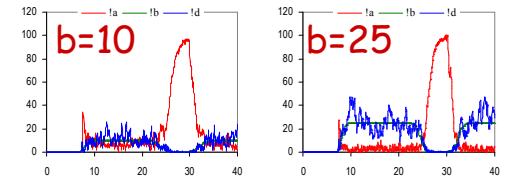
hence $d=b$



"Noninverting Configuration"



$d=b$ analog response!!



$a=100$ may or may not happen

```

directive sample 40.0 1000
directive plot ta tb ld

new a@0:chan new b@1:chan new d@1:chan

let Xor_hi_a(chan, b chan, c chan, d chan) =
  do l: Xor_hi_a(chan, b, chan, c, chan) or pb: Xor_lo_ab(a,b,c,d) or delay@1:0; Xor_lo_a(a,b,c,d)
  and Xor_hi_b(chan, b chan, c chan, d chan) =
  do l: Xor_hi_b(chan, b, chan, c, chan) or pb: Xor_lo_ab(a,b,c,d) or delay@1:0; Xor_lo_b(a,b,c,d)
  and Xor_lo_ab(chan, b chan, c chan, d chan) =
  do pb: Xor_hi_ab(b,c,d) or pb: Xor_lo_ab(a,b,c,d)
  and Xor_lo_ab(chan, b chan, c chan, d chan) =
  do pb: Xor_hi_ab(b,c,d) or pb: Xor_lo_ab(a,b,c,d)
  and Xor_lo_ab(chan, b chan, c chan, d chan) =
  do delay@1:0; Xor_hi_ab(a,b,c,d) or delay@1:0; Xor_hi_b(a,b,c,d)

run 50 of (Xor_hi_a(b,c,d) | Xor_hi_b(b,c,d))

let clock(t:float, tickchan) = (* sends a tick every t time *)
  (val ti := t/200.0 val d := 1.0/ti
  let step(n:int) =
    if n<0 then tickchan.clock(t, tickchan) else delay@d; step(n-1)
  run step(200))

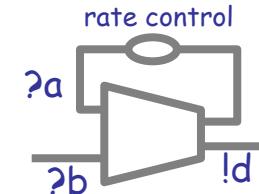
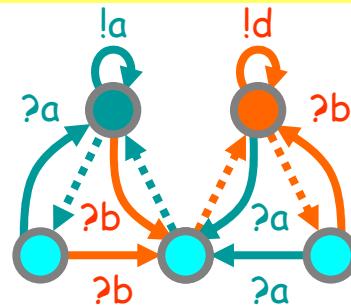
let S_b1(tickchan) = Htick: S_b1(tick)
and S_b2(tickchan) = do l: S_b2(tick) or Htick: S_b3(tick)
and S_b3(tickchan) = do l: S_b3(tick) or Htick: S_b4(tick)
and S_b4(tickchan) = l: S_b4(tick)

run 10 of (new tickchan run (clock@0.0 tick) | S_b(tick)))

```

Changing the OpAmp Gain

An OpAmp provides “infinite” differential amplification, but a stable finite amplification can be obtained by a feedback loop with a load splitter (the follower is a special case of that, which gives gain 1). The equivalent here is simply changing the rate on the feedback link.



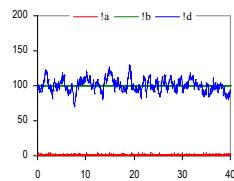
```

directive sample 40.0 1000
directive plot ?a; ?b; !d

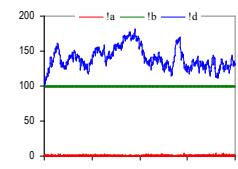
new a@1.0-chan new b@1.0-chan new d@1.0-chan

let Xor_hi_a(chan, bchan, cchan, dchan) =
do !c: Xor_hi_a(b,c,d) or ?b: Xor_lo_a(b,c,d) or delay@1.0: Xor_lo_a(b,c,d)
and Xor_hi_b(chan, bchan, cchan, dchan) =
do !c: Xor_hi_b(b,c,d) or ?a: Xor_lo_b(b,c,d) or delay@1.0: Xor_lo_b(b,c,d)
and Xor_hi_c(chan, bchan, cchan, dchan) =
do !c: Xor_hi_c(b,c,d) or ?b: Xor_lo_c(b,c,d)
and Xor_hi_d(chan, bchan, cchan, dchan) =
do !b: Xor_hi_d(b,c,d) or ?a: Xor_lo_d(b,c,d)
and Xor_lo_ab(chan, bchan, cchan, dchan) =
do delay@1.0: Xor_hi_ab(b,c,d) or delay@1.0: Xor_lo_ab(b,c,d)
run 100 of Xor_lo_a(b,c,d) | Xor_hi_b(b,c,d)
run 100 of replicate !b
  
```

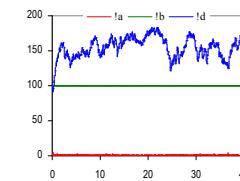
$$[d] = [b]/\text{rate}(a)$$



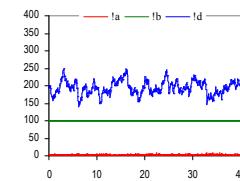
$b=100$
 $a@1.0$
 d gain 1.0
#OpAmp=200



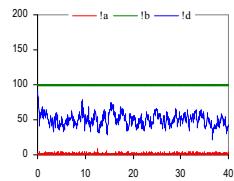
$b=100$
 $a@0.75$
 d gain 1.33
#OpAmp=200



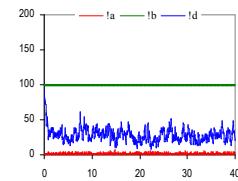
$b=100$
 $a@0.6$
 d gain 1.66
#OpAmp=200



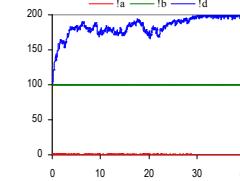
$b=100$
 $a@0.5$
 d gain 2.00
#OpAmp=400
(non saturated)



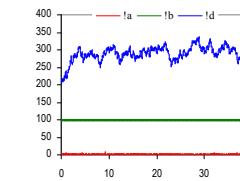
$b=100$
 $a@2.0$
 d gain 0.5
#OpAmp=200



$b=100$
 $a@4.0$
 d gain 0.25
#OpAmp=200



$b=100$
 $a@0.5$
 d gain 2.00
#OpAmp=200
(saturated)



$b=100$
 $a@0.33$
 d gain 3.00
#OpAmp=400

2006-04-21



Bidirectional Polymerization

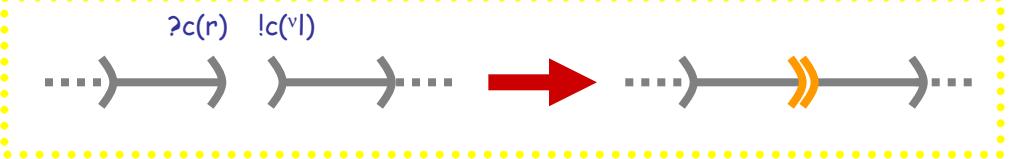
new $c @ \mu$ new stop@1.0

$A_{\text{free}} =$
 $(\text{new } rht @ \lambda; !c(rht); A_{\text{brht}}(rht))$
 $+ ?c(lft); A_{\text{blft}}(lft)$

$A_{\text{blft}}(lft) =$
 $(\text{new } rht @ \lambda; !c(rht); A_{\text{bound}}(lft, rht))$

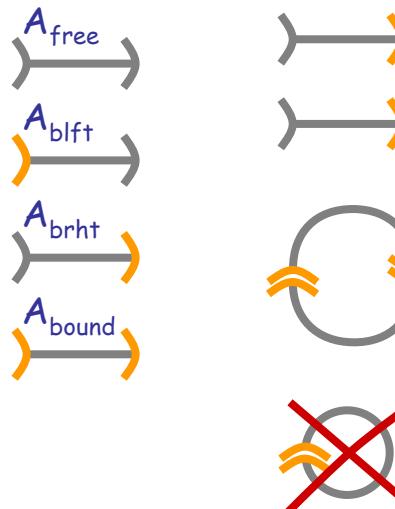
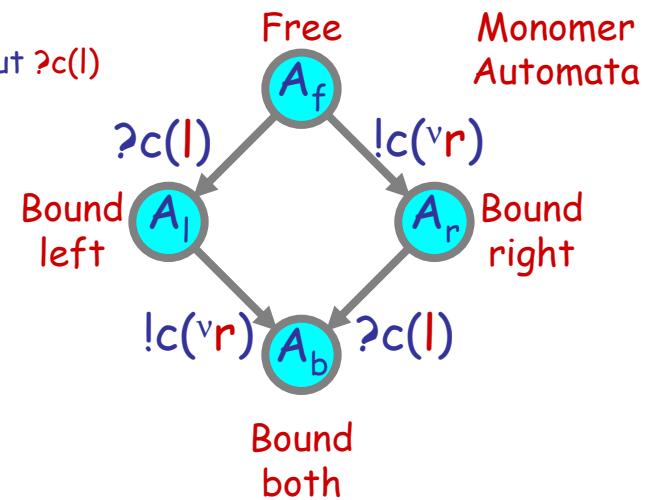
$A_{\text{brht}}(rht) =$
 $?c(lft); A_{\text{bound}}(lft, rht)$

$A_{\text{bound}}(lft, rht) = ?\text{stop}$



Communicating Automata

Bound output $!c(r)$ and input $?c(l)$
on automata transitions
to model complexation



```

directive sample 10000.0
directive plot Afree(), AblfH(), Abrht(), Abound()

val lam : 1.0 val mu = 1.0
new @muchan(chan) new stop@1.0 chan

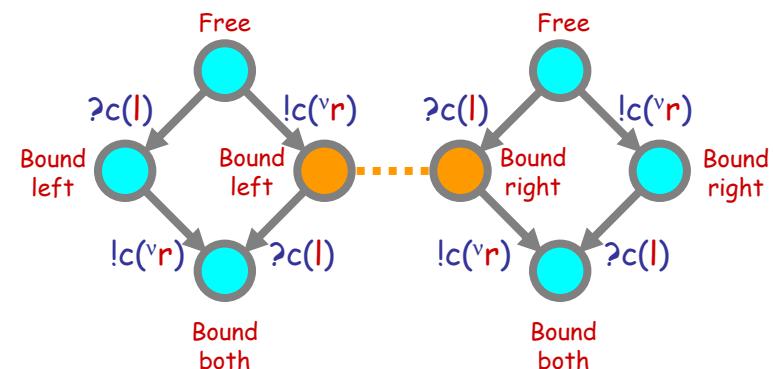
let A_free() =
  (new rht@lam chan run
  do !c(rht); Abrht(rht)
  or ?c(lft); Ablf(lft))

and Ablf(lft chan) =
  (new rht@lam chan run
  !c(rht); Abound(lft, rht))

and Abrht(rht chan) =
  ?c(lft); Abound(lft, rht)

and Abound(lft chan, rht chan) =
  ?stop.

run (2 of Afree())
  
```



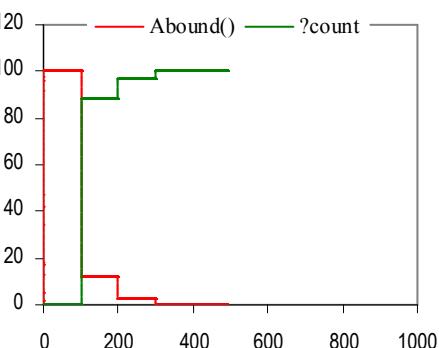
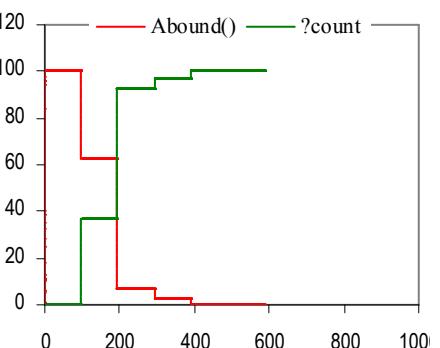
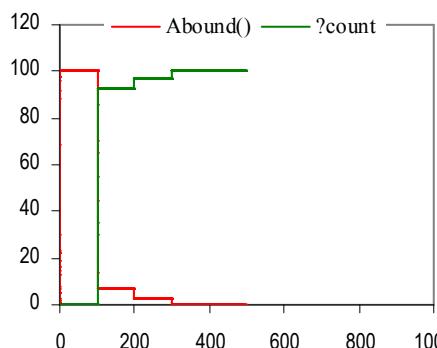
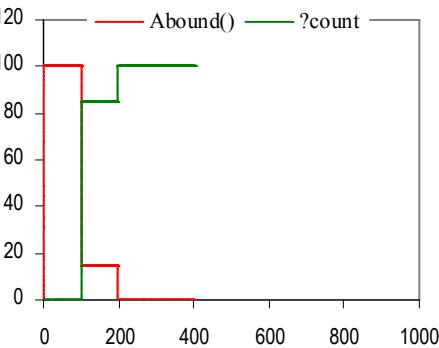
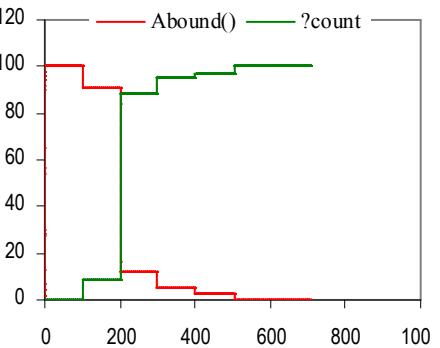
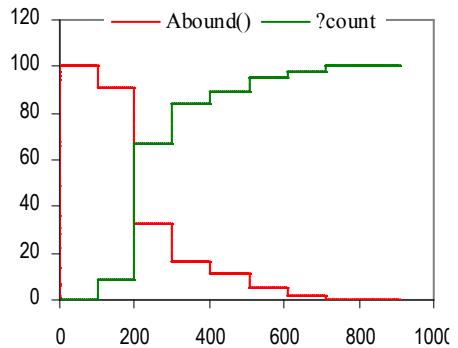
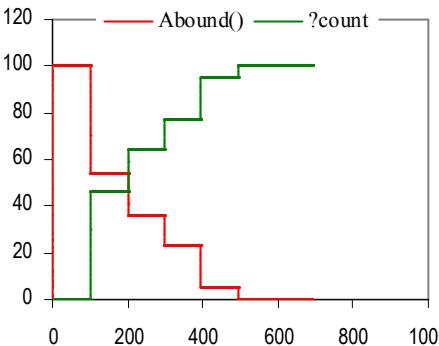
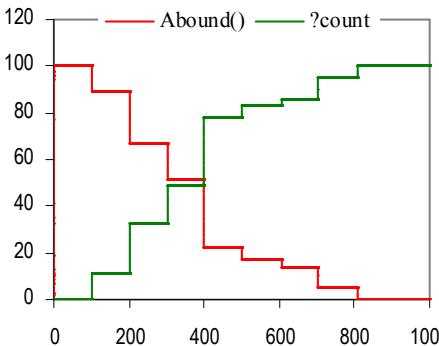
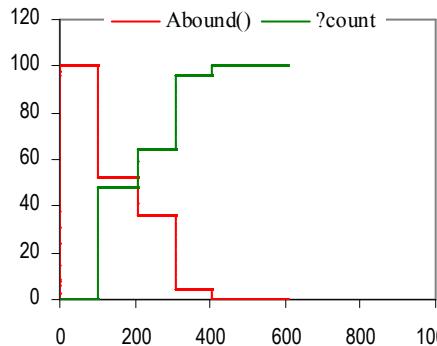
2006-04-21

Bidirectional Polymerization

Circular Polymer Lengths

Scanning and counting the size of the circular polymers (by a cheap trick).

Polymer formation is complete within 10t; then a different polymer is scanned every 100t.



```

directive sample 1000.0
directive plot Abound(); ?count

type Link = chan(chan)
type Barb = chan

val lam = 1000.0 (* set high for better counting *)
val mu = 1.0
new c@mu:chan(Link)
new enter@lam:chan(Barb)
new count@lam:Barb

let Afree() =
  (new rht@lam:Link run
  do lc(rht); Abrht(rht)
  or ?c(lft); Ablf(lft))

and Ablf(lft:Link) =
  (new rht@lam:Link run
  lc(rht); Abound(lft,rht))

and Abrht(rht:Link) =
  ?c(lft); Abound(lft,rht)

and Abound(lft:Link, rht:Link) =
  do ?enter(barb); (barb | !rht(barb))
  or ?lft(barb); (barb | !rht(barb))
(* each Abound waits for a barb, exhibits it, and passes it to
the right so we can plot number of Abound in a ring*)

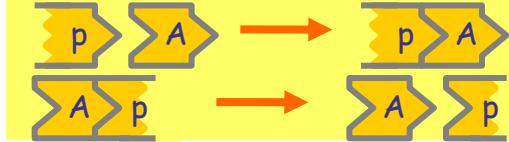
let clock(t:float, tick:chan) = (* sends a tick every t time *)
  (val ti = t/1000.0 val d = 1.0/ti
  let step(n:int) =
    if n>0 then !tick; clock(t,tick) else delay@d; step(n-1)
  run step(1000))

new tick:chan
let Scan() = ?tick; lenter(count); Scan()

run 100 of Afree()
run (clock(100.0, tick) | Scan())

```

100xA_{free}, initially.
The height of each rising
step is the size of a
separate circular polymer.
(Unbiased sample of nine
consecutive runs.)



Actin-like Poly/Depolymerization

new $c @ \mu$

$$A_{\text{free}} = (\text{new } lft @ \lambda; !c(lft); A_{\text{blft}}(lft)) + ?c(rht); A_{\text{brht}}(rht)$$

$$A_{\text{blft}}(lft) =$$

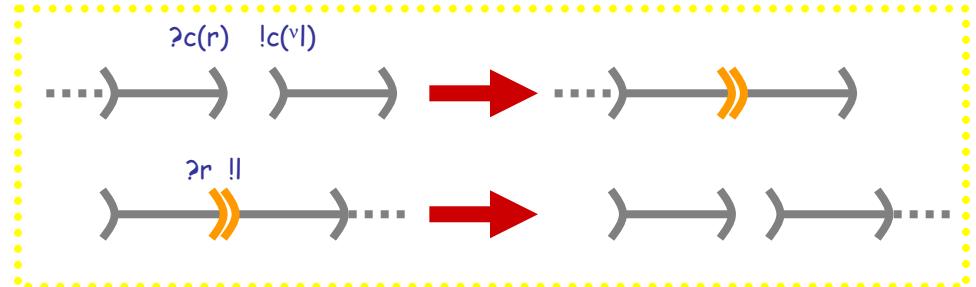
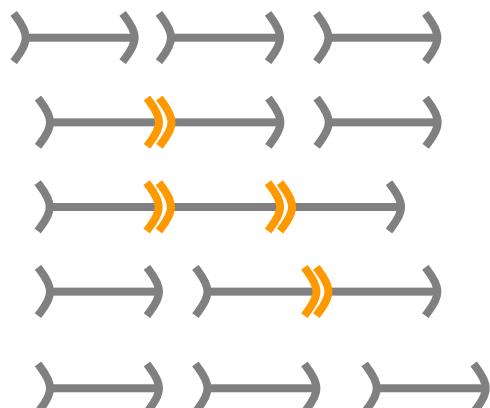
$$!lft; A_{\text{free}} + ?c(rht); A_{\text{bound}}(lft, rht)$$

$$A_{\text{brht}}(rht) =$$

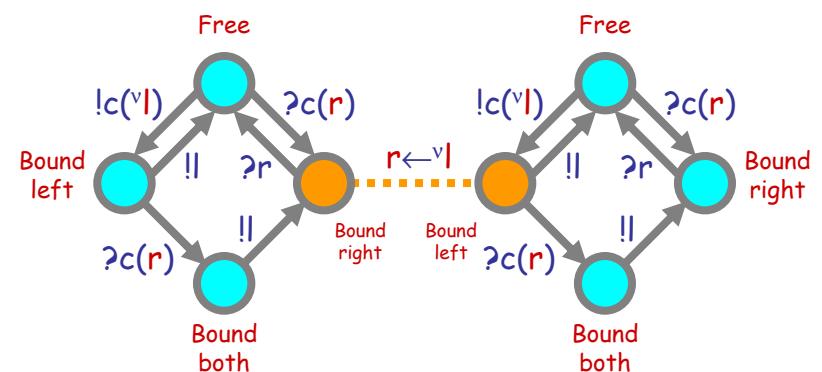
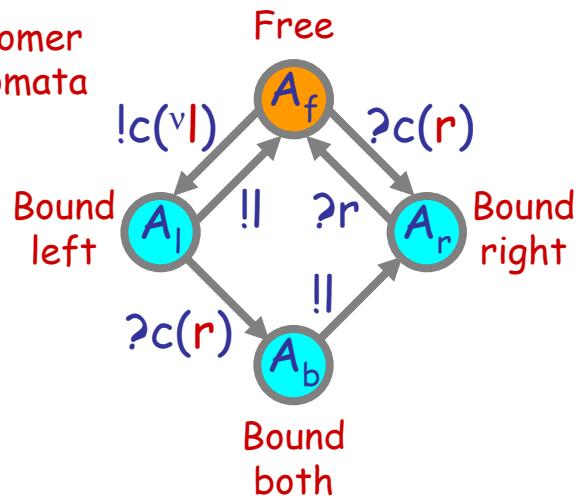
$$?rht; A_{\text{free}}$$

$$A_{\text{bound}}(lft, rht) =$$

$$!lft; A_{\text{brht}}(rht)$$



Monomer
Automata

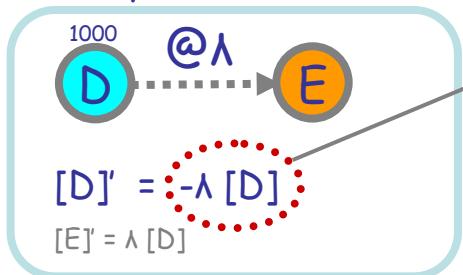


The Law of Mass Interaction

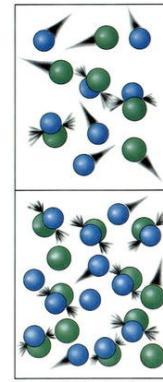
Law of Mass Interaction

The speed of interaction[†] is proportional to the number of possible interactions.

Decay



Exponential Decay law
Rate of change proportional to number of possible decays.



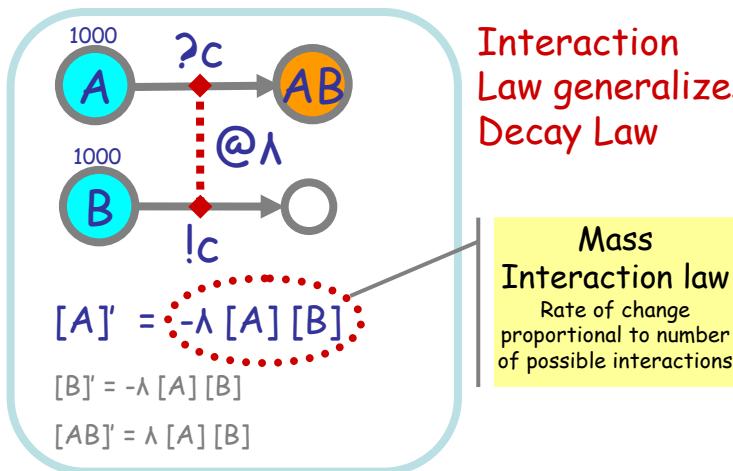
Chemical Law of Mass Action
http://en.wikipedia.org/wiki/Chemical_kinetics
The speed of a chemical reaction is proportional to the activity of the reacting substances.

Activity = concentration, for well-stirred aqueous medium

Concentration = number of moles per liter of solution

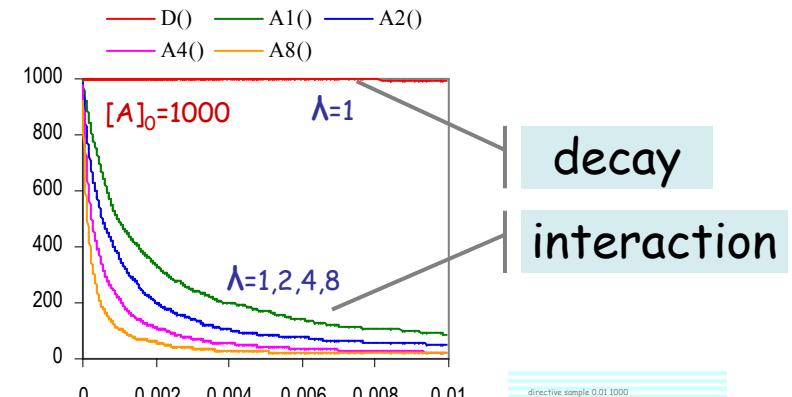
Mole = 6.022141×10^{23} particles

Mass interaction



Interaction Law generalizes Decay Law

Mass Interaction law
Rate of change proportional to number of possible interactions



```
directive sample 0.01 1000
directive plot D(); A1(); A2(); A4(); AB()
new c@0.1: char() new c@0.2: char()
new c@0.4: char() new c@0.8: char()

let D() = delay@1.0
let A1() = x1 and B1() = k1
let A2() = x2 and B2() = k2
let A4() = x4 and B4() = k4
let AB() = x8 and BB() = k8

run 1000 of (D() | A1() | B1() | A2()
| B2() | A4() | B4() | AB() | BB())
2006-04-21
```

[†] speed of interaction (formally definable)
= number of interactions over time

not proportional to the number of interacting processes!

[P] is the number of processes P (this is informal; it is only meaningful for a set of processes offering a given action, but a set of such processes can be counted and plotted)

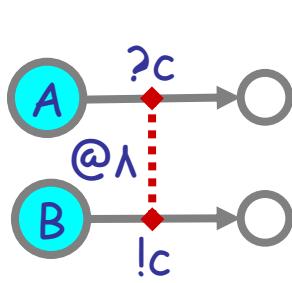
Activity and Speed

stochastic algebras disagree!

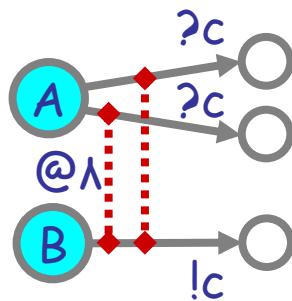
The speed of interaction is proportional to the number of possible interactions.

- = The activity (= "concentration") on a channel is the number of possible interactions on that channel.

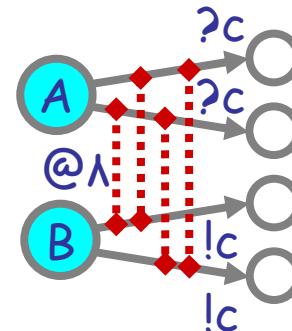
The speed of interaction on a channel, is the activity multiplied by the base rate of the channel.



c activity: 1
speed: λ



c activity: 2
speed: 2λ



c activity: 4
speed: 4λ

The mass interaction law [Buchholz] [Priami-Regev-Shapiro-Silverman] is compatible with chemistry [Gillespie] and incompatible with any other stochastic algebra in the literature! (including [Priami]; see [Hermanns])

Other algebras assign rates to actions, not channels, with speed laws:
 $2\lambda * 2\lambda = 4\lambda^2$
 $\max(2\lambda, 2\lambda) = 2\lambda$ [Goetz]
 $\min(2\lambda, 2\lambda) = 2\lambda$ [Priami]
 $1/(1/(2\lambda)+1/(2\lambda)) = \lambda$ [PEPA]
 $2\lambda * 1 = 2\lambda$ (passive inputs)

```
directive sample 0.01 10000
directive plot A1(); A2(); A3()

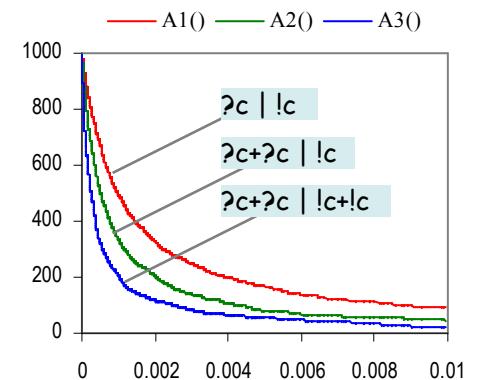
new c1@1.0:chan
new c2@1.0:chan
new c3@1.0:chan

let A1() = ?c1
and B1() = !c1

let A2() = do ?c2 or ?c2
and B2() = !c2

let A3() = do ?c3 or ?c3
and B3() = do !c3 or !c3

run 1000 of (A1() | B1())
| A2() | B2() | A3() | B3())
```



Possible Interactions

The speed of interaction is proportional to the number of possible interactions.

But a process cannot interact with itself.

Assume each process P is in restricted-sum-normal-form. For each channel x:

$$In(x, P) = \text{Num of active } ?x \text{ in } P$$

$$Out(x, P) = \text{Num of active } !x \text{ in } P$$

$$Mix(x, P) = In(x, P) * Out(x, P)$$

#interactions that cannot happen
in a given summation P

$$In(x) = \text{Sum P of } In(x, P)$$

$$Out(x) = \text{Sum P of } Out(x, P)$$

$$Mix(x) = \text{Sum P of } Mix(x, P)$$

total #interactions that cannot happen

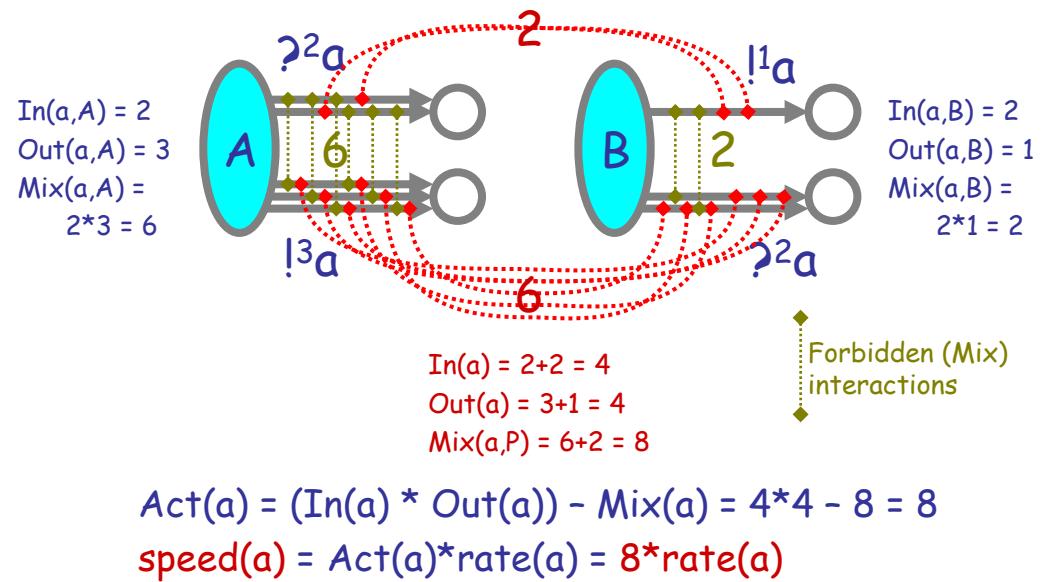
The global **Activity** on channel x:

$$Act(x) = (In(x) * Out(x)) - Mix(x)$$

total cross product of inputs and outputs
minus total #interactions that cannot happen

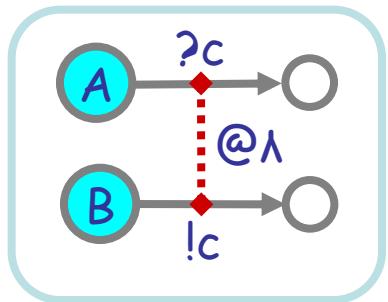
The global **speed** of interaction on a channel x:

$$speed(x) = Act(x) * rate(x)$$



Deriving Back Interaction Laws

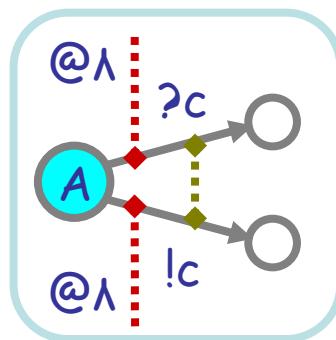
The mass action law:



$$\begin{aligned}[A]' &= -\text{speed}(c) = -\lambda \text{Act}(c) \\ \text{Act}(c) &= (\text{In}(c) * \text{Out}(c)) - \text{Mix}(c) \\ &= ([A] * [B]) - 0\end{aligned}$$

hence $[A]' = -\lambda [A][B]$

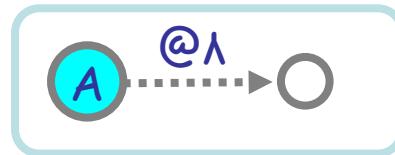
The mixed interaction law:



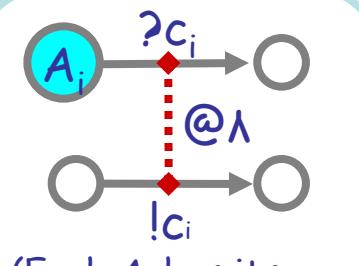
$$\begin{aligned}[A]' &= -\text{speed}(c) = -\lambda \text{Act}(c) \\ \text{Act}(c) &= (\text{In}(c) * \text{Out}(c)) - \text{Mix}(c) \\ &= ([A] * [A]) - [A] = [A] * ([A]-1)\end{aligned}$$

hence $[A]' = -\lambda [A] ([A]-1)$

The decay law:



= def



(Each A_i has its own private channel c_i)

$$\begin{aligned}[A]' &= \sum(c_i) -\text{speed}(c_i) \\ &= \sum(c_i) -\lambda \text{Act}(c_i) \\ \text{Act}(c_i) &= (\text{In}(c_i) * \text{Out}(c_i)) - \text{Mix}(c_i) \\ &= (1 * 1) - 0 = 1\end{aligned}$$

hence $[A]' = -\lambda [A]$

$\text{Act}(x) = (\text{In}(x) * \text{Out}(x)) - \text{Mix}(x)$

Conclusions

Conclusions

- Stochastic Collectives
 - Complex global behavior from simple components
 - Emergence of collective functionality from “non-functional” components
 - (Cf. “swarm intelligence”: simple global behavior from complex components)
- Artificial Biochemistry
 - Stochastic collectives with Law of Mass Interaction kinetics
 - Connections to classical Markov theory, chemical Master Equation, and Rate Equation
- The agent/automata/process point of view
 - “Individuals” that transition between states (vs. transmutation between “unrelated” chemical species)
 - More appropriate for Systems Biology
 - Stochastic π -calculus (SPiM) for investigating stochastic collectives
 - Restriction+Communication \Rightarrow Polymerization: FSA that “stick together”