The auxiliary region method for coupling **PDE and Brownian**based dynamics for reaction-diffusion systems

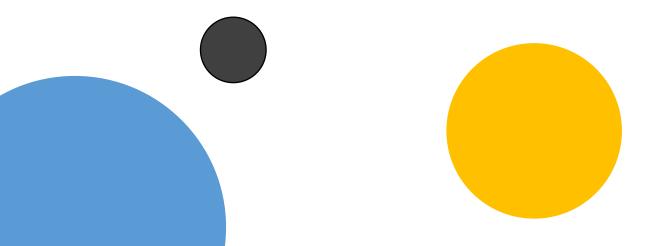


Cameron Smith (Cohort 3)

SAMBa Conference 2019 08/07/19

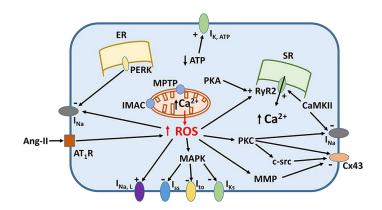


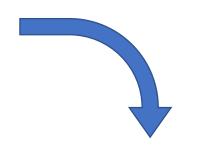
- Reaction-diffusion systems
- Spatially extended hybrid methods
- The auxiliary region method (ARM)

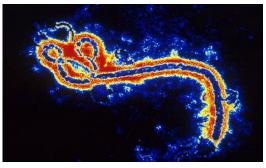


### Reaction-diffusion systems (SAMBa)

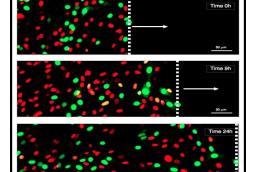


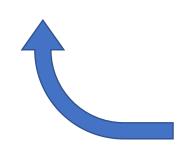




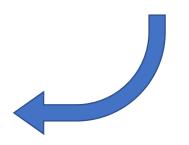


#### Reactiondiffusion

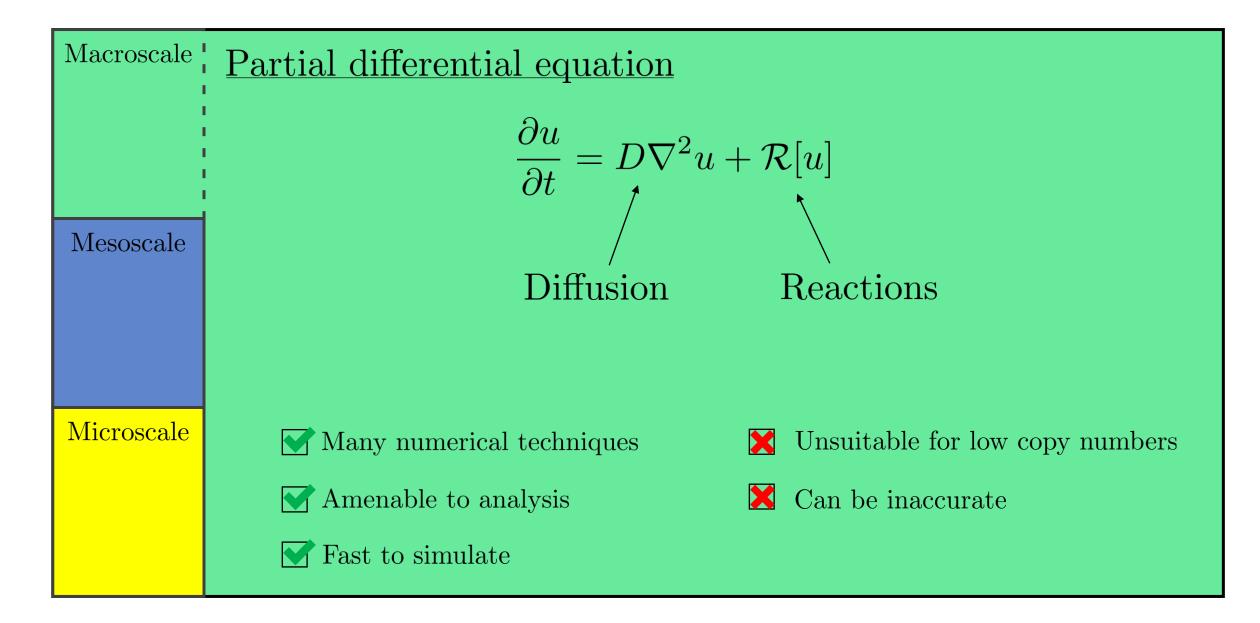


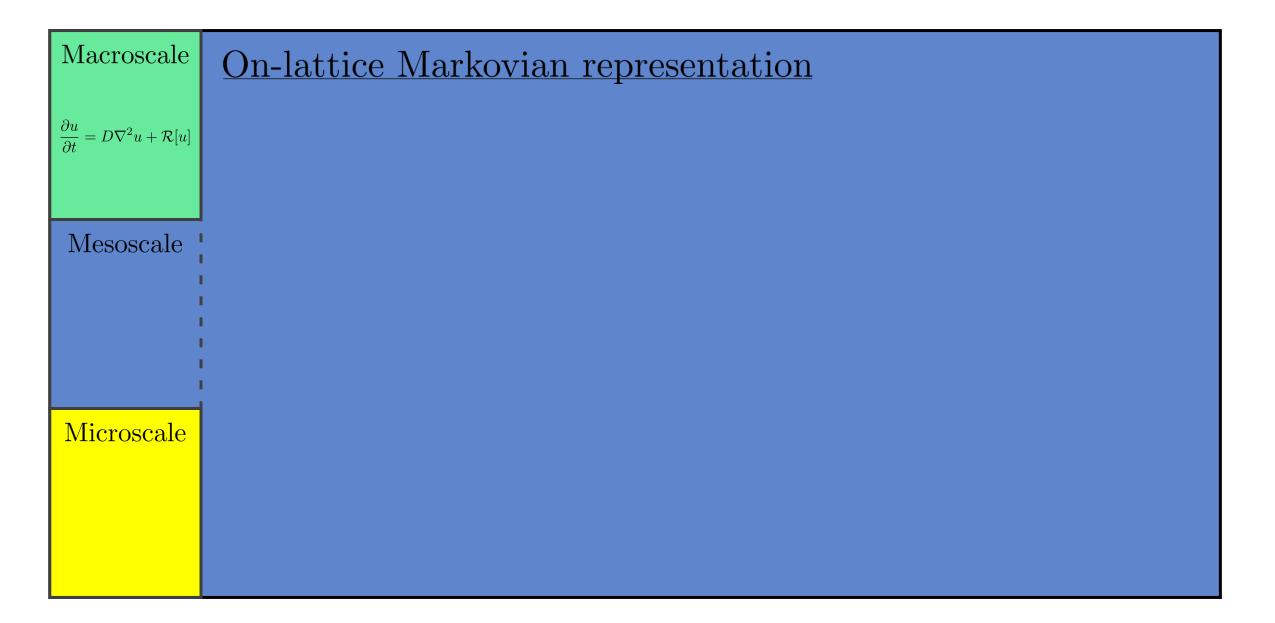


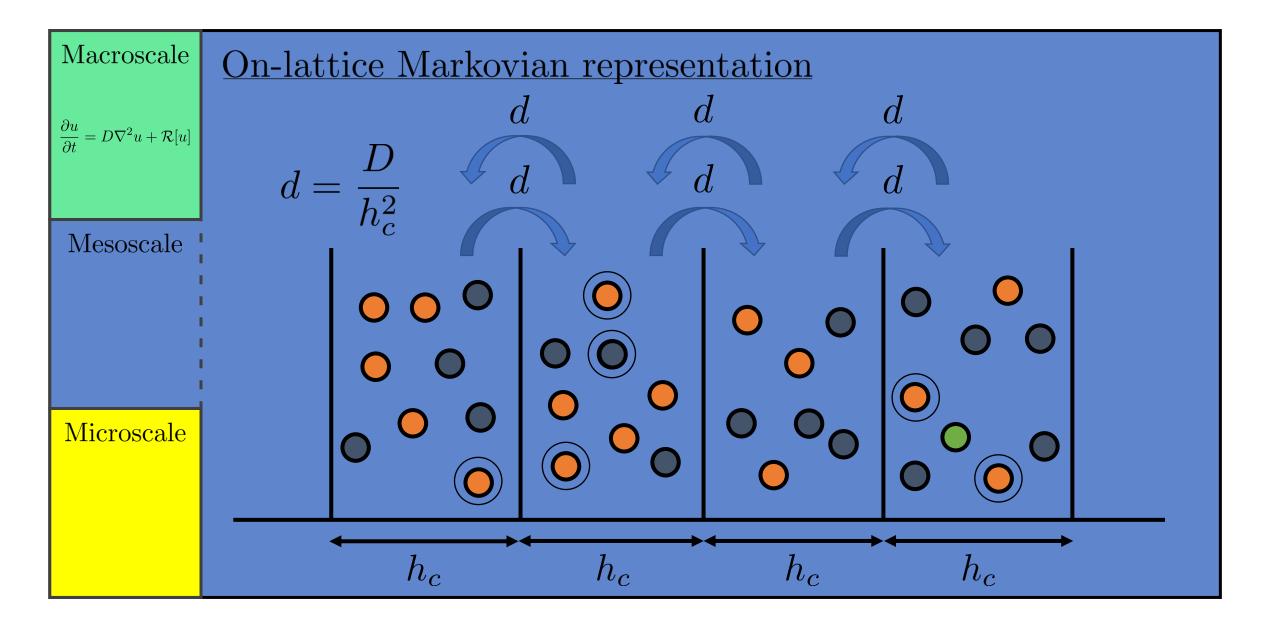


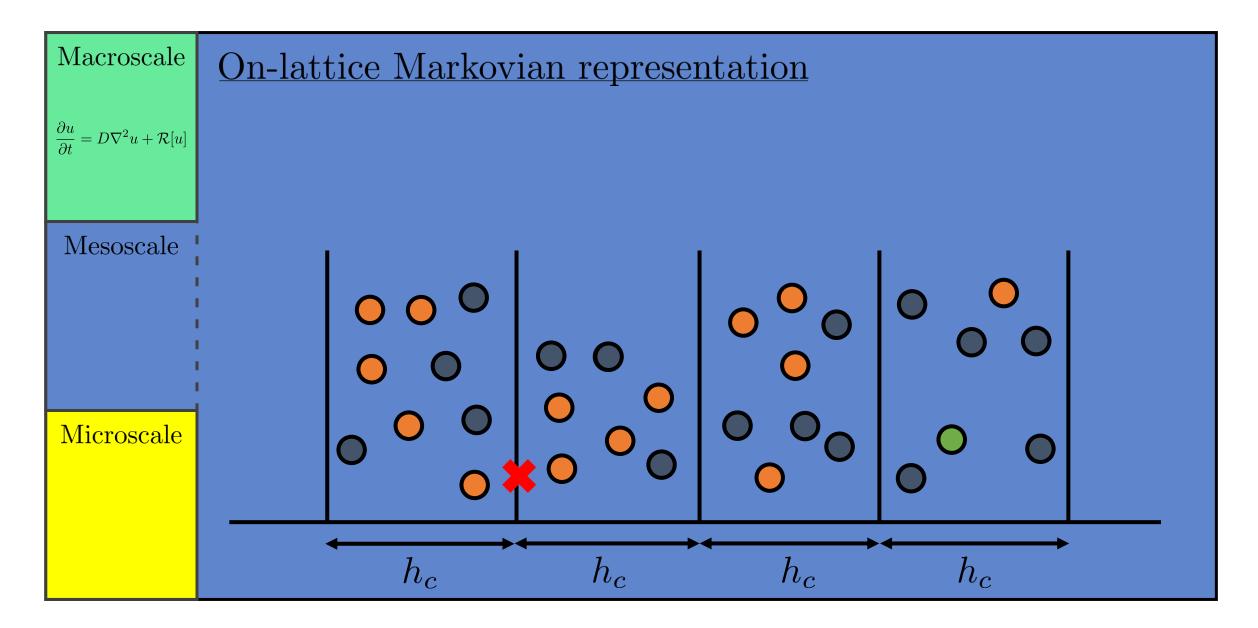


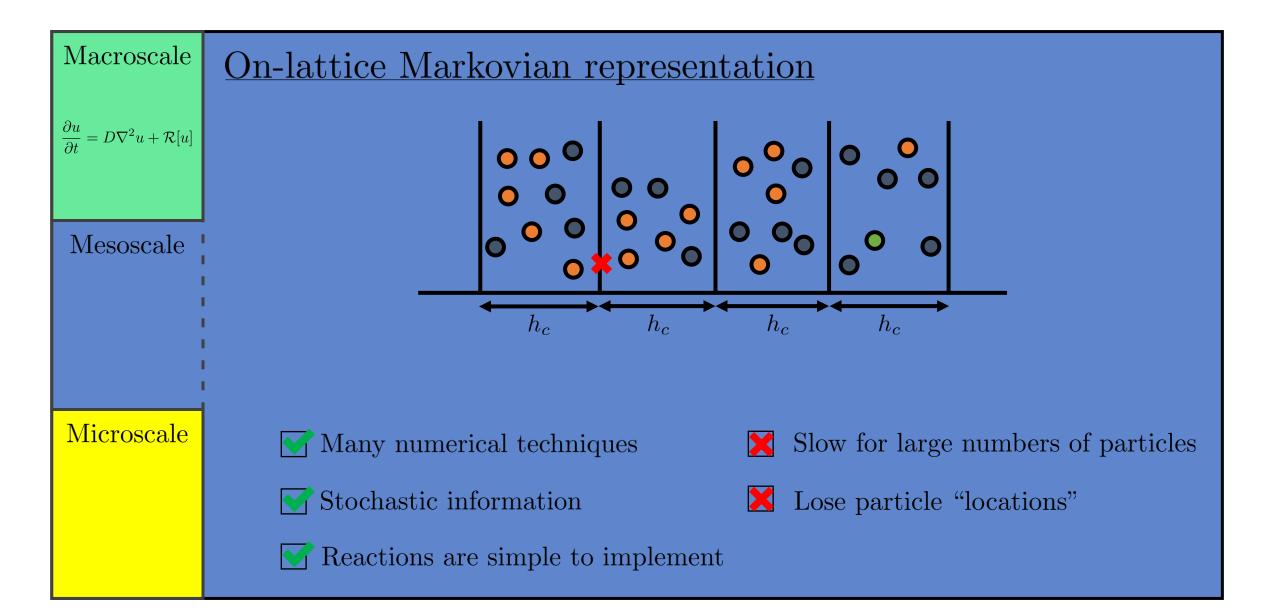


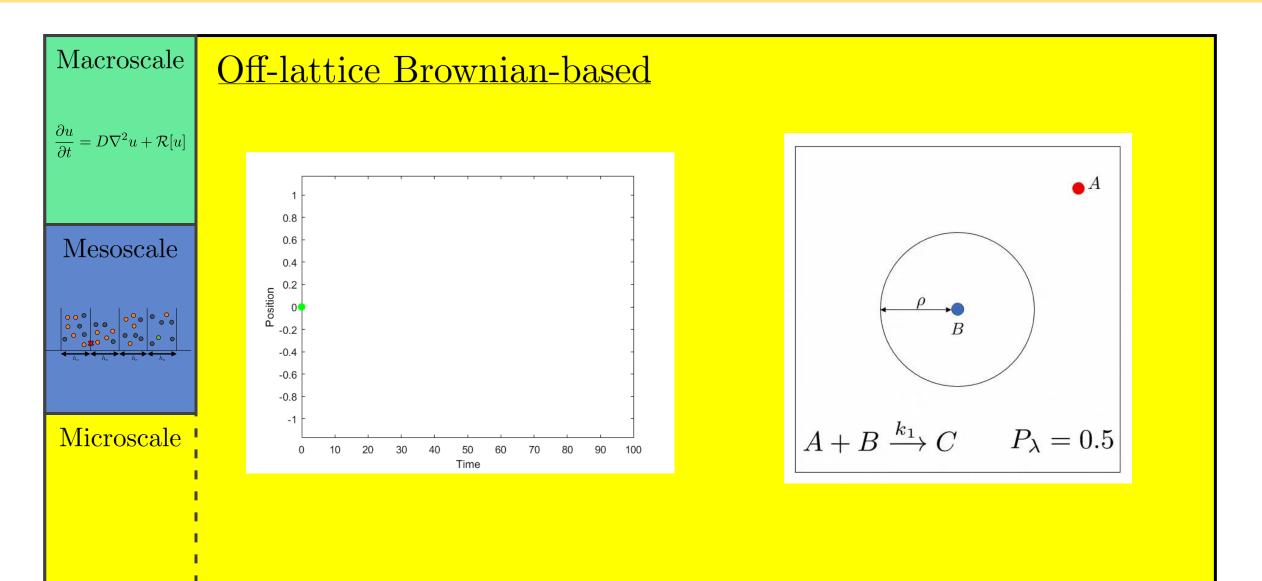


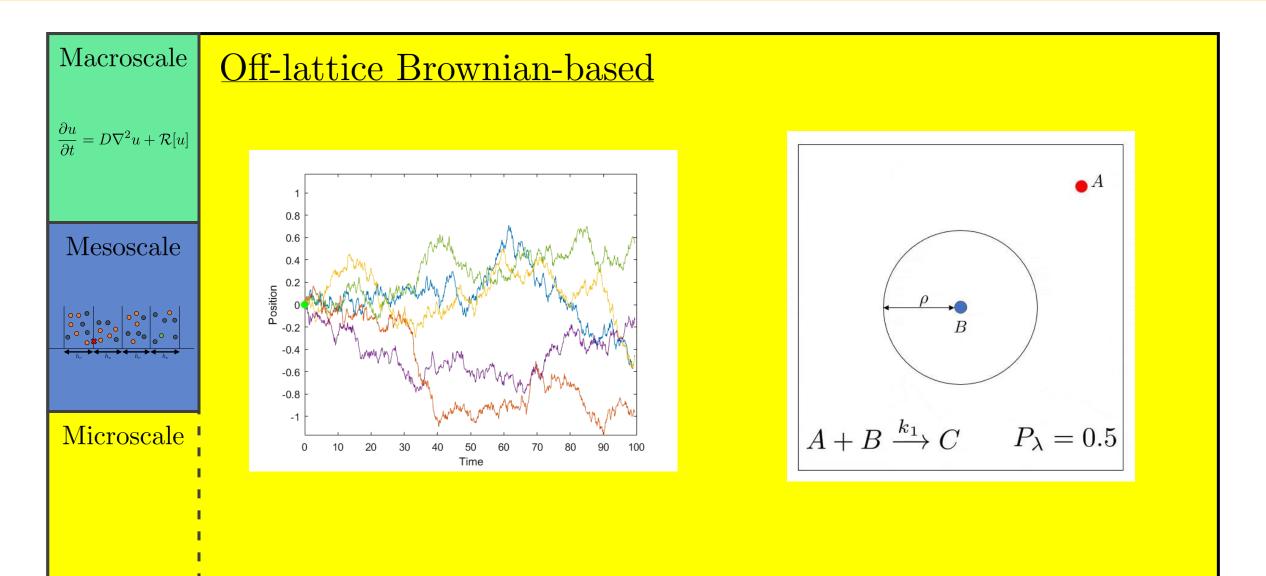


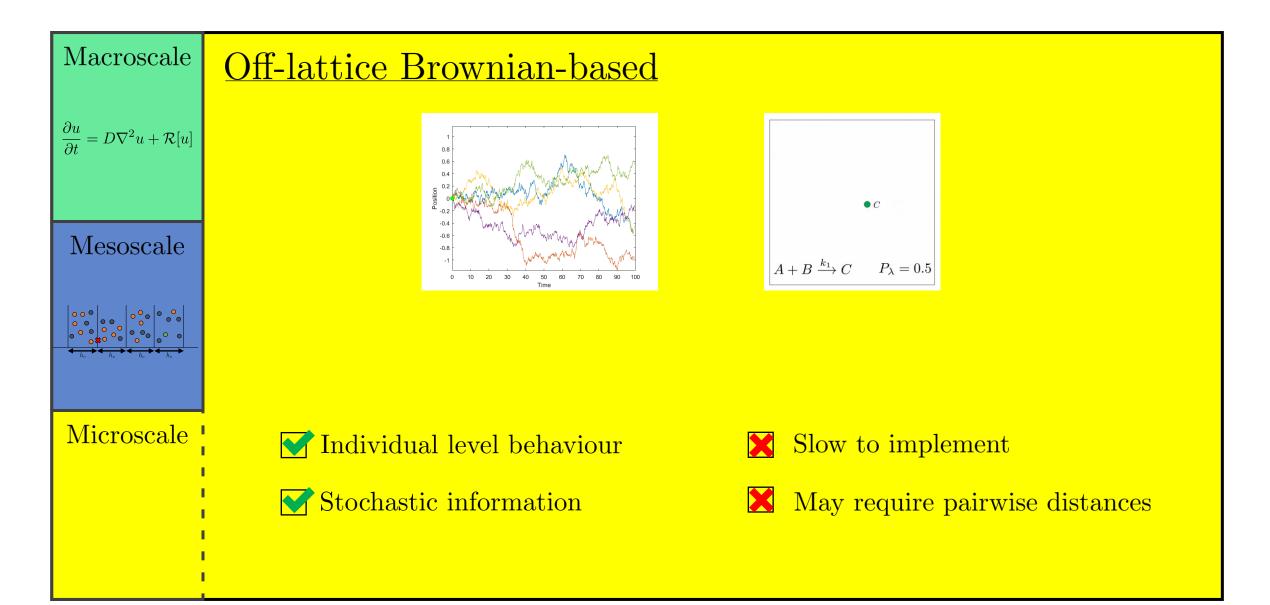


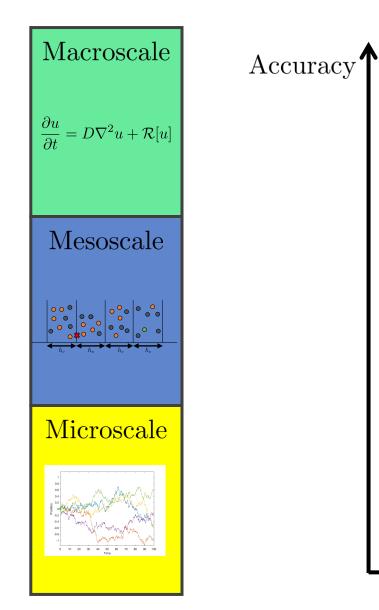


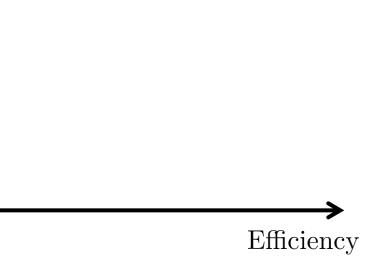


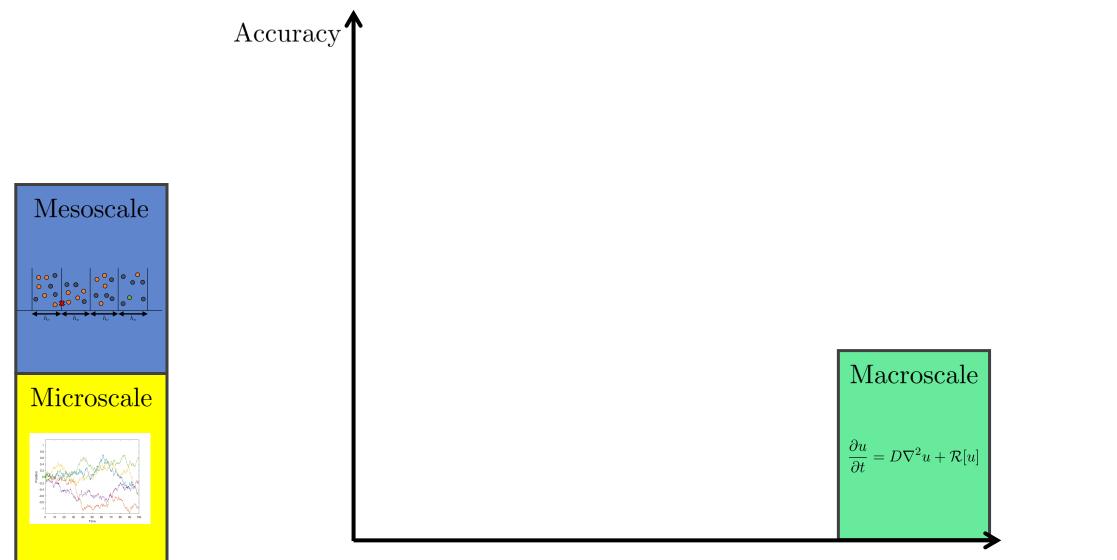




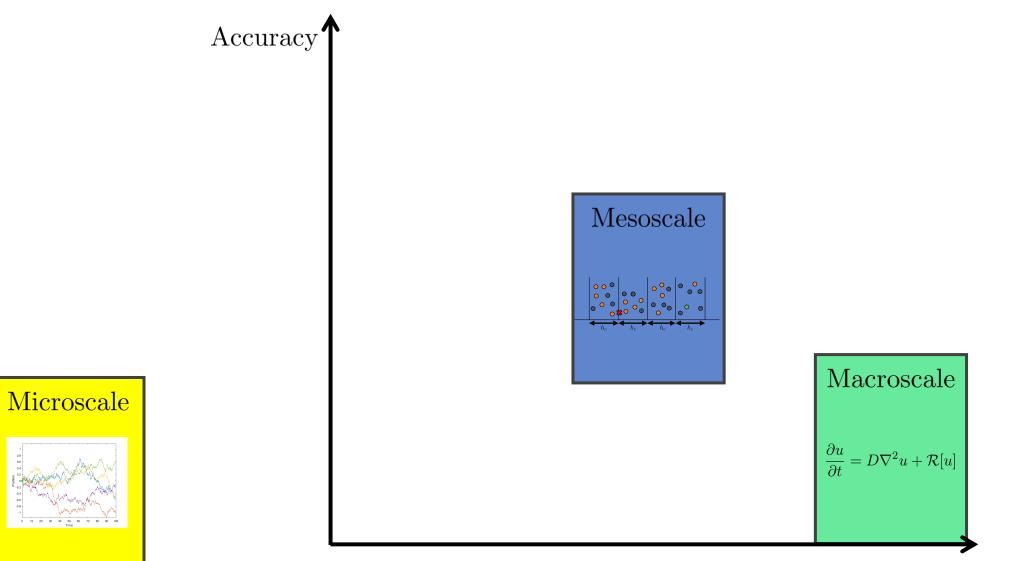




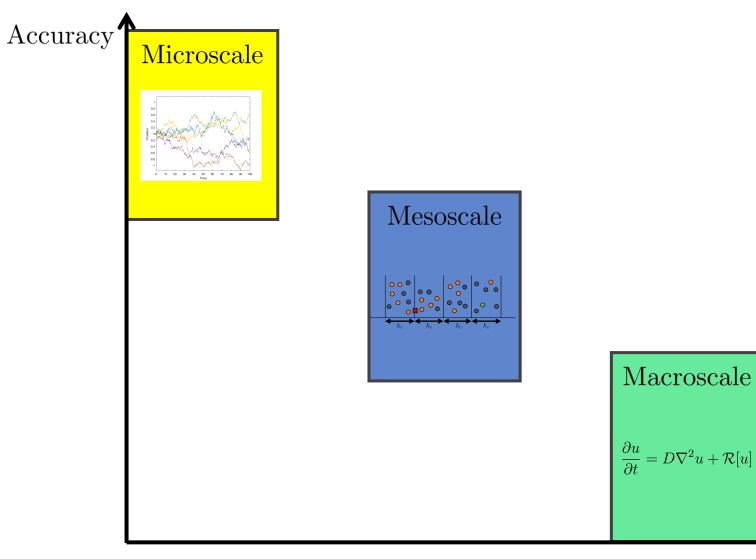




Efficiency



Efficiency



Efficiency

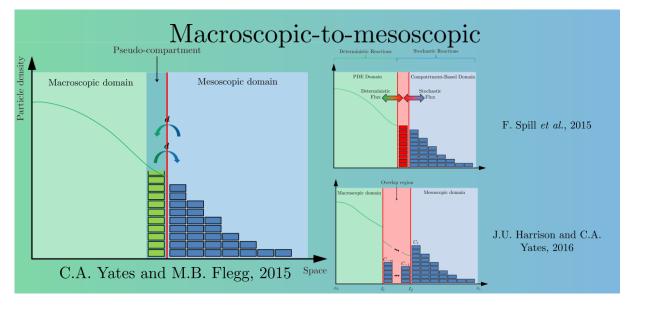


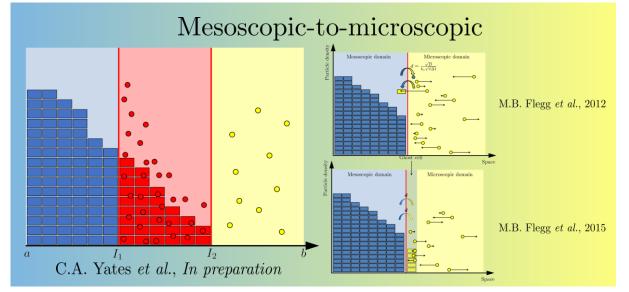


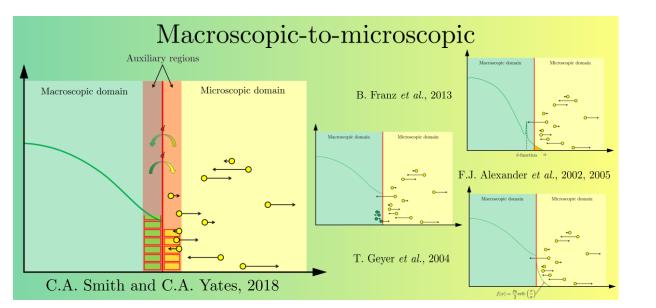
... spatially extended hybrid methods employ different modelling paradigms at different scales in order to compliment the strengths and negate the weaknesses of each.

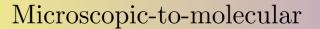
> C.A. Smith and C.A. Yates, 2018 Spatially extended hybrid methods: a review

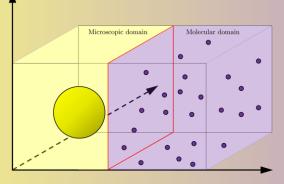
#### Many, many examples!









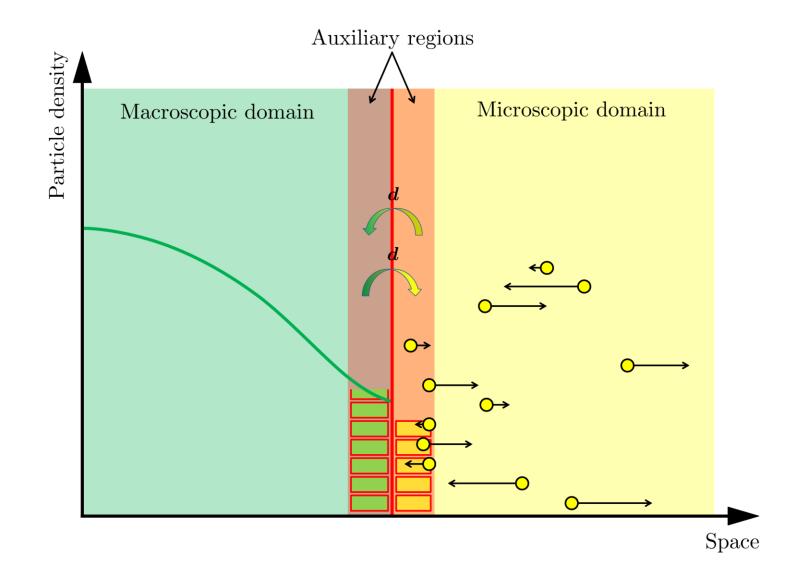


R. Erban, 2014

# The auxiliary region method (SAMBa



#### The auxiliary region method

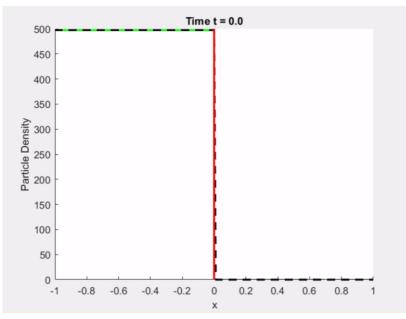


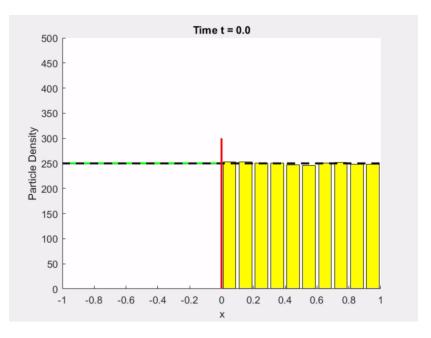
#### **Basic algorithm**

### Let the PDE/Brownian update step be $\Delta t$ , and $t_{\Delta}$ be the next PDE/Brownian update time.

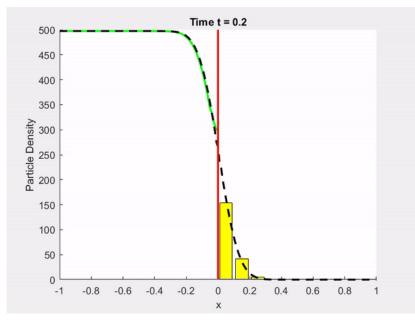
- 1) Find the time until the next event within the auxiliary regions occurs. Call this  $t_a$ .
- 2) If this is less than the time until the next PDE/Brownian update (i.e. if  $t_a < t_{\Delta}$ ), find the corresponding event and enact it.
- 3) Otherwise (i.e. if  $t_{\Delta} < t_a$ ), evolve the PDE and Brownian subdomains.
- 4) Update time and return to step 1.

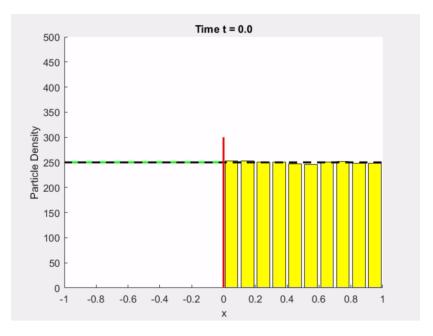
# Pure diffusion $\begin{aligned} \frac{\partial u}{\partial t} &= D\nabla^2 u \\ \frac{\partial u}{\partial x}\Big|_{x=-1} &= 0 \text{ and } \left. \frac{\partial u}{\partial x} \right|_{x=1} &= 0 \\ u(x,0) &= \begin{cases} 500 & x \in [-1,0) \\ 0 & x \in [0,1] \end{cases} \end{aligned}$



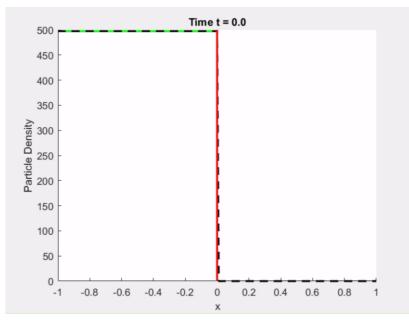


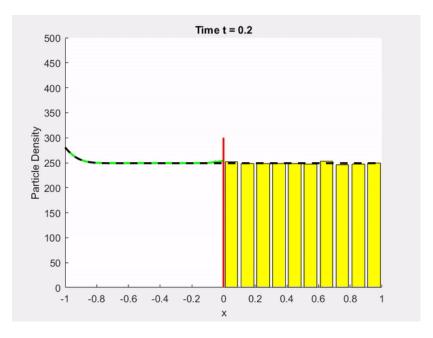
# Pure diffusion $\begin{aligned} \frac{\partial u}{\partial t} &= D\nabla^2 u \\ \frac{\partial u}{\partial x}\Big|_{x=-1} &= 0 \text{ and } \left. \frac{\partial u}{\partial x} \right|_{x=1} &= 0 \\ u(x,0) &= \begin{cases} 500 & x \in [-1,0) \\ 0 & x \in [0,1] \end{cases} \end{aligned}$



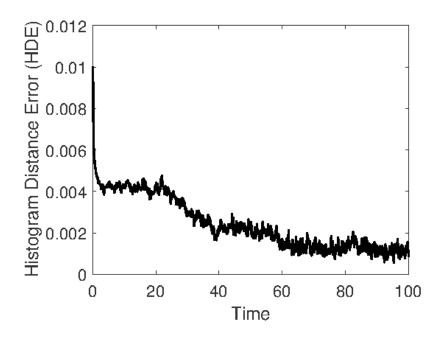


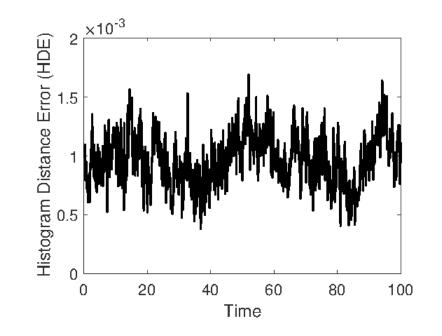
# Pure diffusion $\begin{aligned} \frac{\partial u}{\partial t} &= D\nabla^2 u \\ \frac{\partial u}{\partial x}\Big|_{x=-1} &= 0 \text{ and } \left. \frac{\partial u}{\partial x} \right|_{x=1} &= 0 \\ u(x,0) &= \begin{cases} 500 & x \in [-1,0) \\ 0 & x \in [0,1] \end{cases} \end{aligned}$





### Pure diffusion $\begin{aligned} \frac{\partial u}{\partial t} &= D\nabla^2 u \\ \frac{\partial u}{\partial x}\Big|_{x=-1} &= 0 \text{ and } \left. \frac{\partial u}{\partial x} \right|_{x=1} &= 0 \\ u(x,0) &= \begin{cases} 500 & x \in [-1,0) \\ 0 & x \in [0,1] \end{cases} \end{aligned}$





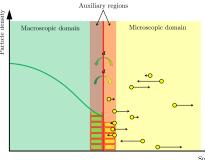
Reaction-diffusion systems may be modelled in different ways, each with complimentary advantages and disadvantages.

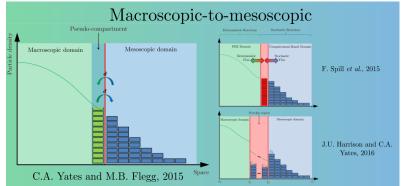
### Summary

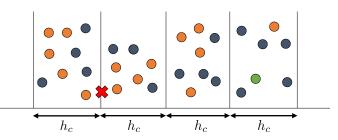
Macroscopic-to-mesoscopic Mesoscopic domain F. Spill et al., 2015

Hybrid methods combine these to form accurate and efficient methods. There are many examples.

auxiliary region method combines PDE and The Brownian-based approaches. Able to simulate reactiondiffusion systems accurately.







# Thank you for your attention.

### **Any questions?**

This work is joint with Kit Yates.





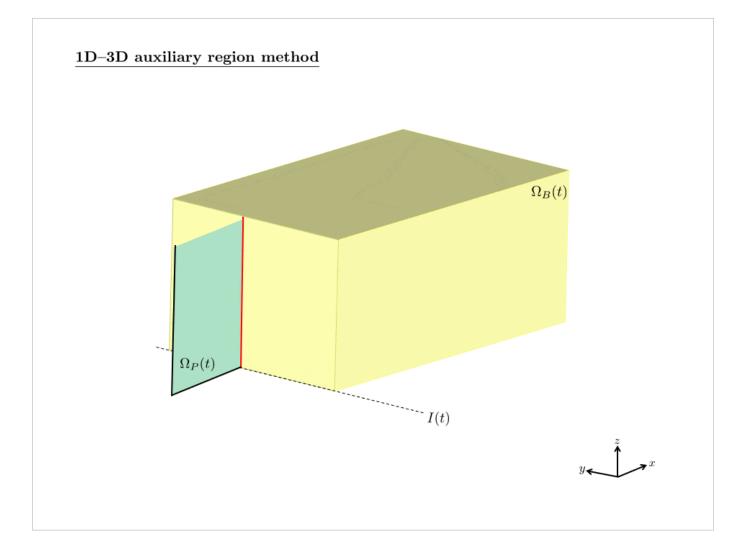
#### Get in touch:

@C\_A\_Smith50

c.smith3@bath.ac.uk

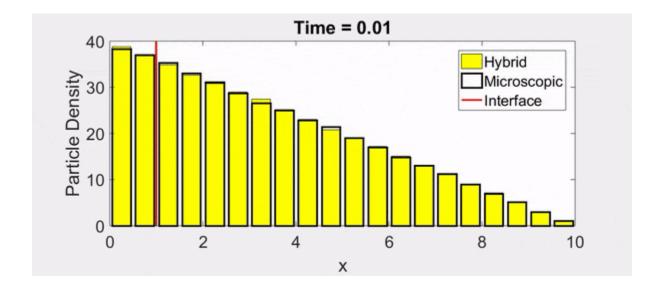
https://people.bath.ac.uk/cs640/

#### 1D-3D ARM



#### Results: 1D-3D ARM

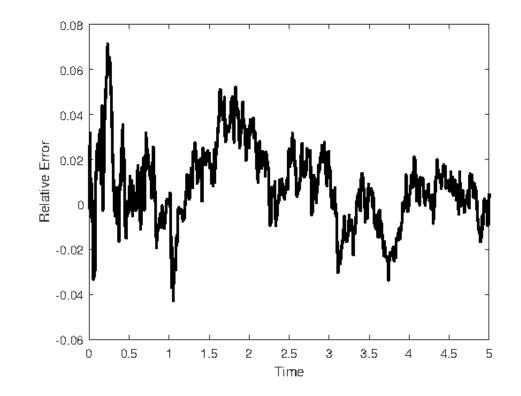
Reaction system: $2A \xrightarrow{\kappa_1} \emptyset \xrightarrow{\kappa_2} A$ Corresponding PDE: $\frac{\partial u}{\partial t} = D\nabla^2 u - \frac{\kappa_1}{L_y L_z} u^2 + \kappa_2 L_y L_z$ Moment closure (Poisson) $\langle A \rangle = \mathbb{V}\mathrm{ar}(A)$ 



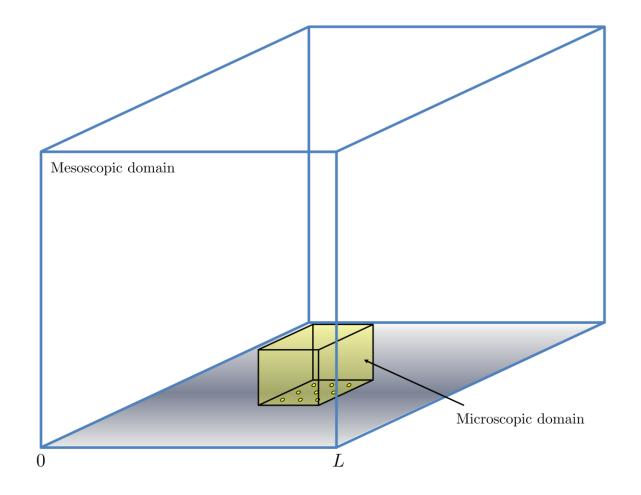
#### Error: 1D-3D ARM

$$E_{\rm Rel}(t) = rac{N_{\rm M}(t) - N_{\rm H}(t)}{N_{\rm M}(t)}.$$

 $N_{\rm H}(t)$  is the average number of particles in the final binning width of the hybrid method at time t, and  $N_{\rm M}(t)$  is the same for the fully microscopic simulation.

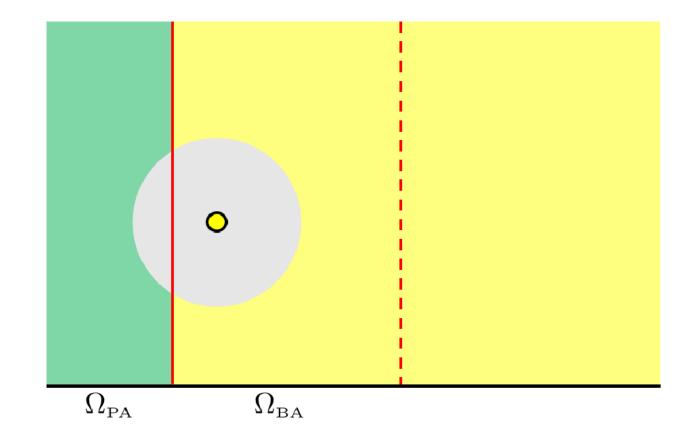


#### Calcium induced calcium release





#### **Reactions in the Brownian AR**



#### Summary of models

Scale	Advantages	Disadvantages
Macroscopic (Mean-field)	Fast to compute solutions. Suitable for high copy numbers. Amenable to analysis.	Inaccurate for low particle numbers. Mean-field dynamics diverge from individual-level behaviour for high-order reactions.
Mesoscopic (Compartments)	Fast for low copy numbers. Represents the individual-level behaviour.	Can be slow for large copy numbers. Does not retain precise locations of particles or particle identity.
Microscopic (Brownian-based)	Most accurate representation of the three.	Slow to compute reactions. Impractical for large numbers of particles.