

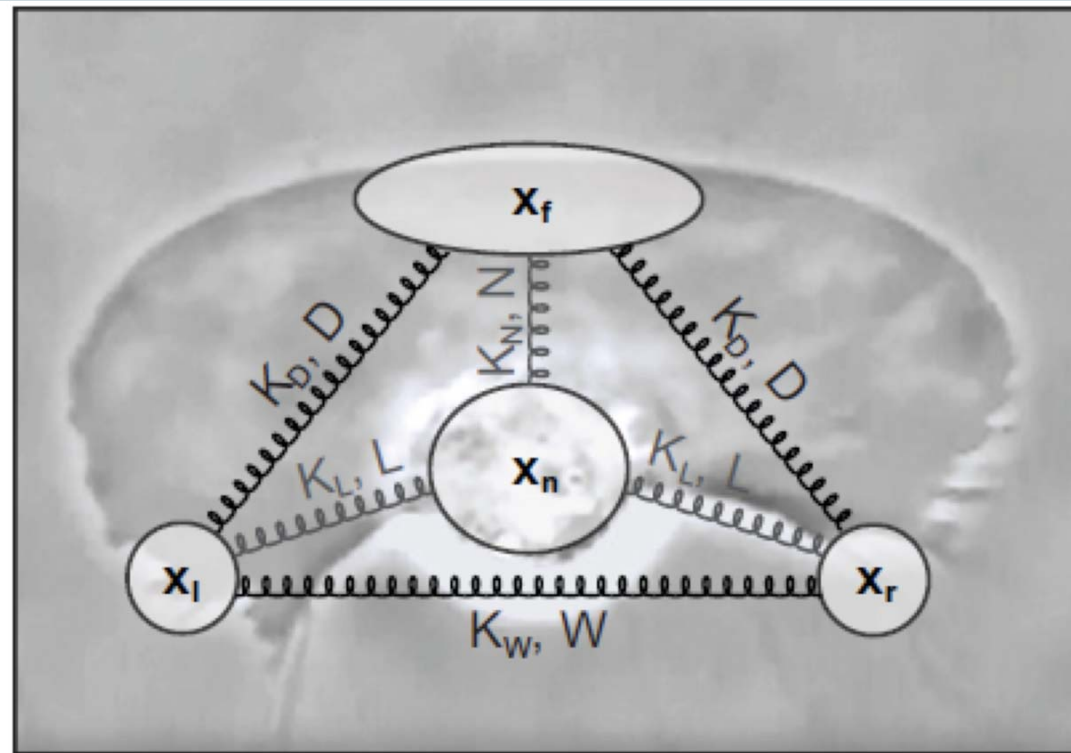
A Multiple Spring Model that Predicts Bipedal Locomotion of Crawling Cells



Alex Loosley and Jay X. Tang, Physics Department
Brown University, Providence RI

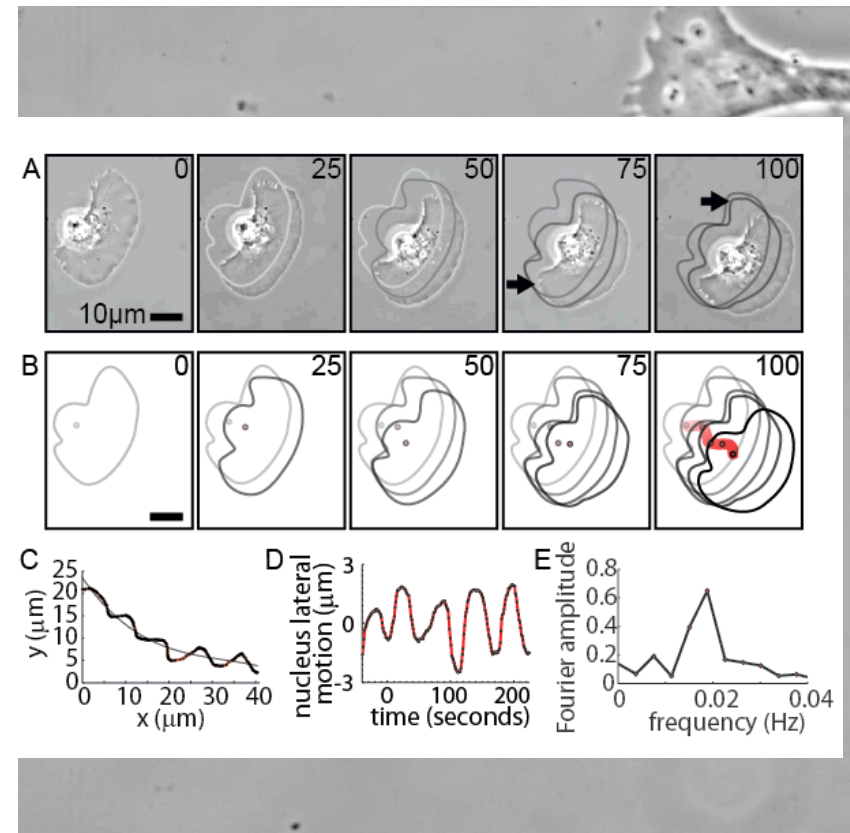


BROWN



Fish Keratocyte Dynamics

- Front edge extends forward at constant speed (here $0.17 \mu\text{m/s}$)
- Cell body contracts in alternate steps from both left and right sides of trailing edge
- Nucleus swings from side to side as the cell crawls forward



E. Barnhart , G. Allen, F. Julicher, J. Theriot, *Biophys J*, **98**, 933-942 (2010)

Crawling and the 1D Model

- 1) Actin polymerization occurs at the leading edge of cell to protrude the cell forward
- 2) Contractile forces generated within cytoskeleton pull the rear forward

Leading Edge: $x_1, \dot{x}_1 = v_f$

Trailing Edge: x_2

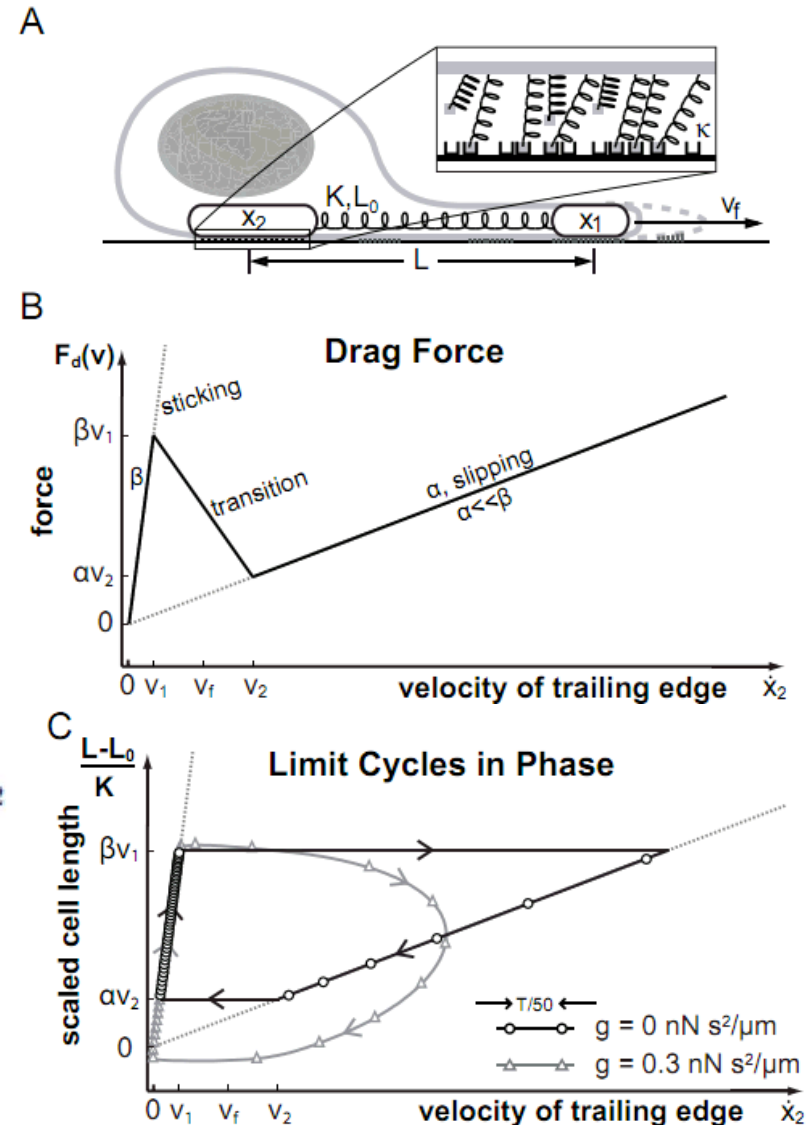
Spring represents elastic coupling between elements x_1 and x_2

Expression of the Drag Force

$$F_d[\dot{x}_2] = \begin{cases} -\beta\dot{x}_2, & \dot{x}_2 < v_1 \\ \frac{v_1 - \dot{x}_2}{v_1 - v_2}(\beta v_1 - \alpha v_2) - \beta v_1, & v_1 < \dot{x}_2 < v_2 \\ -\alpha\dot{x}_2, & v_2 < \dot{x}_2 \end{cases}$$

Equations of Motion

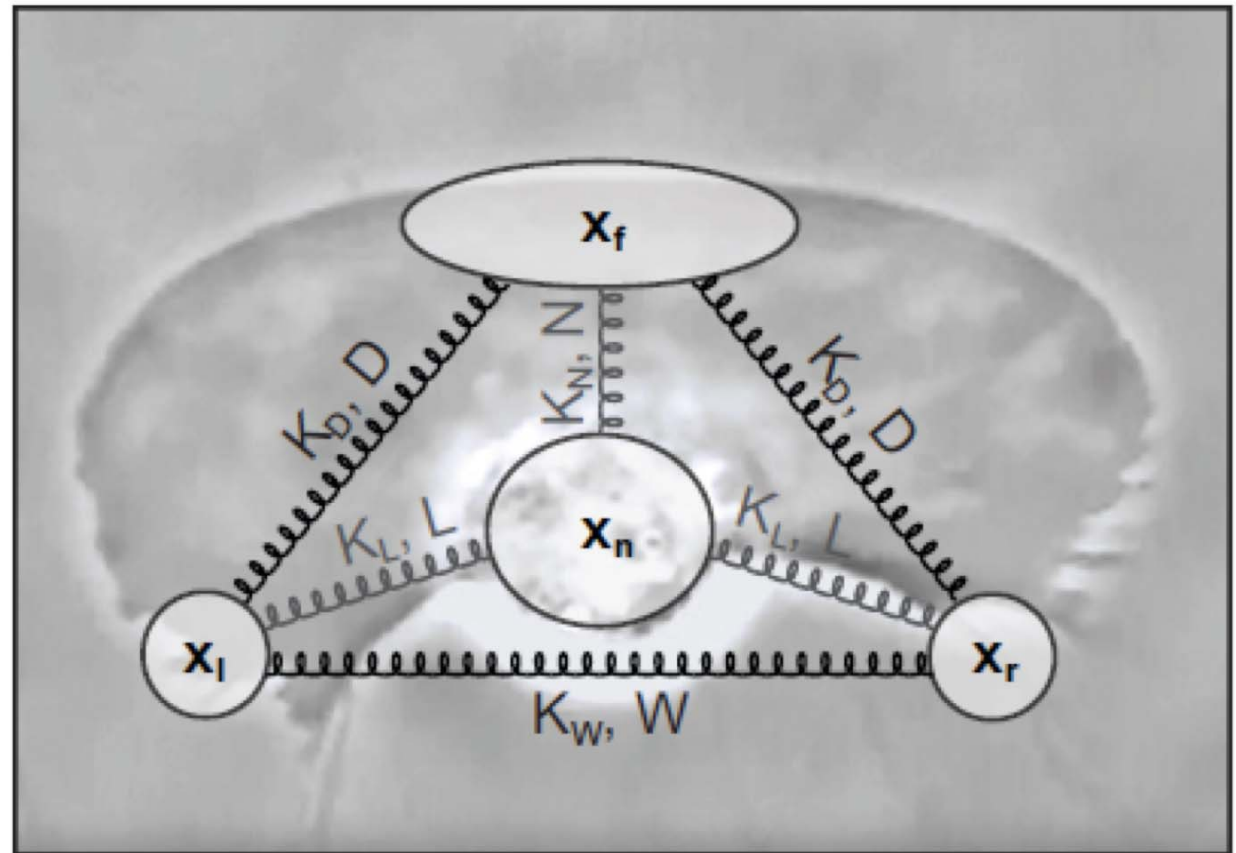
$$\begin{aligned} \dot{x}_1 &= v_f \\ g\ddot{x}_2 - F_d[\dot{x}_2] - K(L - L_0) &= 0 \end{aligned}$$



The 2D Model

General criteria

- Minimal number of components are introduced in the model
- Cell front moves with constant speed
- No asymmetry between left and right is required
-
- Left and right rears can move out of phase
- The nuclear position is shown explicitly



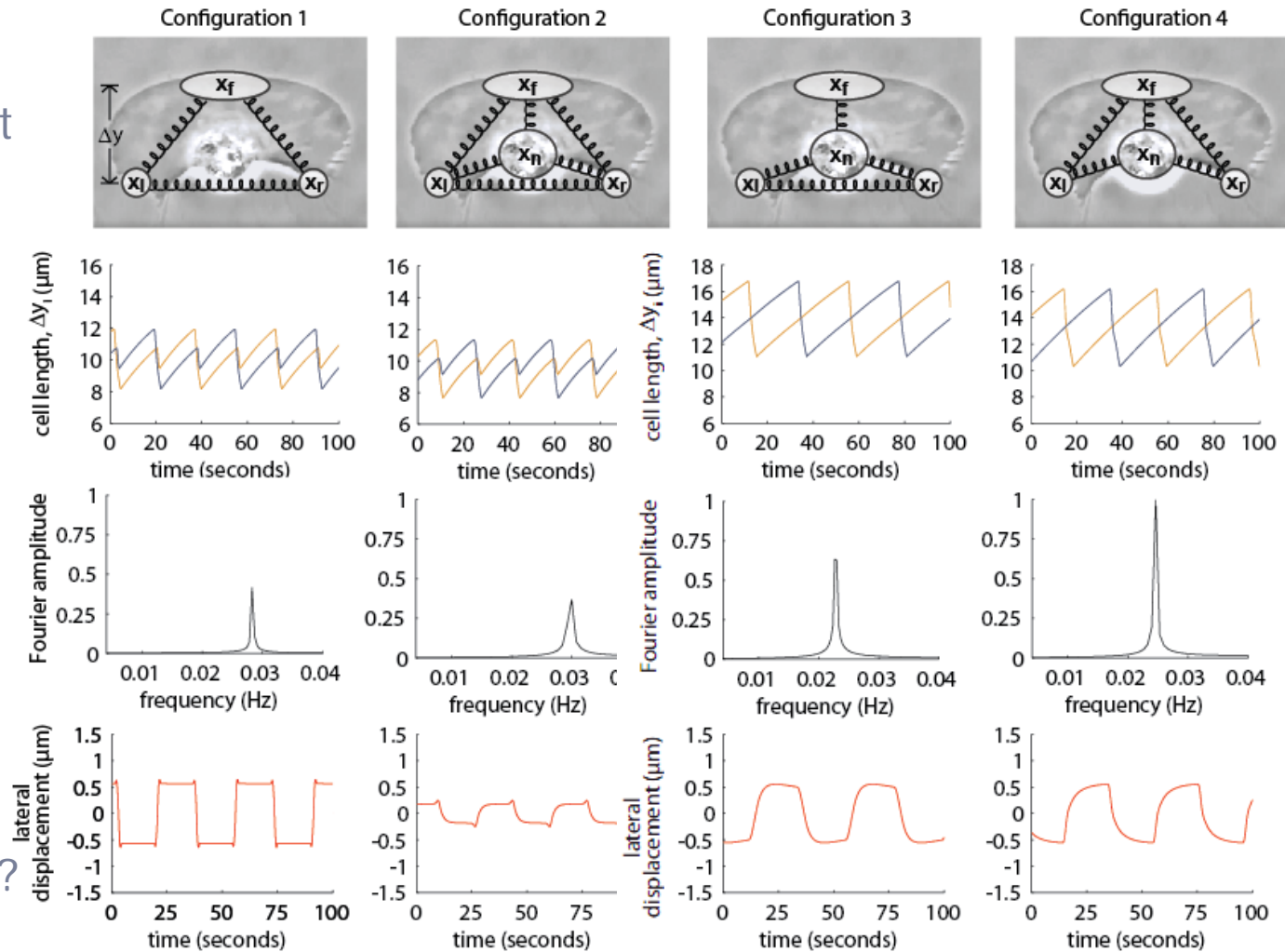
Configurations of the 2D Model

Configuration 1

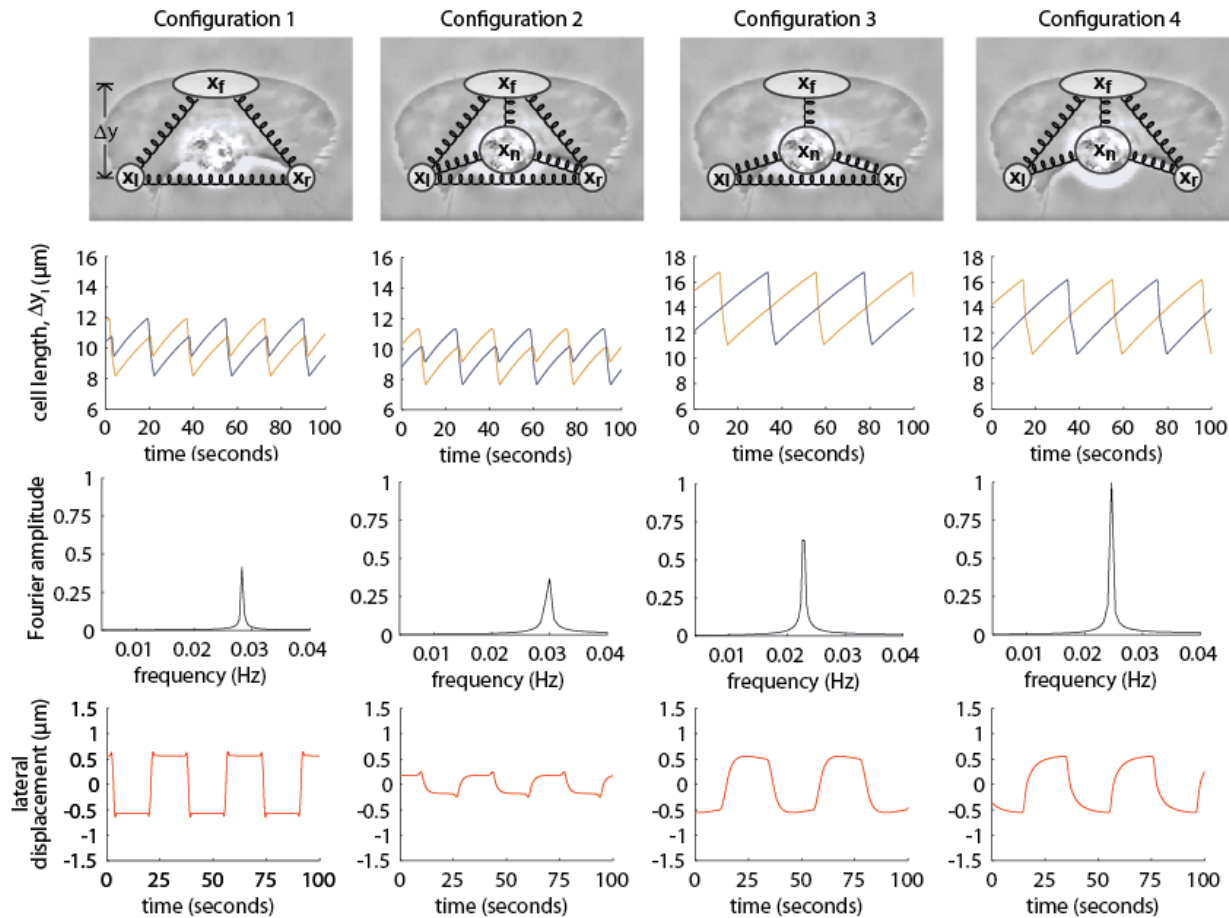
- Simplest configuration that can generate bipedal locomotion
- Problems with cell shape and dynamics

Configuration 2

- Element representing the nucleus added
- System over-constrained by too many springs?



Configurations of the 2D Model



Configuration 3

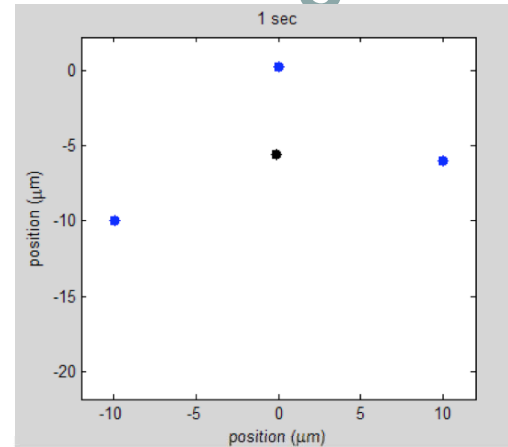
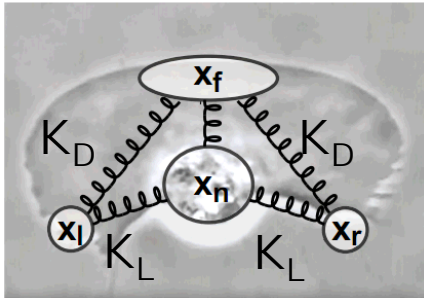
- Configuration proposed by Barnhart *et al.*
- The addition of spring K_{nf} gives realistic cell length

Configuration 4

- An alternative spring configuration based on assumptions of cell symmetry and element localization

A detailed analysis of Configuration 4

Configuration 4



Equations of Motion:

$$\dot{x}_f = v_f \hat{y}$$

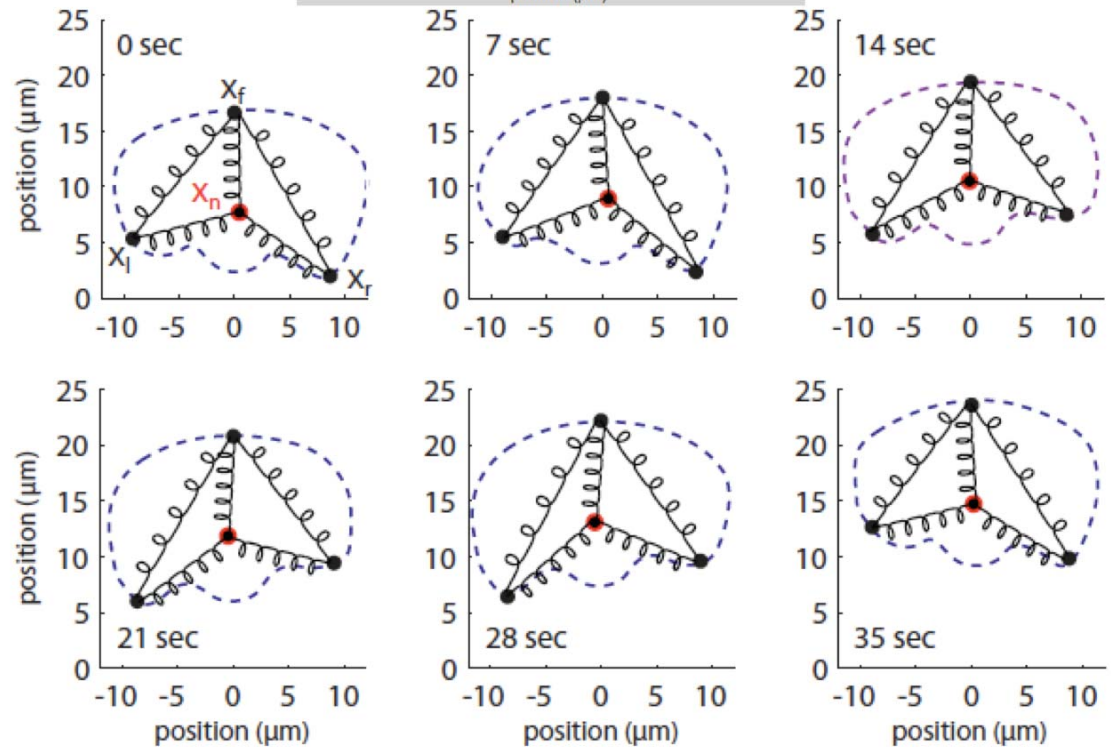
$$\gamma \dot{x}_n - \sum_{i=f,l,r} K_{in} (|x_n - x_i| - L_{in,0}) \frac{x_n - x_i}{|x_n - x_i|} = 0$$

$$g \ddot{x}_l - F_d(|\dot{x}_l|) \frac{\dot{x}_l}{|\dot{x}_l|}$$

$$- \sum_{i=f,n,r} K_{il} (|x_l - x_i| - L_{il,0}) \frac{x_l - x_i}{|x_l - x_i|} = 0$$

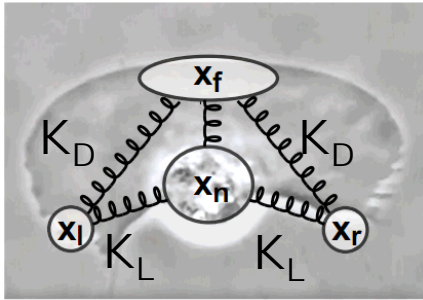
$$g \ddot{x}_r - F_d(|\dot{x}_r|) \frac{\dot{x}_r}{|\dot{x}_r|}$$

$$- \sum_{i=f,n,l} K_{ir} (|x_r - x_i| - L_{ir,0}) \frac{x_r - x_i}{|x_r - x_i|} = 0$$

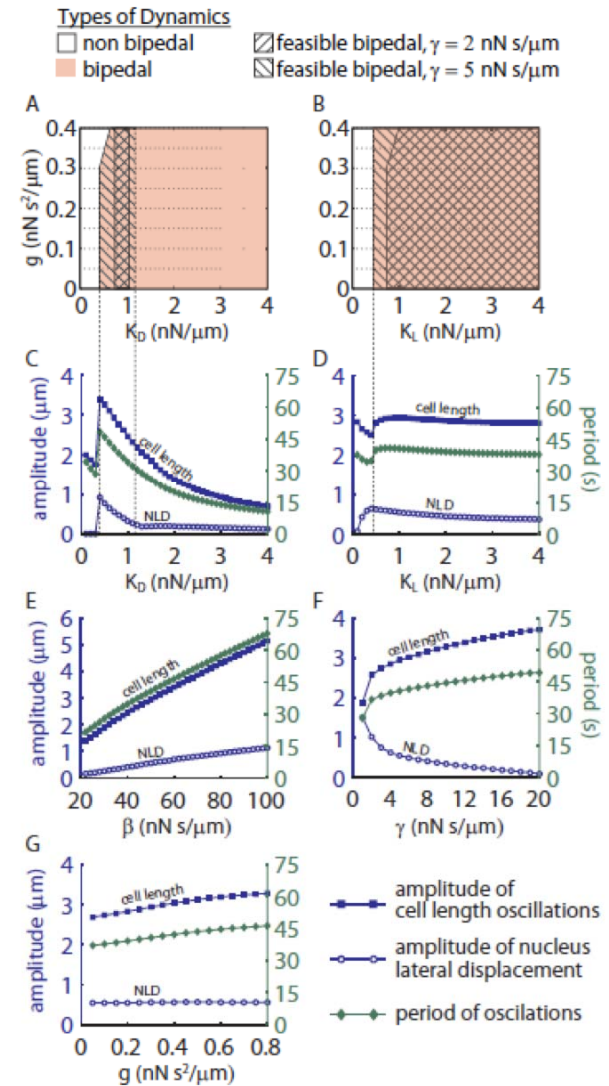


Model predictions of Configuration 4

Configuration 4



parameter	meaning	range investigated	units
α	slipping drag coefficient (viscous shear)	0.15 - 0.5	$nN s/\mu m$
β	sticking drag coefficient (adhesion at trailing edge)	20 - 100	$nN s/\mu m$
γ	nuclear drag coefficient (adhesion at cell nucleus)	1 - 20	$nN s/\mu m$
g	inertial term (sets the time-scale of switching between stick and slip dynamics)	0 - 0.8	$nN s^2/\mu m$
v_1	critical sticking velocity (upper limit of the sticking domain)	0.08	$\mu m/s$
v_2	critical slipping velocity (lower limit of the slipping domain)	1	$\mu m/s$
v_f	leading edge velocity ($v_f > v_1$ required for stick-slip dynamics)	0.2	$\mu m/s$
K_N	spring constant	0.1 - 10	$nN/\mu m$
K_D	spring constant	0.1 - 10	$nN/\mu m$
K_L	spring constant	0.1 - 10	$nN/\mu m$
K_W	spring constant	0.1 - 10	$nN/\mu m$
N	spring length	conforms to cell shape	μm
D	spring length	conforms to cell shape	μm
L	spring length	conforms to cell shape	μm
W	spring length	conforms to cell shape	μm



Advantage of Configuration 4

It recapitulates the correct:

- Timescale
(period ~ 40 sec)
- Amplitudes
(bipedal ~2.5 μm)
(lateral disp. ~ 0.5 μm)
- Smooth lateral motion of nucleus

This configuration robustly generates bipedal locomotion and appropriate lateral displacement of the nucleus over realistic variation of parameters.

Summary

- Two 4 element configurations with centralized mechanical coupling produce the realistic length and timescales of bipedal locomotion
- Both configurations robustly generate bipedal locomotion over a range of realistic parameter variations
- The new configuration we propose explicitly couples the nucleus with the cytoskeletal elements, thus the model reproduces the nuclear lateral motion observed experimentally

Acknowledgements



**Jingjing
Wang**



**Angus J
McMullen**

This work is supported by NSF CMMI 0825873, and by an PGS D fellowship from the Natural Sciences and Engineering Research Council of Canada.

