

# Important CPM parameters

temperature

J values

$\lambda$

targetvolume: too small vs too large

Guideline: multiplying  $J$ ,  $\lambda$  and  $T$  by the same constant keeps the dynamics the same  $T$  and  $J$  in order 10,  $\lambda$  in order one typically yields decent dynamics.

# Objections and limitations

volume fluctuations are required for dynamics

no explicit membrane

When  $J$  is lower: more adhesion, more fluctuations

no explicit time scale

computational costs

Do due diligence

## “minimal” energy configurations

Depending on initial conditions and parameters, reaching the minimal energy configuration may take very long.



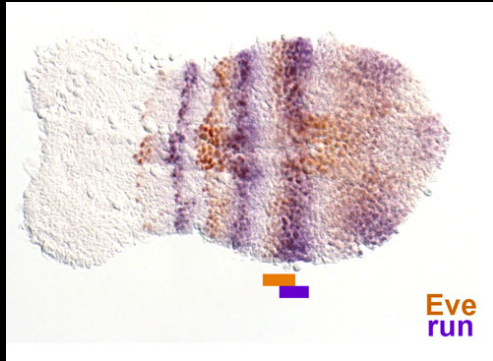
we can also use this in our favour

**Convergent extension**

or

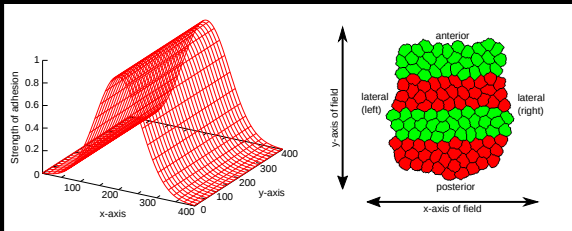
the importance of where you start from

# Differential adhesion and convergent extension

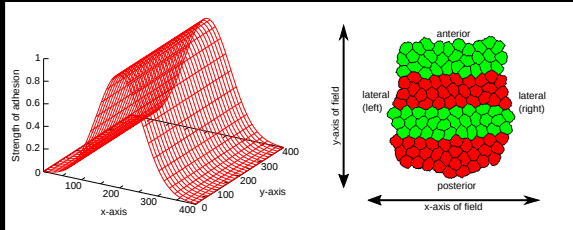


from Choe *et al.* 2006

# Convergent extension could potentially mess up segments



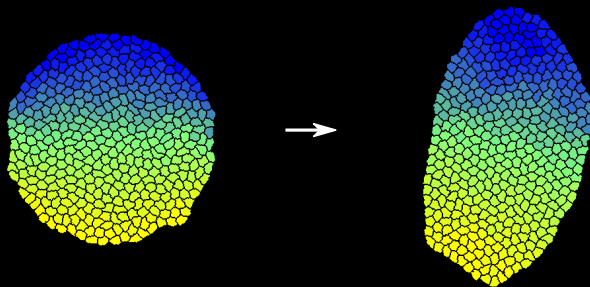
# segment-specific adhesion solves this



# Convergent extension **by** differential adhesion



# Convergent extension by graded expression of adhesive proteins



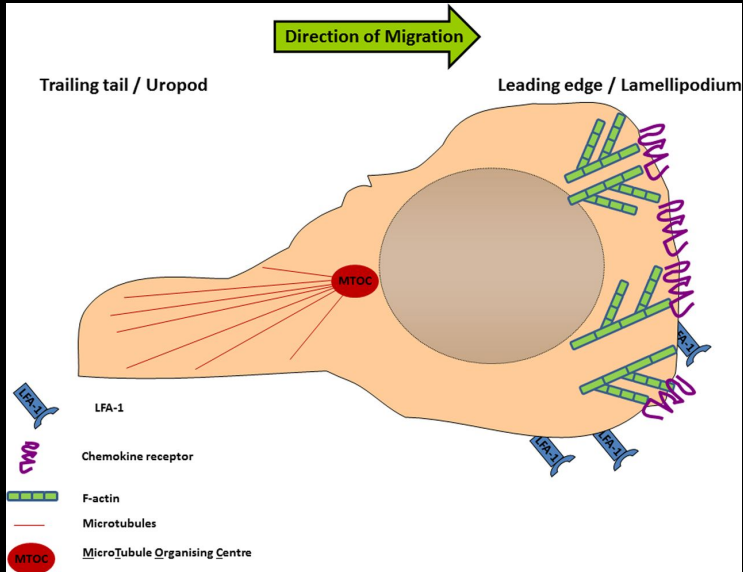
## **Cell migration**

# Persistent random walk in T cells

From Beltman *et al*, 2007

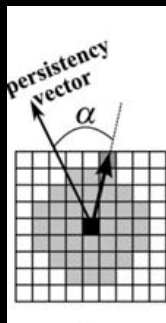
“stop-and-go motion”

# Underlying mechanism in brief



# Persistent random walk in CPM

$$\Delta H_{+} = -\mu \cos(\alpha)$$



The angle is updated every  $x$  timesteps, to reflect the actual direction of motion of the cell

We define this mechanism as change in the Hamiltonian:  
directly add an extra bias to copy probability

## Exercises part 2

