

Outline

1. Motivations: Large-scale models vs. small-scale heterogeneity
2. Observations: Statistics of movement at the individual level
3. Derivations: Spatially-explicit models of populations
4. Approximations: Large-scale effects of unresolved heterogeneity

Quantifying organism movement: gps data logger

Charlotte Boyd

Peruvian Boobies (*Sula variegata*)



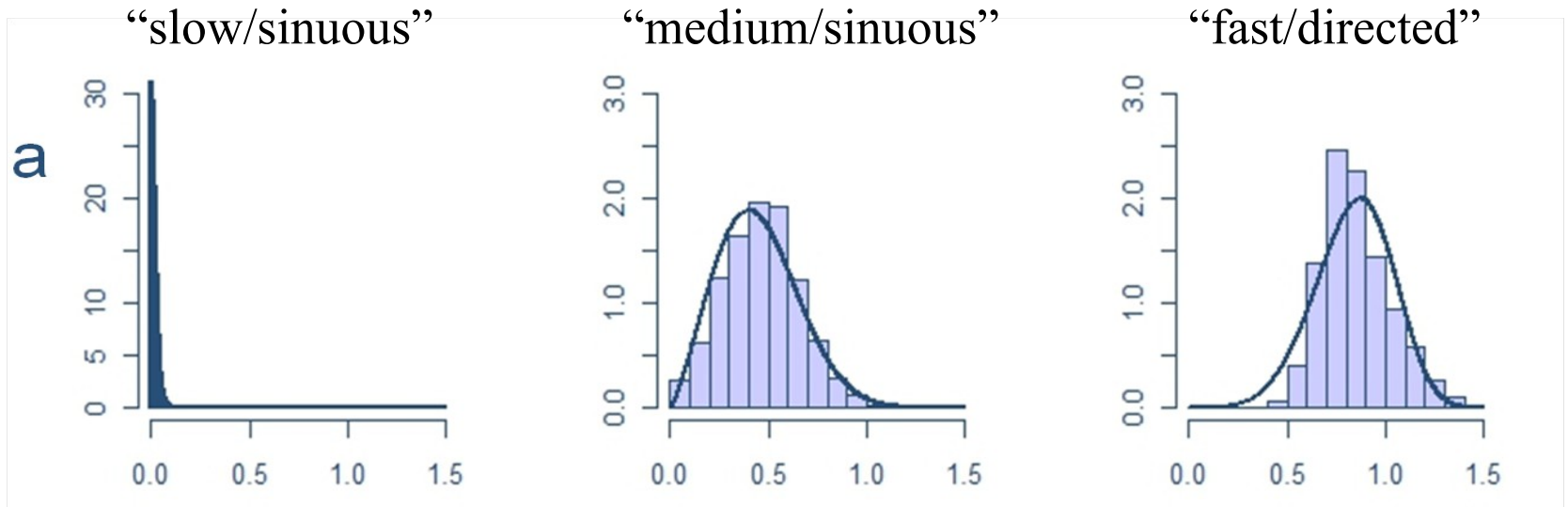
Photo: S. Bertrand, IRD

Quantifying organism movement: gps data logger

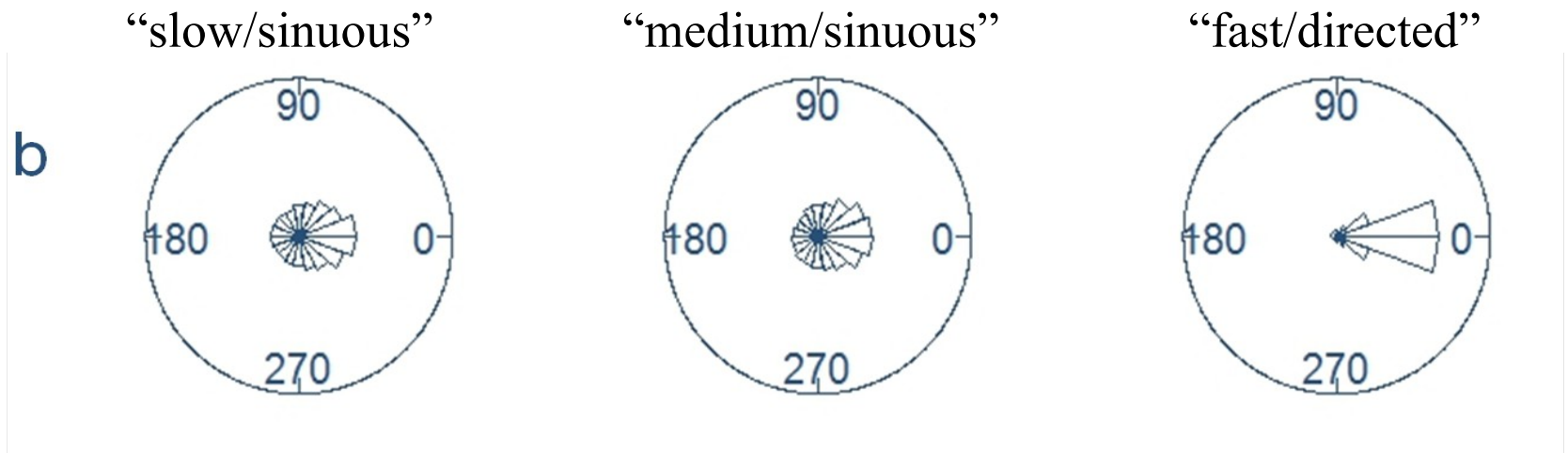
Charlotte Boyd

Hidden Markov Model / Maximum Likelihood Estimation

Observed distances (km over 1 minute intervals) assigned by maximum likelihood

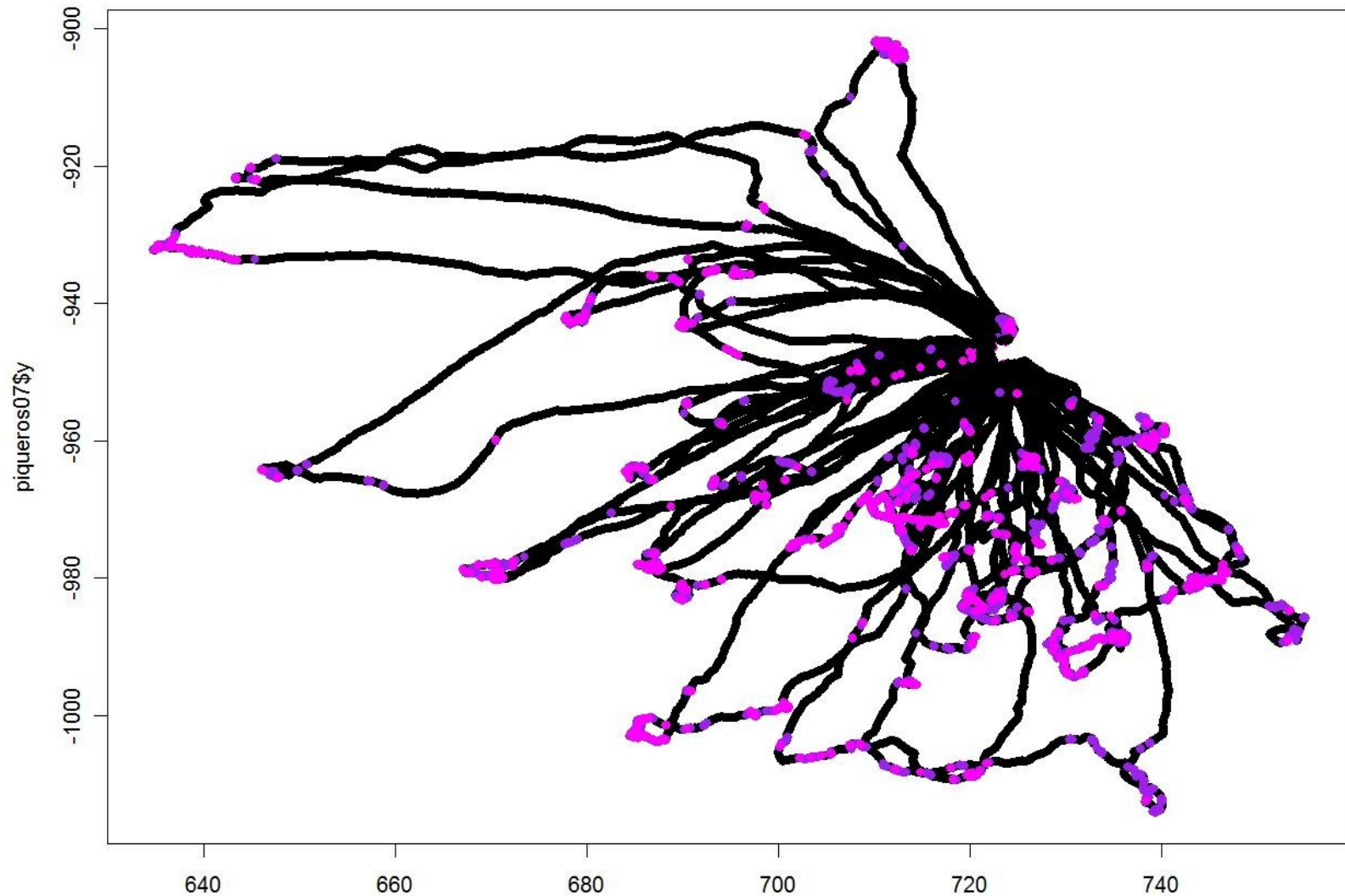


Observed turning angles assigned by maximum likelihood



Quantifying organism movement: gps data logger

Charlotte Boyd



Central-place foraging near breeding colonies: slow/sinuous modes indicative of fine-scale search or foraging in magenta

Quantifying organism movement: satellite tracking

Elizabeth Skewgar



Behavior of Magellanic Penguins at Sea

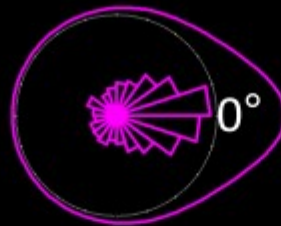
Quantifying organism movement: satellite tracking

Elizabeth Skewgar

Hidden Markov Model / Maximum Likelihood Estimation constrained by prior data

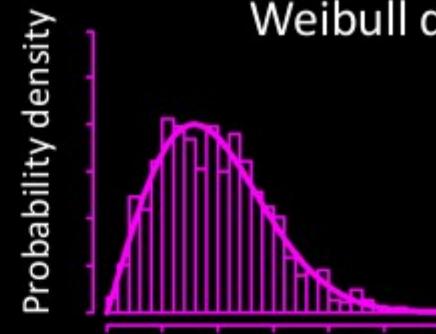
Outbound

Wrapped Cauchy distribution



$$\rho = 0.62$$

Weibull distribution

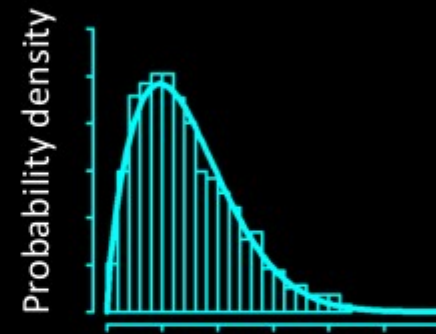


scale = 109
shape = 2.0

Foraging

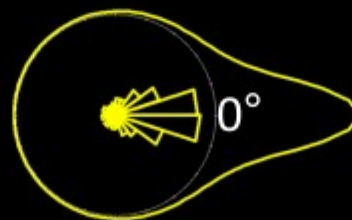


$$\rho = 0.35$$

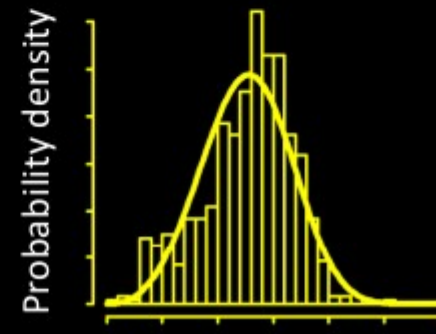


scale = 81
shape = 1.7

Inbound



$$\rho = 0.80$$



scale = 141
shape = 3.6

Turning angles

0 100 200 300

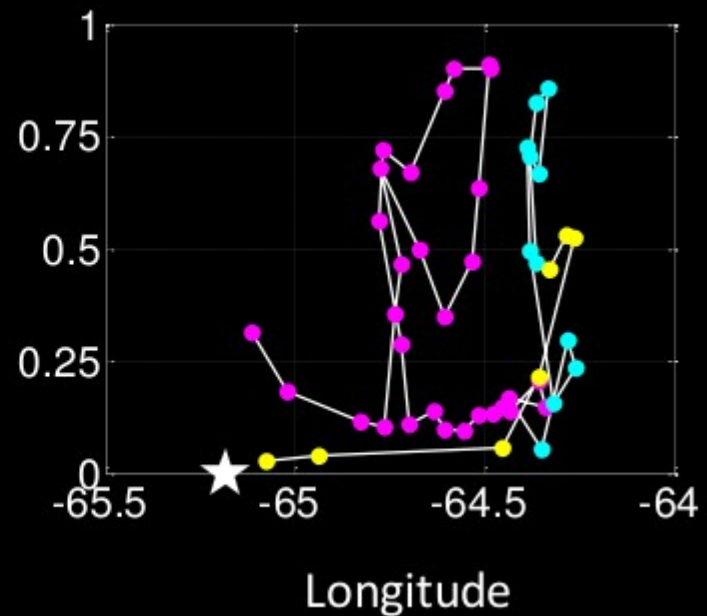
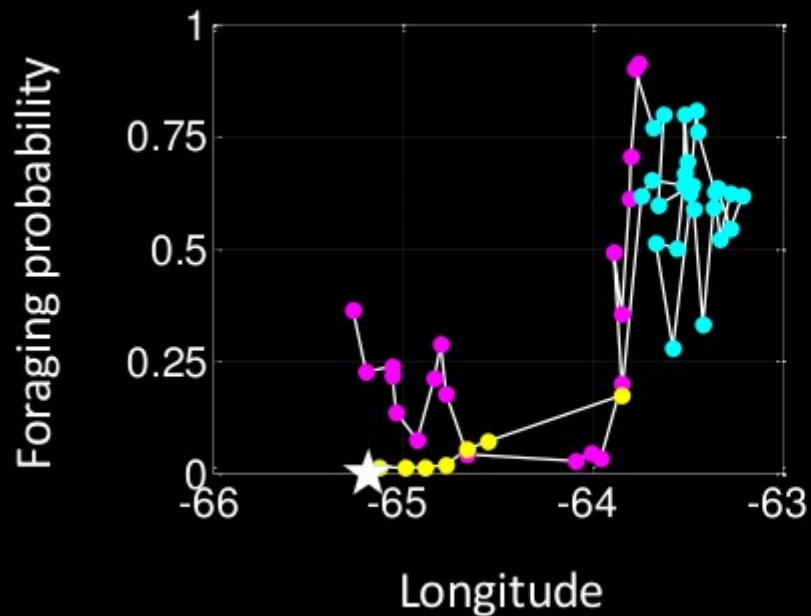
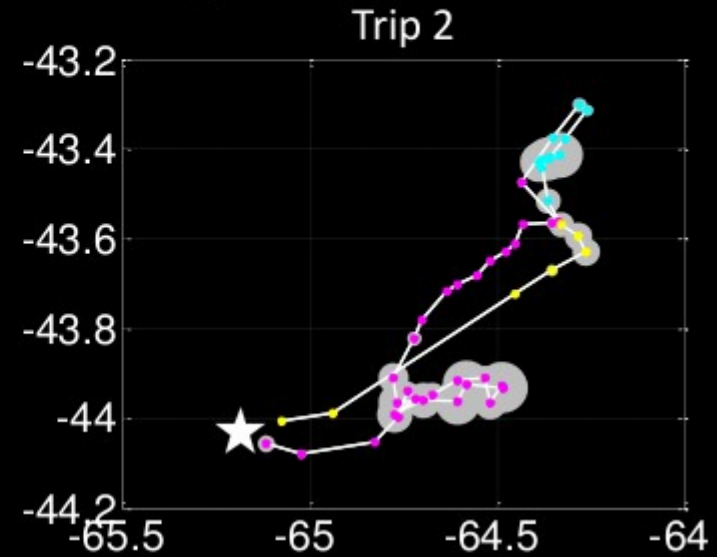
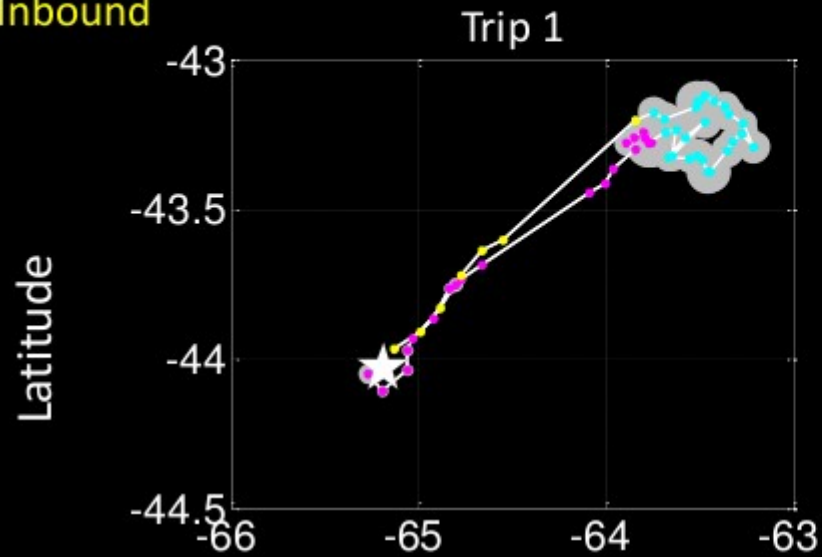
Speed (km/day)

Quantifying organism movement: satellite tracking

Elizabeth Skewgar

Chick rearing

Outbound
Foraging
Inbound

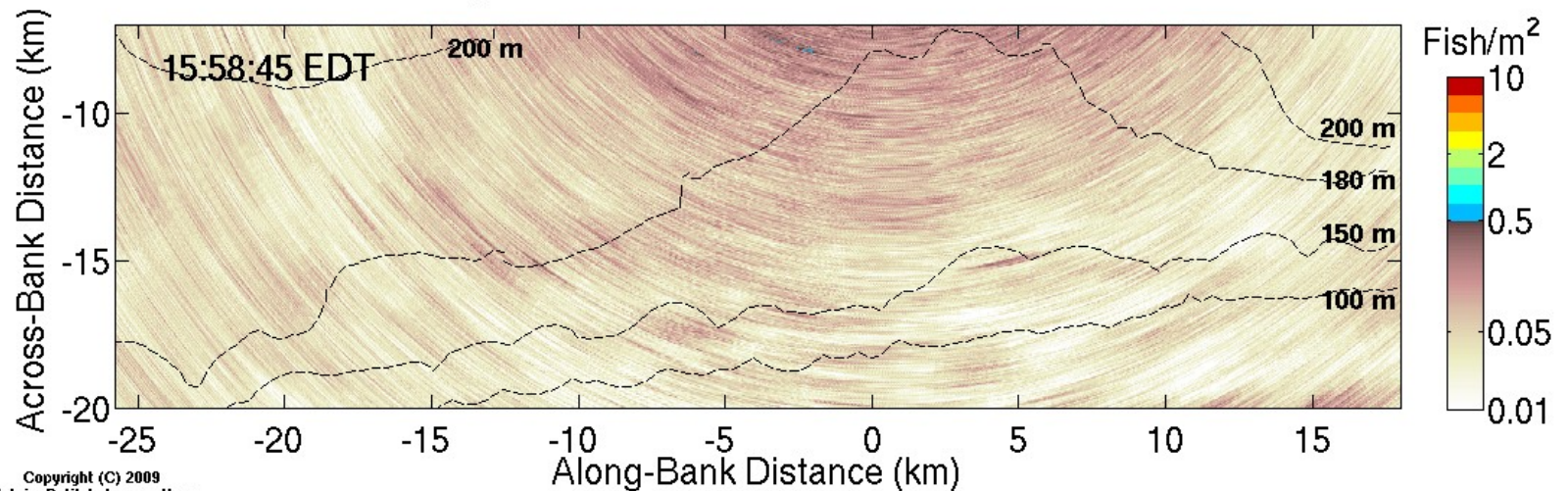


Herring school dynamics from acoustic tracking



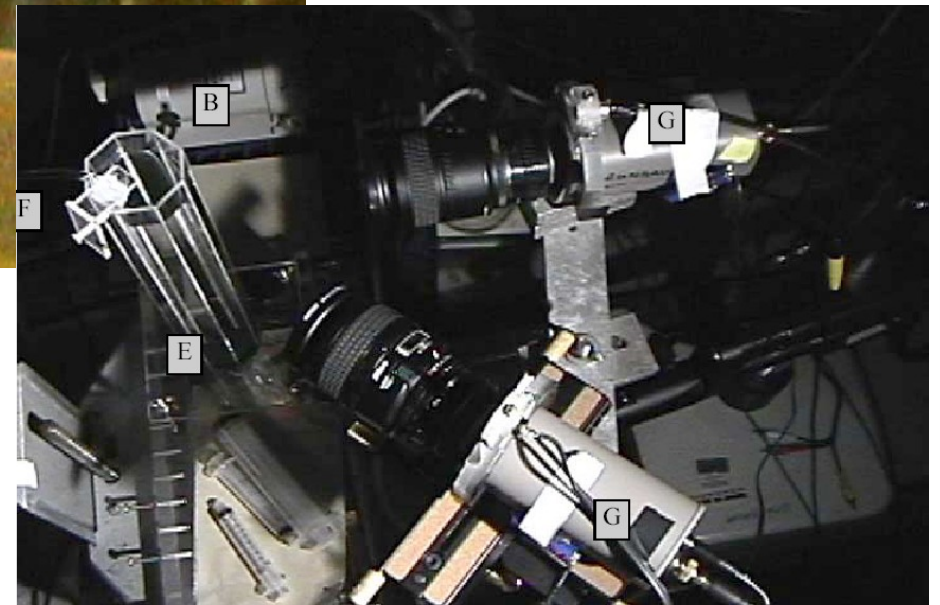
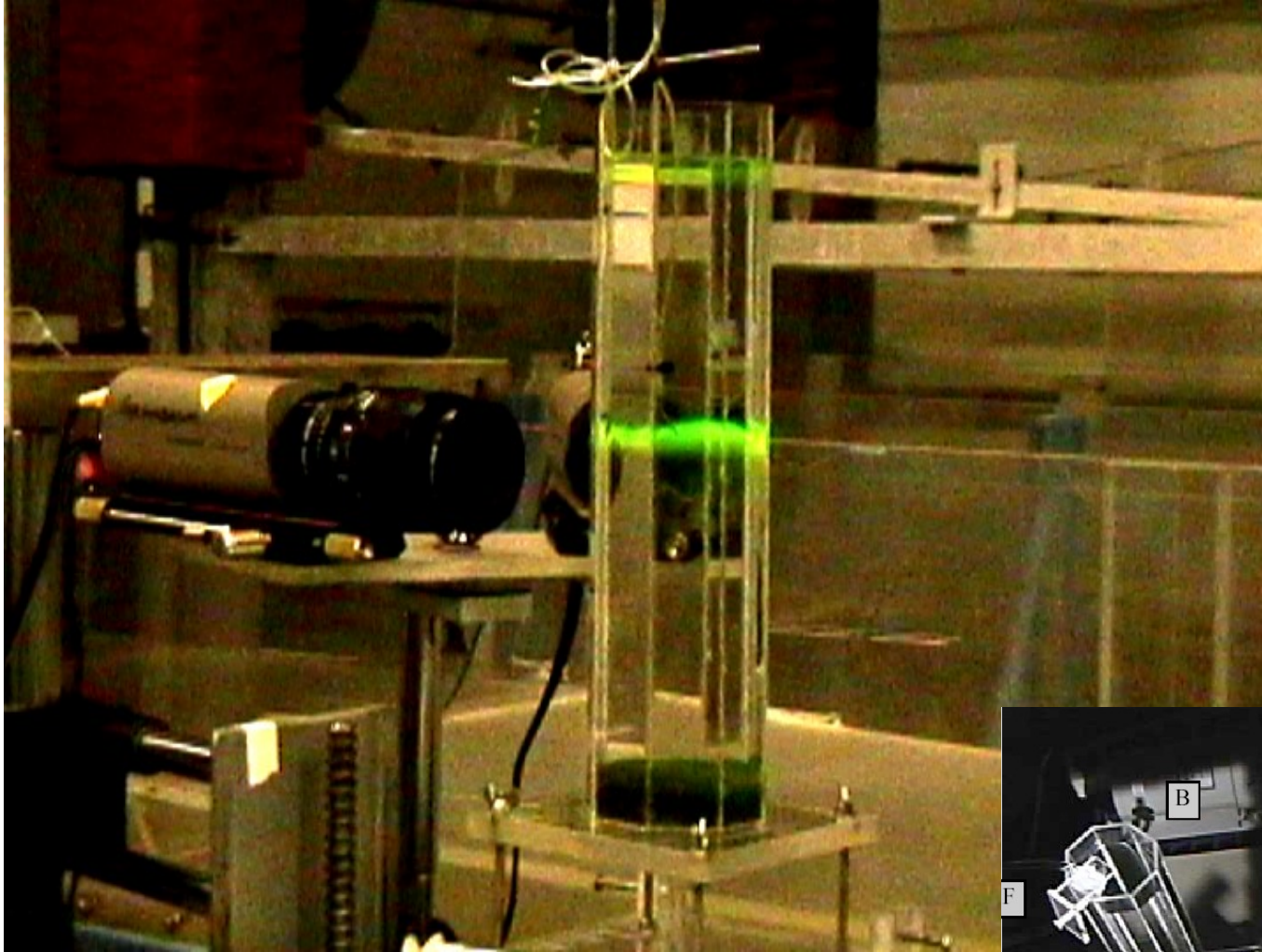
Handegard
et al. 2005

Formation and Migration of Herring Shoals on October 3, 2006

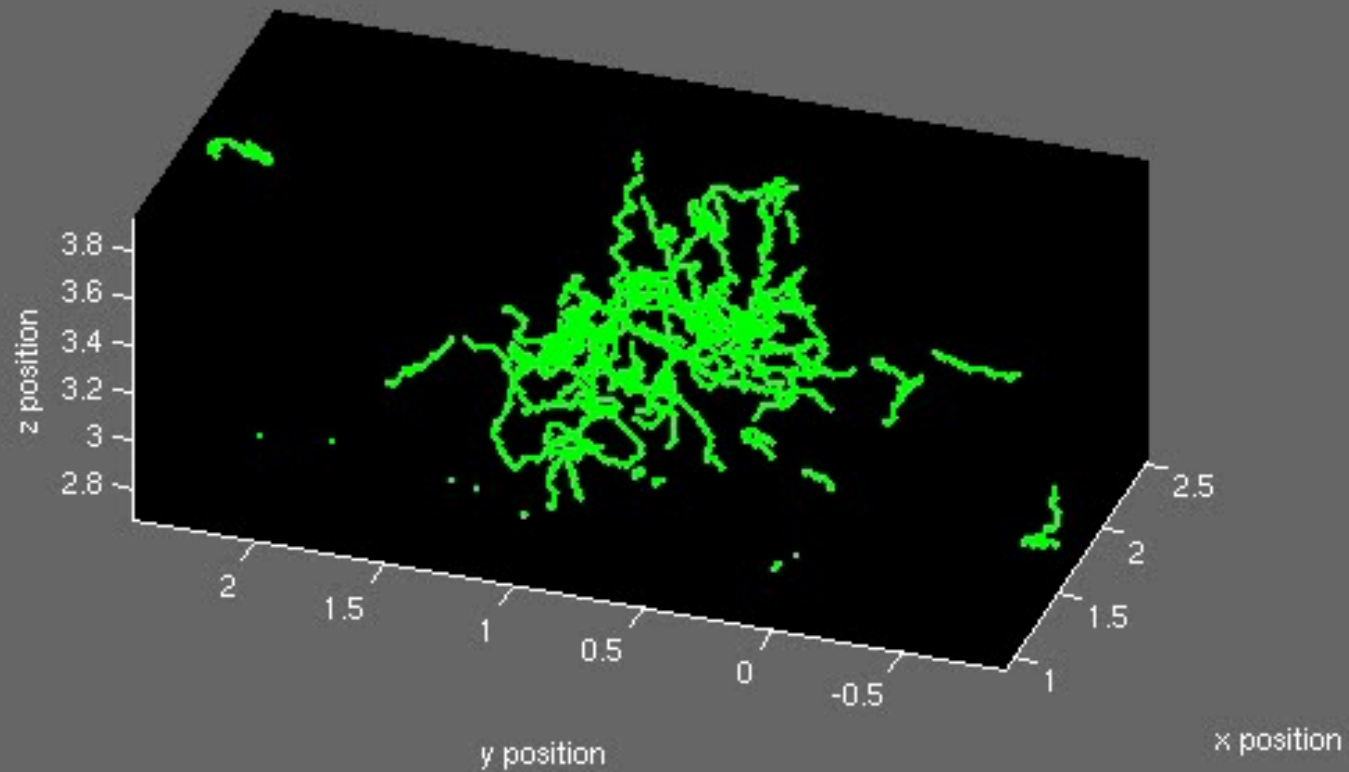


Makris et
al. 2009

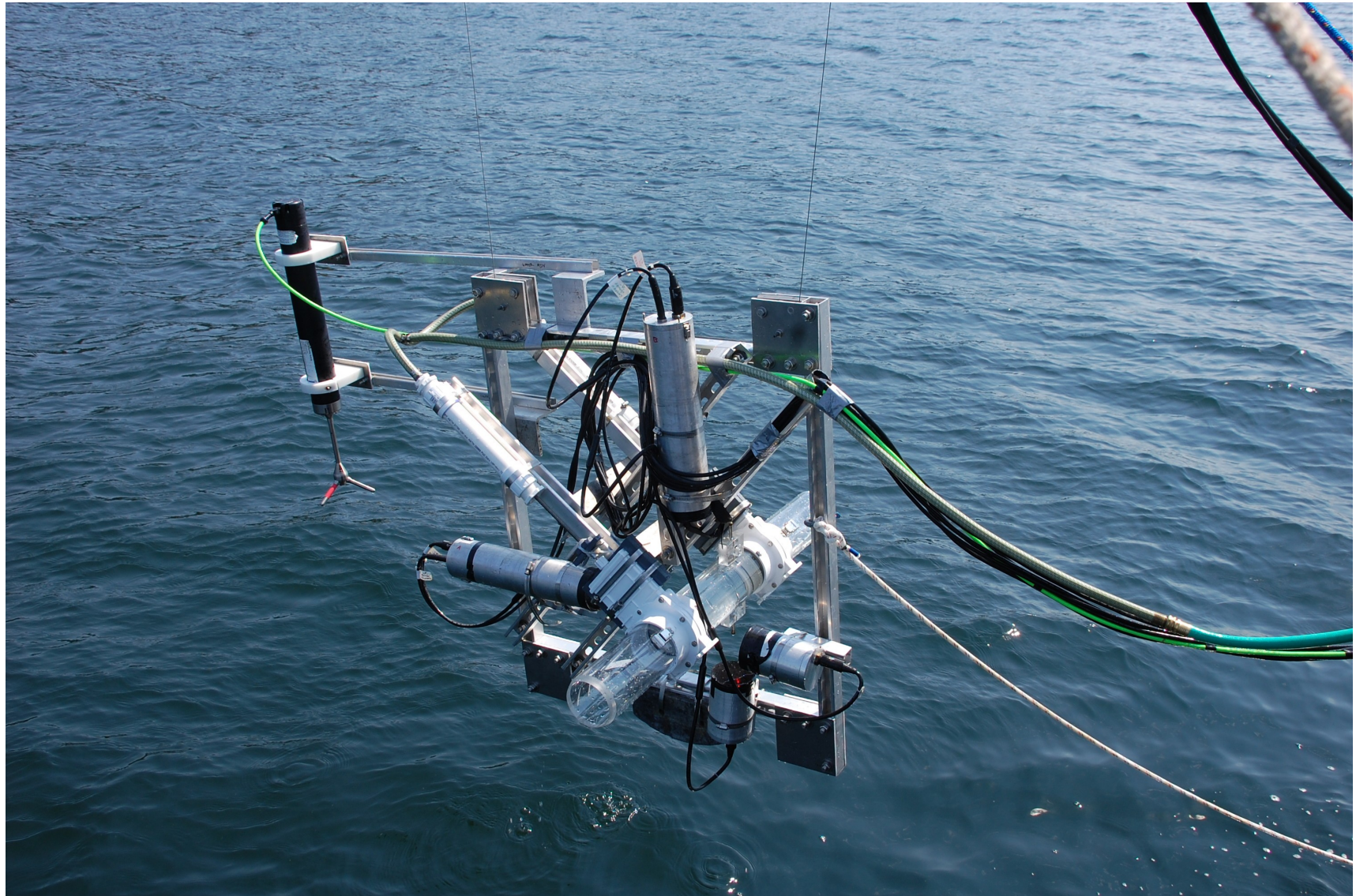
Quantifying organism movement: video tracking



Video tracking for quantifying distributions and movements of plankton
Swimming trajectories, *Strombidinopsis* sp., Susanne Menden-Deuer



Low cost, field-deployable *in situ* sampler



3D Footage of Natural Communities



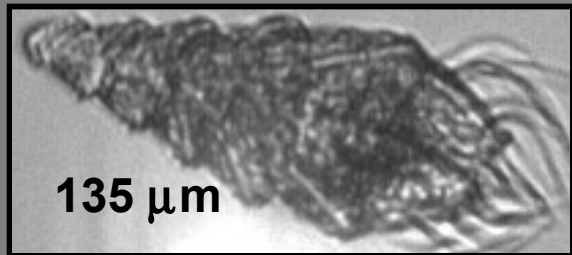
20 mm

30 mm

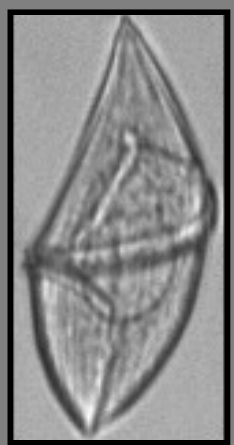


265 μm

Laboea strobila



135 μm



45 μm

Heterosigma akashiwo Harmful Algal Blooms



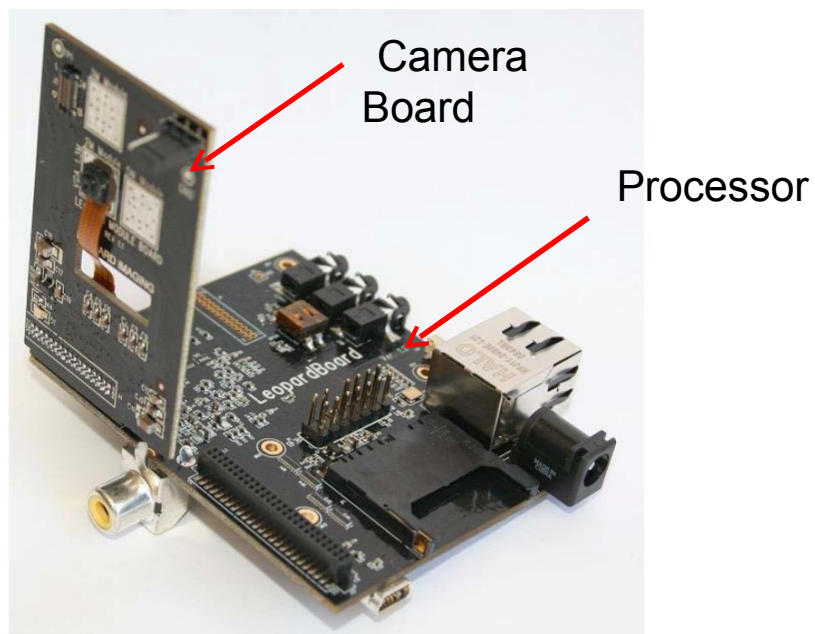
Heterosigma cell. Photo: M. Black



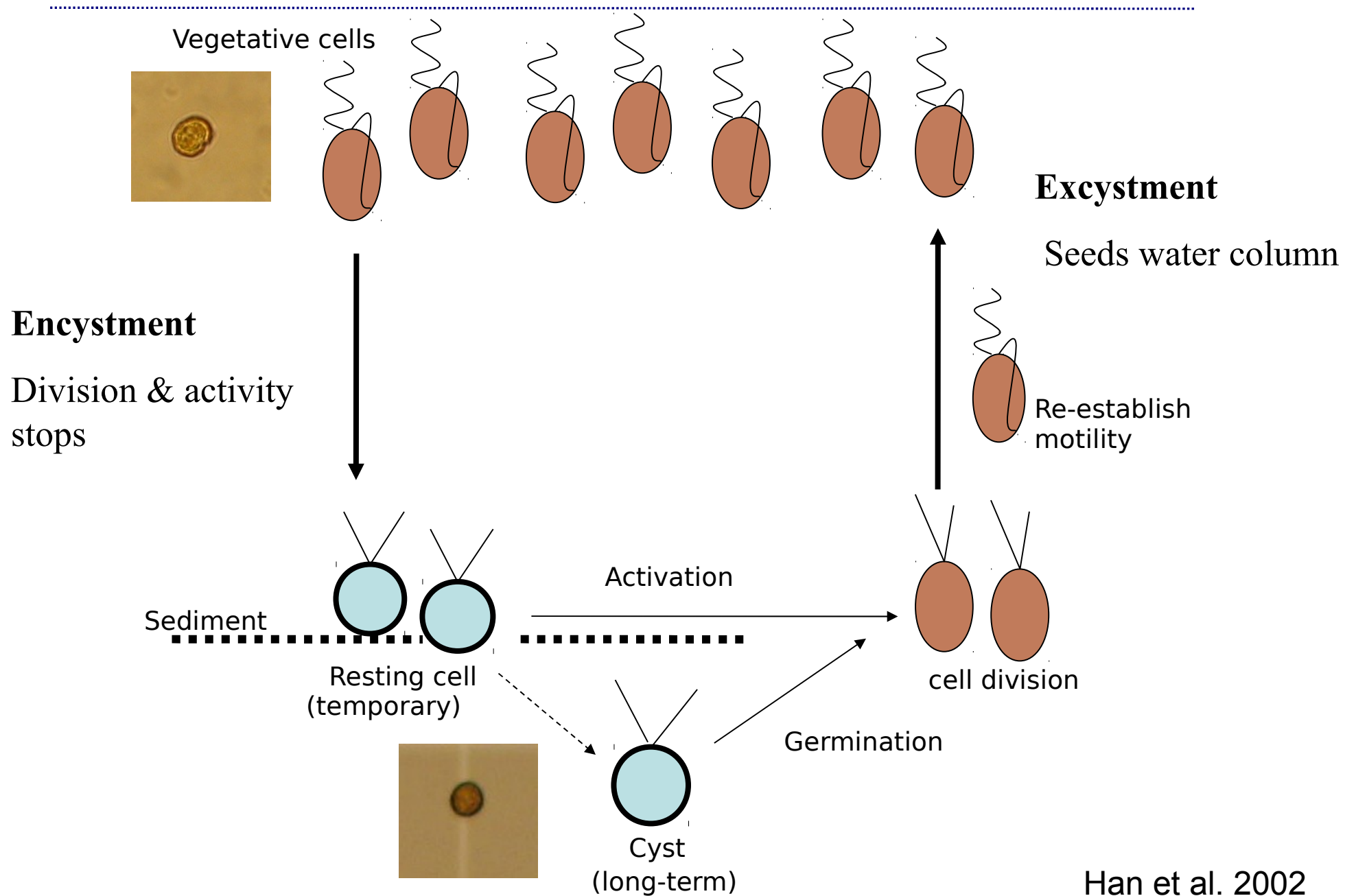
Heterosigma Harmful Algal Bloom. Photo: K. Fredrickson

Lower cost, field-deployable *in situ* sampler

“LeopardBoard”:
ARM computer-based
high-definition imager

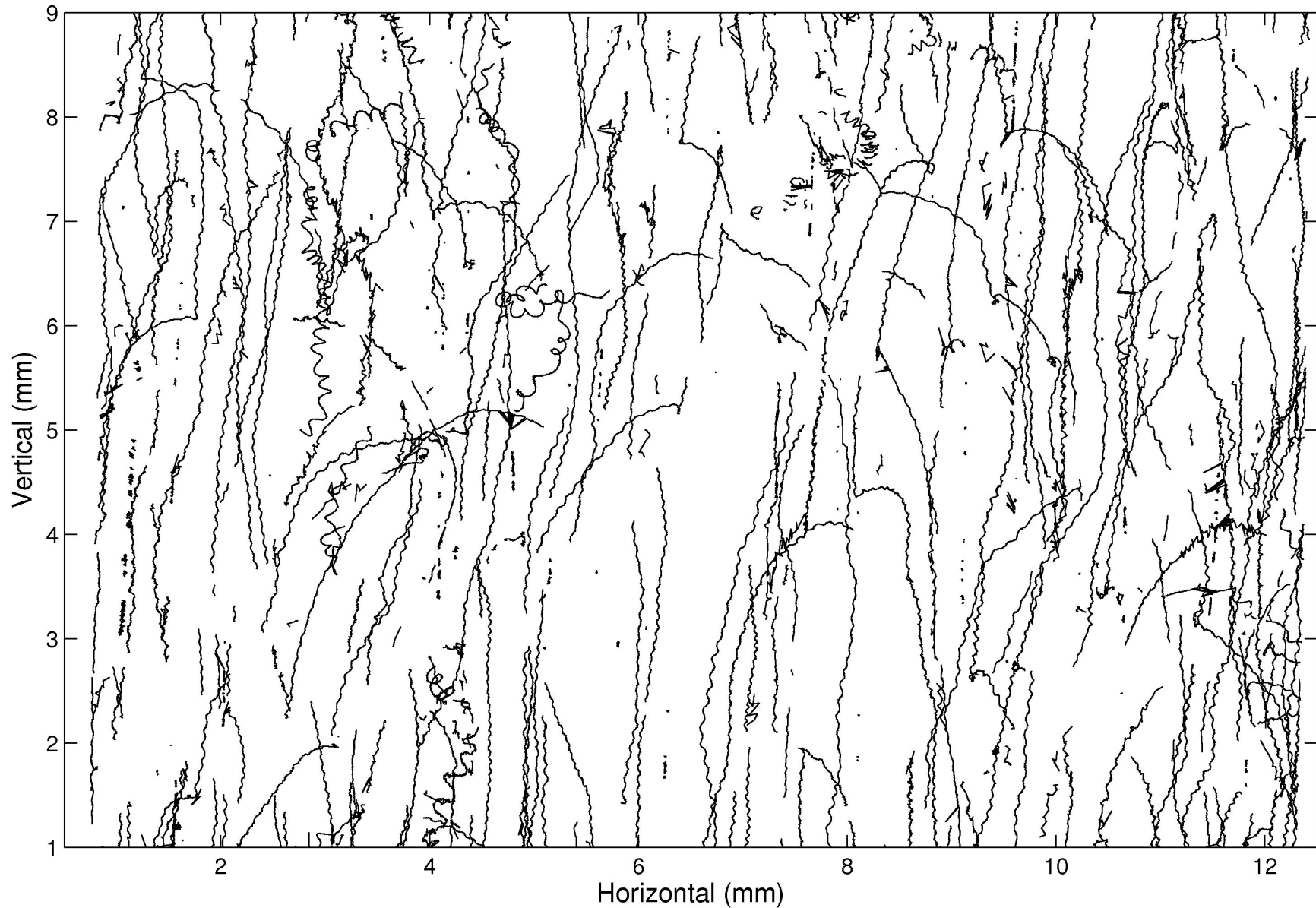


Generalized Lifecycle of *Heterosigma*



Vegetative *Heterosigma* cell swimming

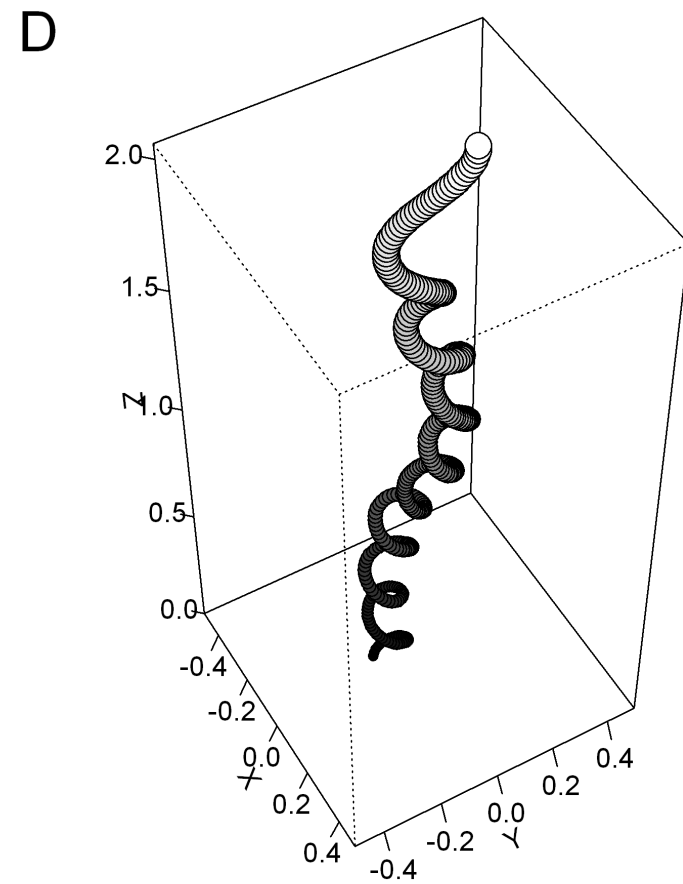
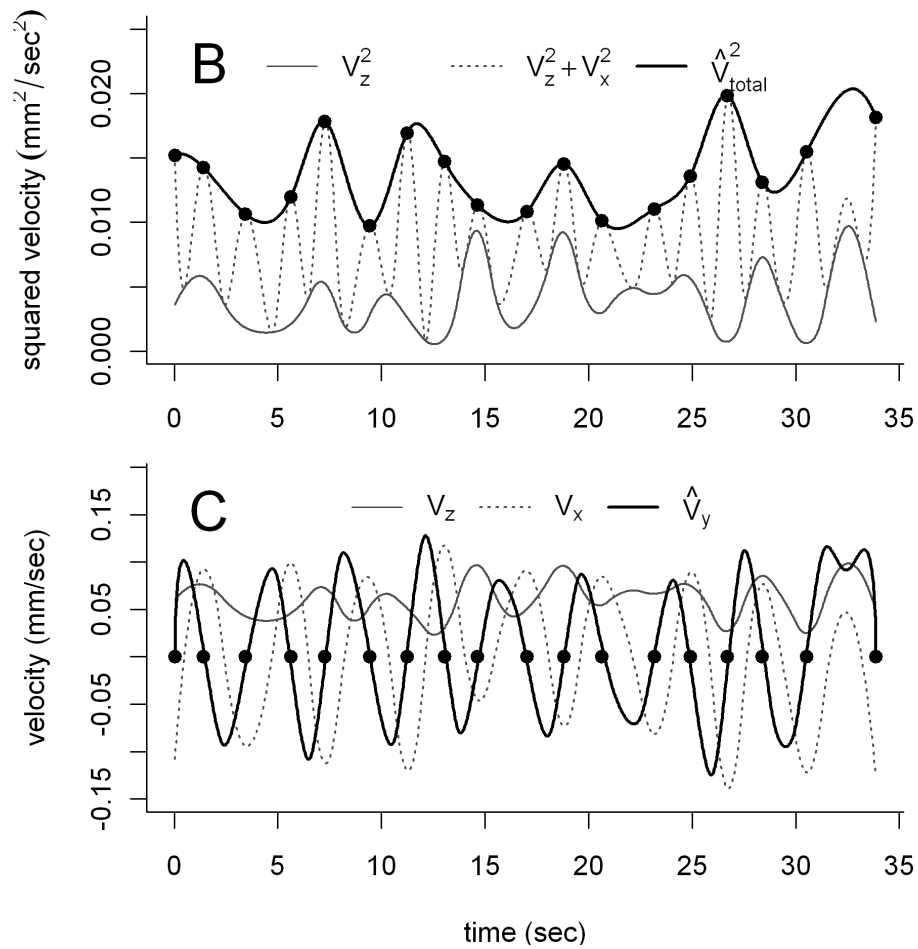
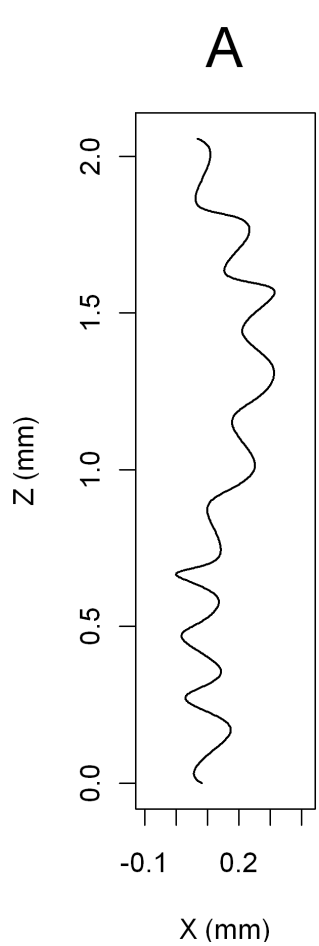
Strain = CCMP452, NY, USA



M. T. Nishizaki

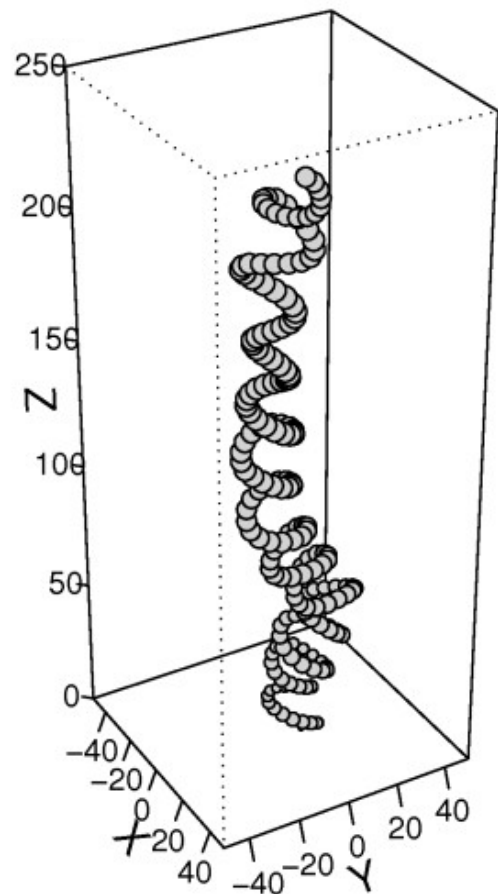
Estimating 3D helical trajectories from 2D data

Gurarie et al. 2010



Helical characteristics are diagnostic of strain (putative genotype)

Gurarie et al. 2010



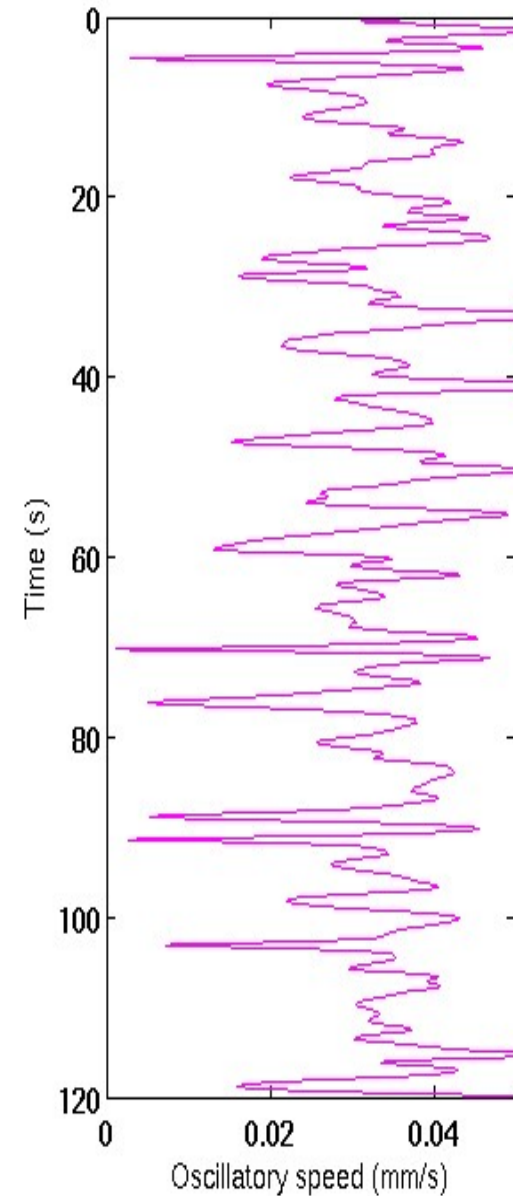
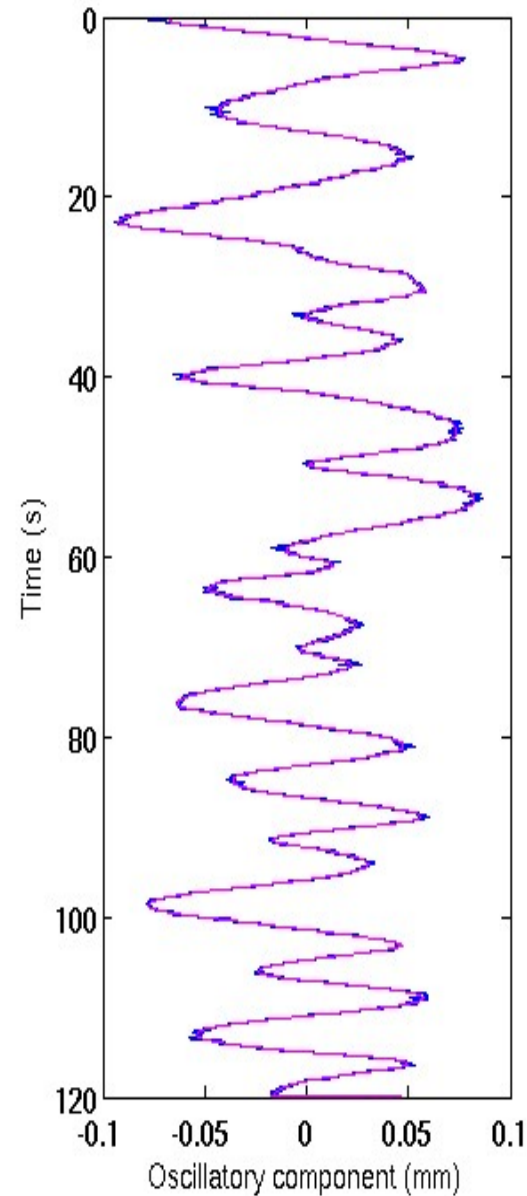
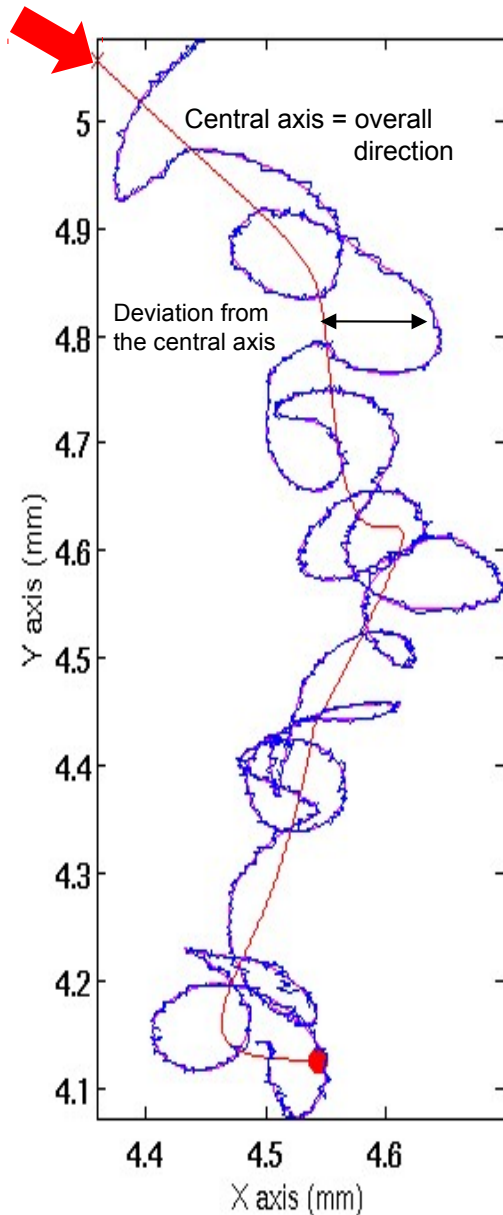
<i>Parameters</i>		
μ_a		mean vertical velocity of \mathbf{V}^a
τ_a		characteristic time scale of \mathbf{V}^a
σ_a		magnitude of stochasticity for \mathbf{V}^a
ω_o		characteristic angular velocity of \mathbf{V}^o
P_o	$1/\omega_o$	characteristic period of rotation of \mathbf{V}^o
τ_o		characteristic time scale of \mathbf{V}^o
σ_o		magnitude of stochasticity for \mathbf{V}^o
\bar{V}_t		mean tangential velocity
θ	$\sin^{-1}(\mu_a/\bar{V}_t)$	mean tangential angle

A) Strains					
Location	Narraganset Bay, RI	Long Island Sound, NY	Puget Sound, WA	Puget Sound, WA	Onagawa Bay, Sea of Japan
Year	1991	1952	2002	2007	1984
N	55	131	31	54	33
B) Parameters					
μ_a (mm s ⁻¹)	0.069 (0.022-0.175)	0.036 (0.005-0.129)	0.047 (0.007-0.088)	0.049 (0.021-0.118)	0.043 (0.013-0.094)
\bar{V}_t (mm s ⁻¹)	0.141 (0.085-0.383)	0.114 (0.05-0.155)	0.055 (0.022-0.103)	0.097 (0.053-0.12)	0.101 (0.061-0.153)

Swimming Statistics of Individual Transitional Cells

Elizabeth Tobin

Start of the path



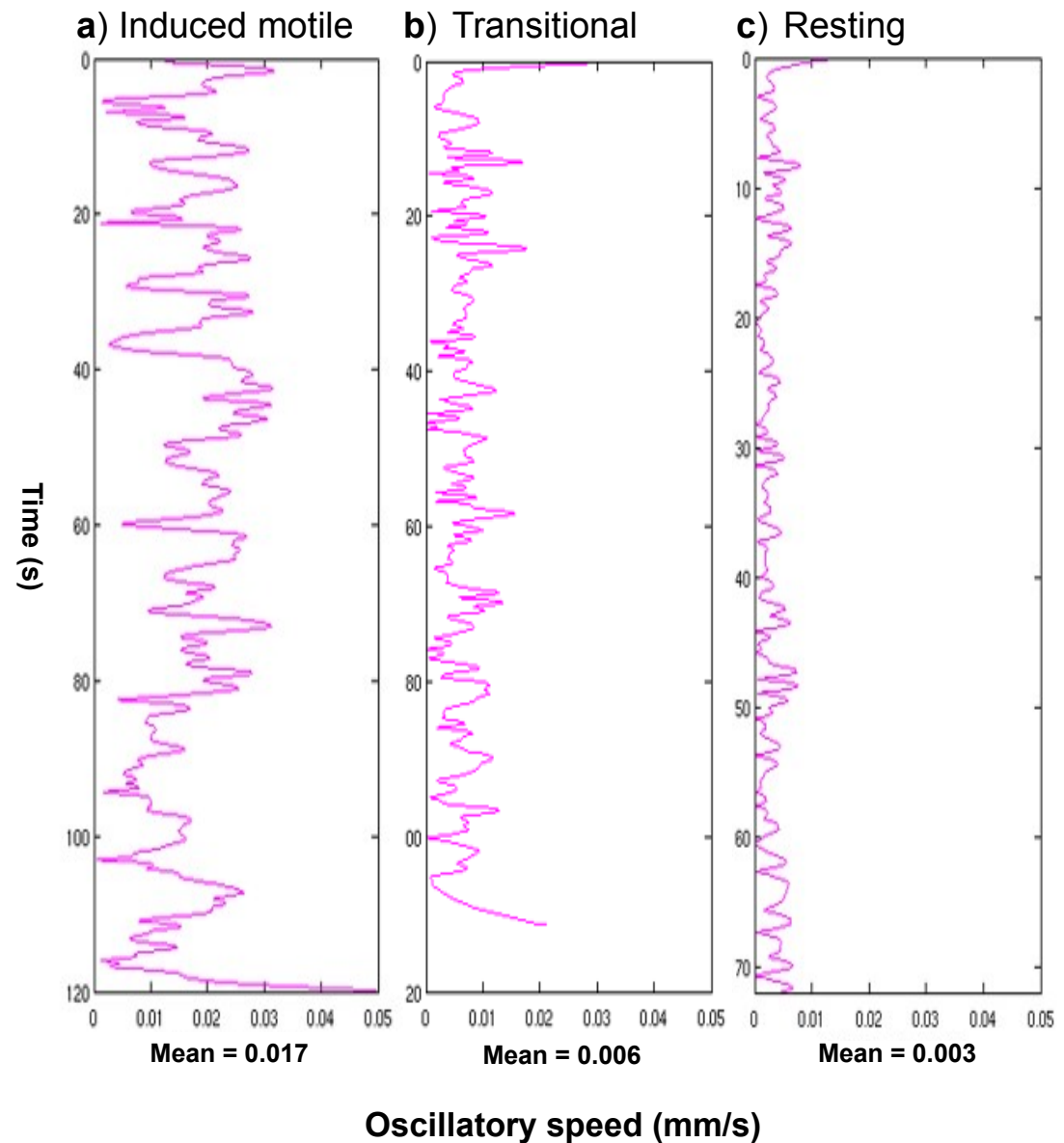
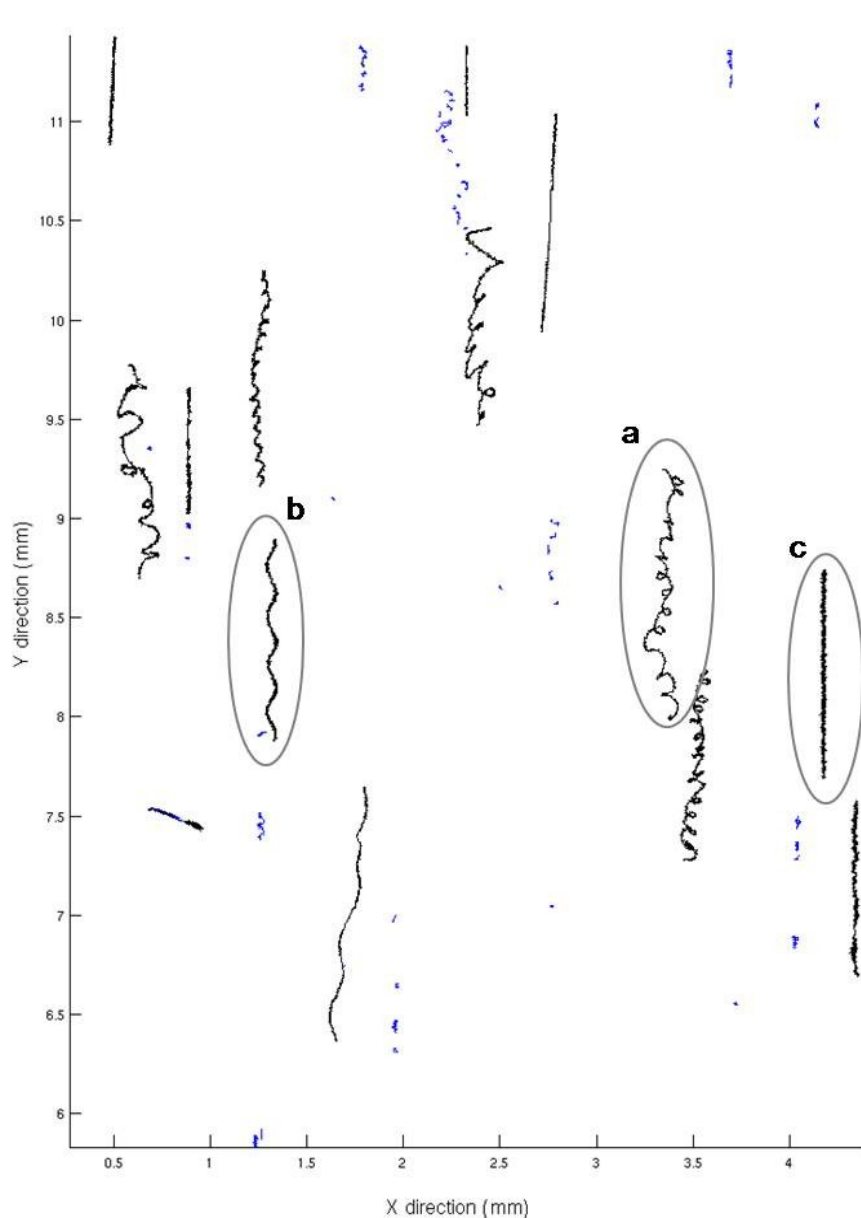
Mean Oscillatory Speed

Mean Gross Speed

Mean Vertical and Horizontal Velocity

Helical characteristics are characteristic of cell state

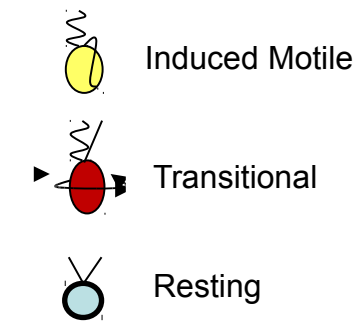
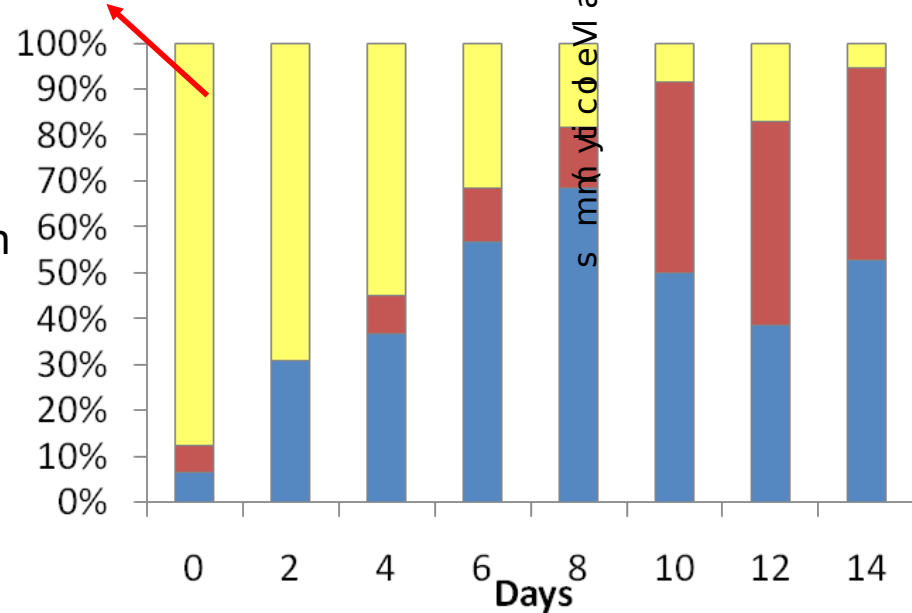
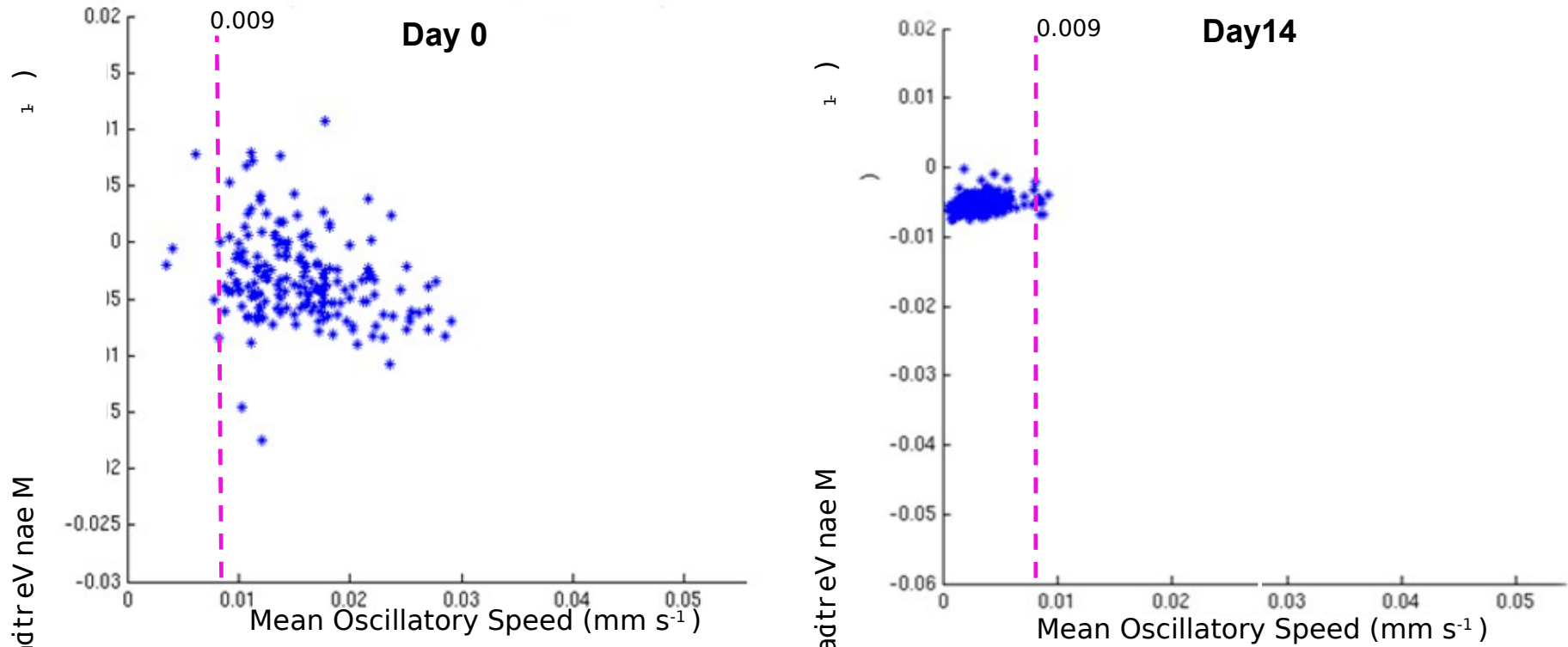
Elizabeth Tobin



- Mostly type **a** on Days 0 and 2
- Mostly types **b/c** on Days 10 -14

Establishing Motion-based Criteria to Classify Cell State

Elizabeth Tobin



Video assay, verified with microscope counts

Identifying Net Movement for Each Cell State

	Induced motile	Transitional	Resting	Vegetative motile	Settling velocity
Mean Oscillatory Speed ($\mu\text{m s}^{-1}$)	13.90 ± 6.1	6.79 ± 2.1	3.77 ± 0.9		
Mean Gross Speed ($\mu\text{m s}^{-1}$)	9.30 ± 4.8	5.87 ± 2.0	4.83 ± 1.7	49 – 66	
Mean Vertical Velocity ($\mu\text{m s}^{-1}$)	- 4.25 ± 3.5	- 4.16 ± 2.2	- 4.47 ± 1.8	35 – 60	$\approx 4 \mu\text{m s}^{-1}$
Mean Horizontal Velocity ($\mu\text{m s}^{-1}$)	5.45 ± 4.1	2.46 ± 1.7	0.80 ± 0.6	(Bearon et al. 2004)	(Stokes' flow)
N	5712	5161	8434		

Stokes flow (Low Reynolds number)


$\rho_p = 1.105 \text{ g cm}^{-3}$ (Wada et al. 1985)

$\rho_f = 1.025 \text{ g cm}^{-3}$ (28 psu)

$R = 5 \mu\text{m}$

$\mu = 1.376 \times 10^{-3}$ (28 psu, 10°C)

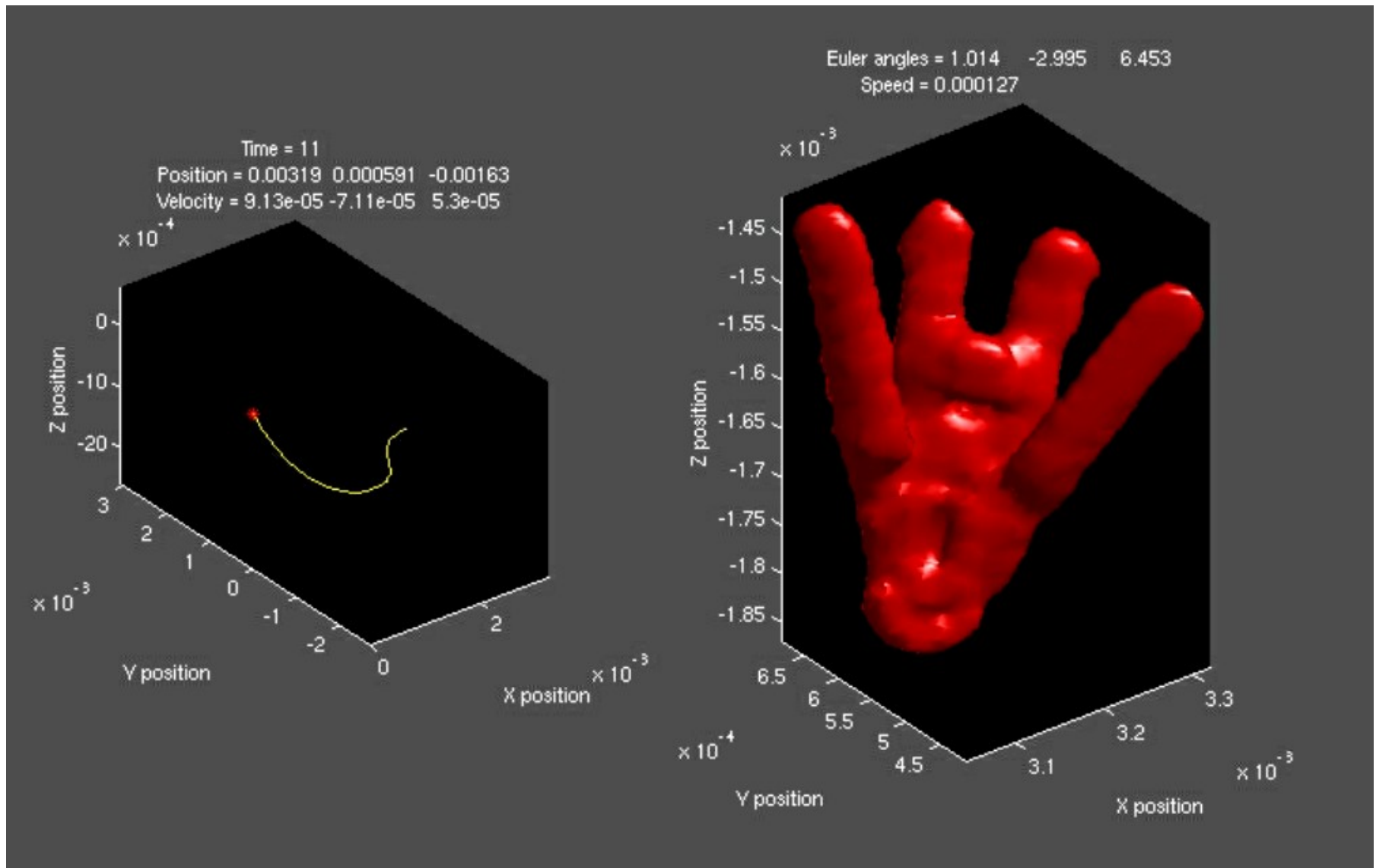
$$V_s = \frac{2(\rho_p - \rho_f)}{9\mu} g R^2$$

 *Heterosigma* cells under going life-stage transition have net downward movement comparable to passive sinking.

Echinoderm larva (*Dendraster excentricus* – sand dollar)
Karen Chan



Confocal-based low-Re model of swimming biomechanics

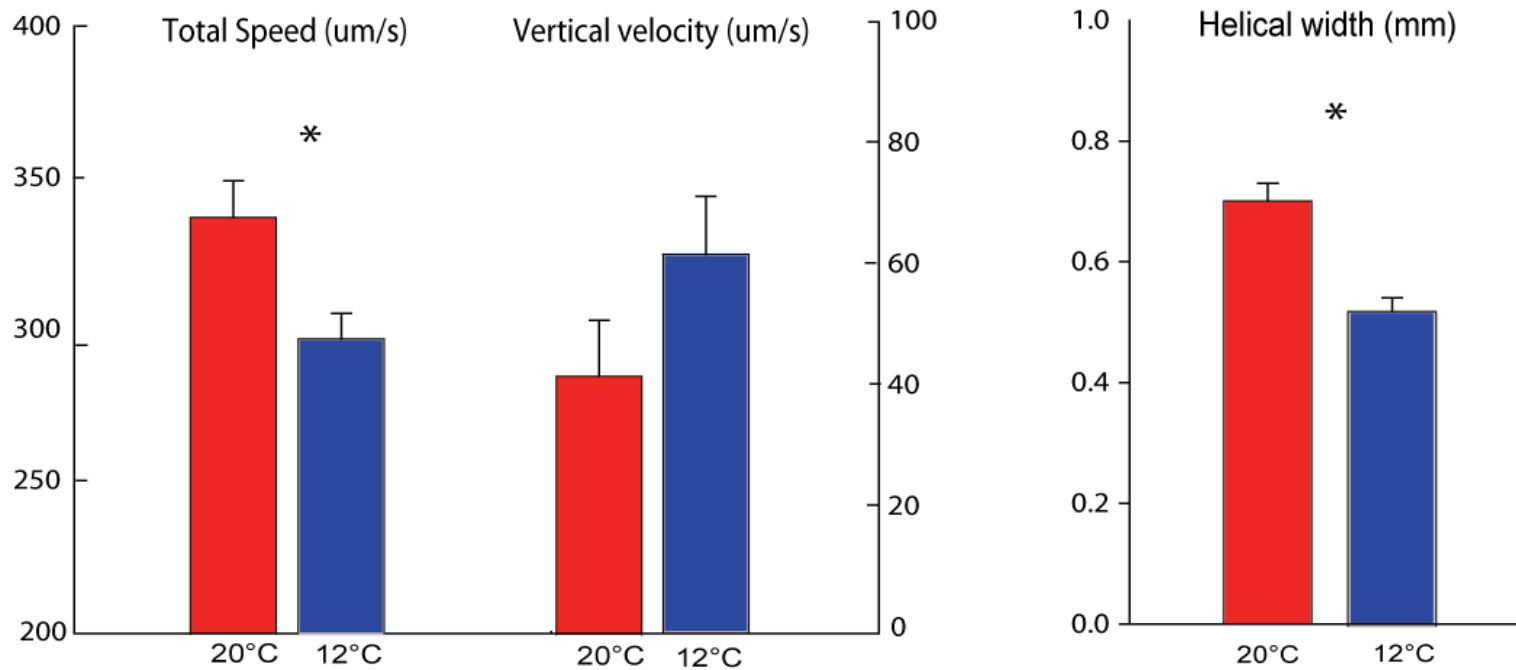
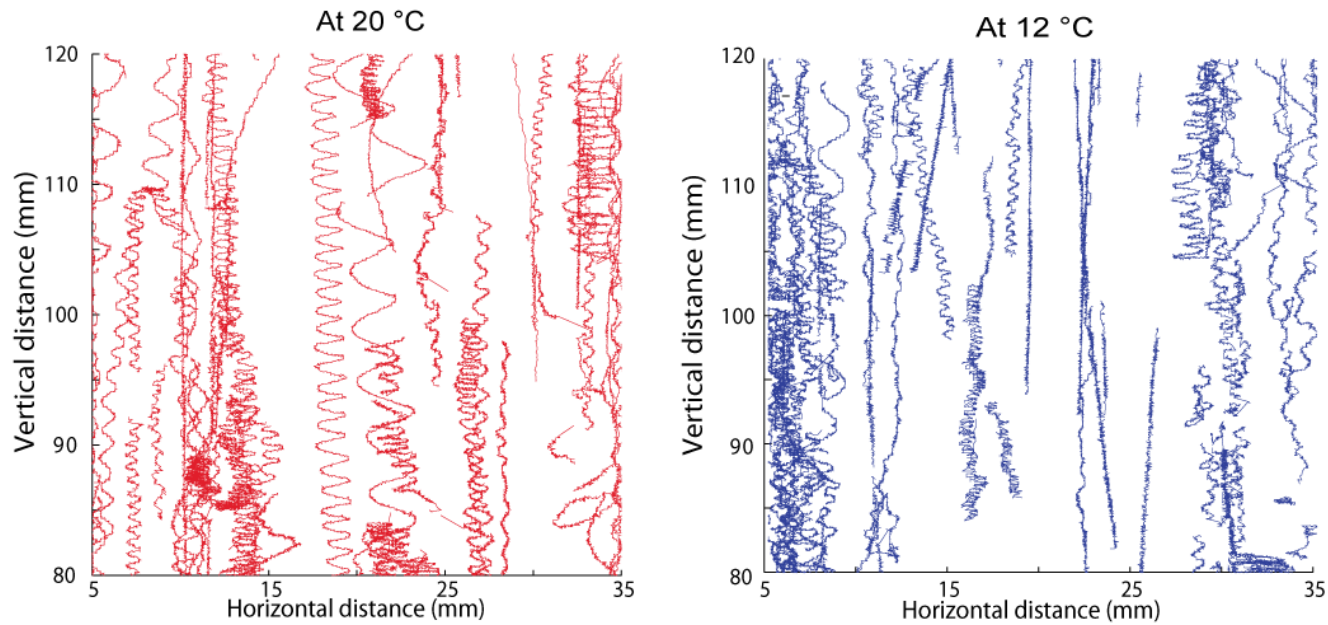


Confocal-based low-Re model of swimming biomechanics



Sand dollar larvae decrease helical widths to conserve upward speed in colder water

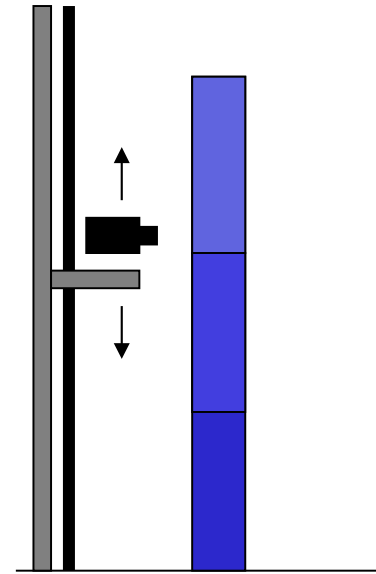
Karen Chan



Thin layer formation by natural plankton communities - 2-halocline experiment

Experimental design:

1. Plankton pump water from 1 m and 4 m (Friday Harbor Laboratories dock)
2. Dilute with R.O. water to create two 1 psu haloclines
3. Sequential vertical profiles with video tracker



Whole
unfiltered
sea water
29.5 psu



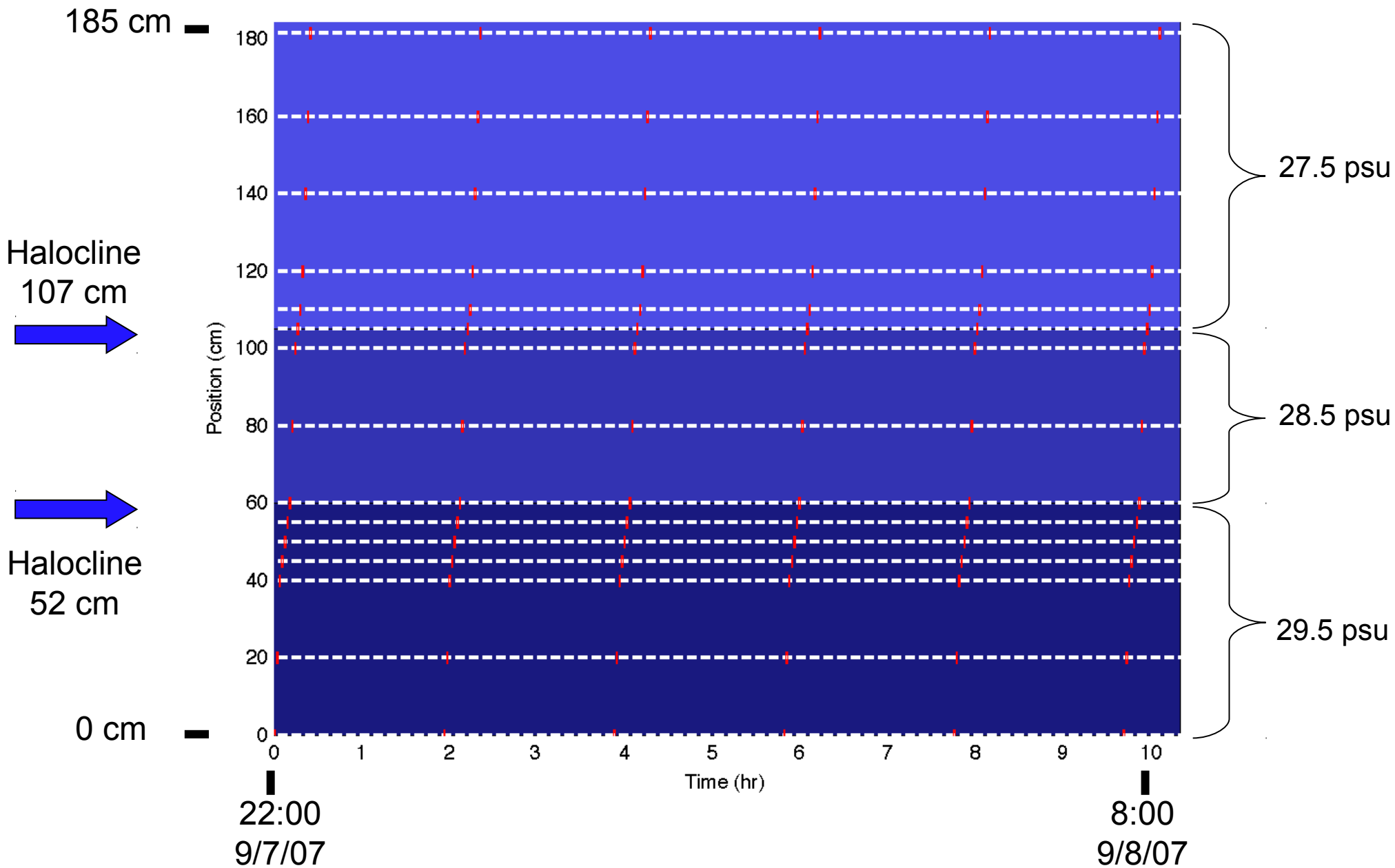
R.O. water +
unfiltered
sea water,
28.5 psu



R.O. water +
unfiltered
sea water,
27.5 psu

Thin layer formation by natural plankton communities - 2-halocline experiment

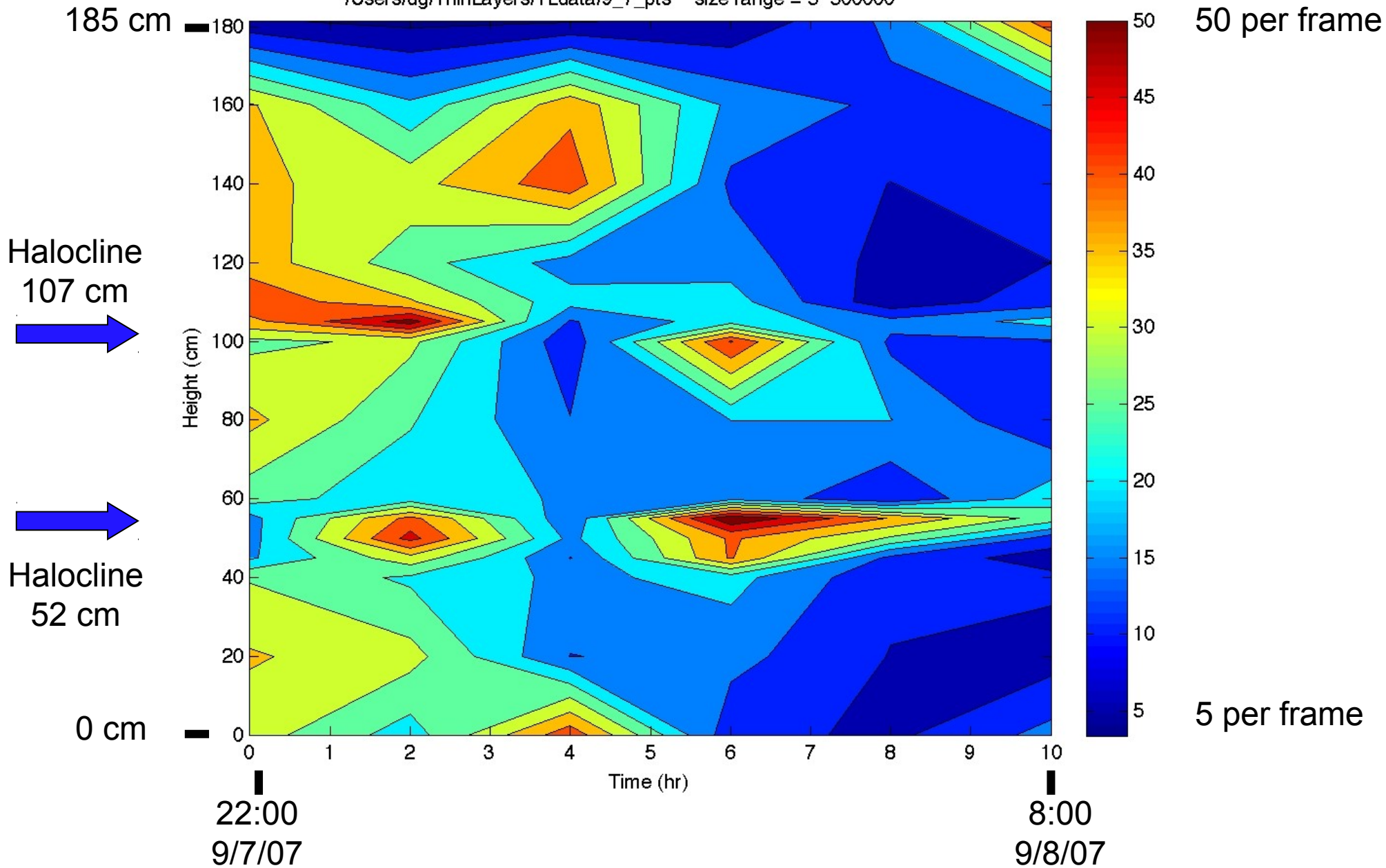
15 vertical stations, 6 profiles, 90 video clips, $> 1.8 \cdot 10^6$ particle localizations



Thin layer formation by natural plankton communities - 2-halocline experiment

Total organism/particle counts

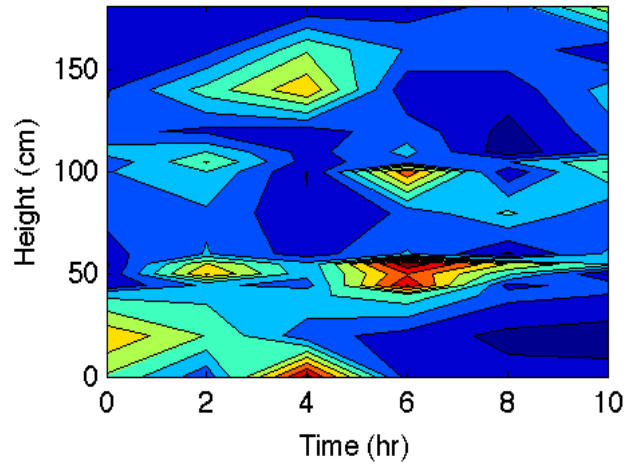
/Users/dg/ThinLayers/TLdata/9_7_pts size range = 5 500000



Abundance and vertical flux for motile and passive particles

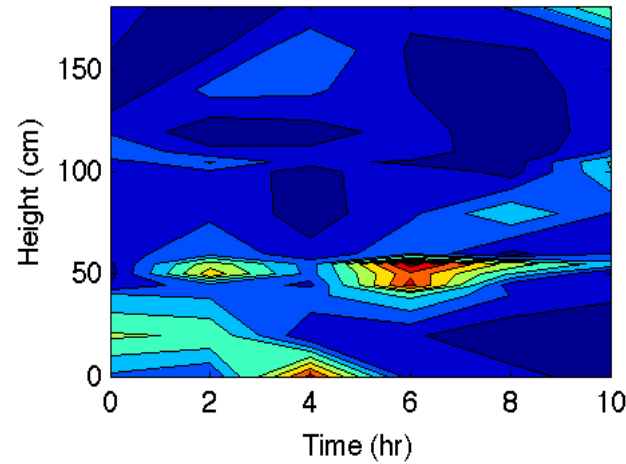
All paths

size range = 20 50



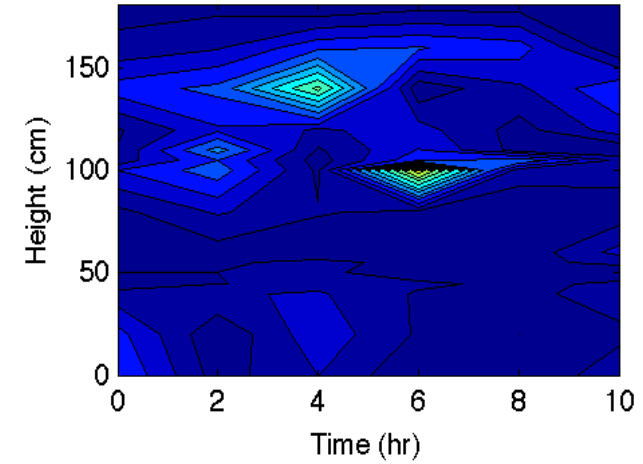
Motile organisms

size range = 20 50

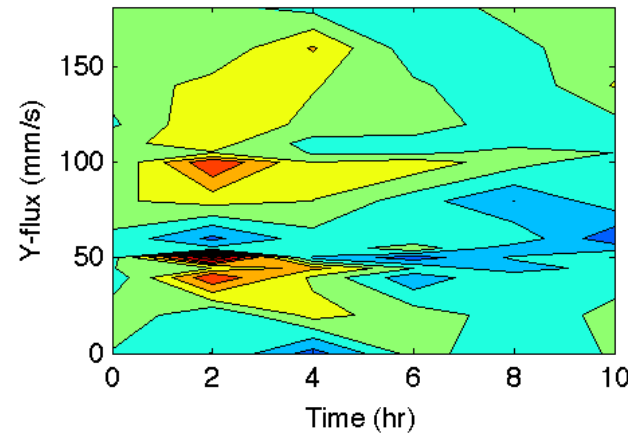


Passive particles

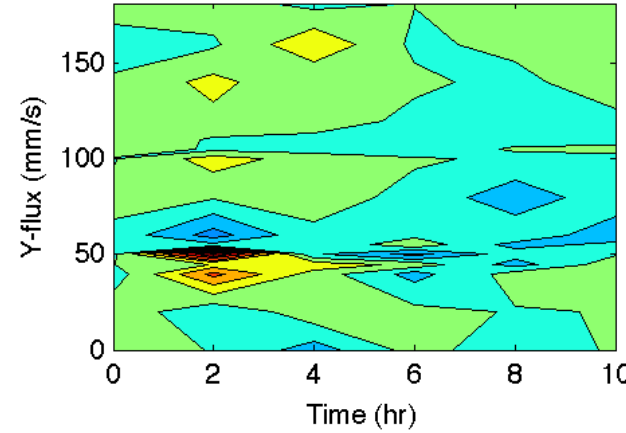
size range = 20 50



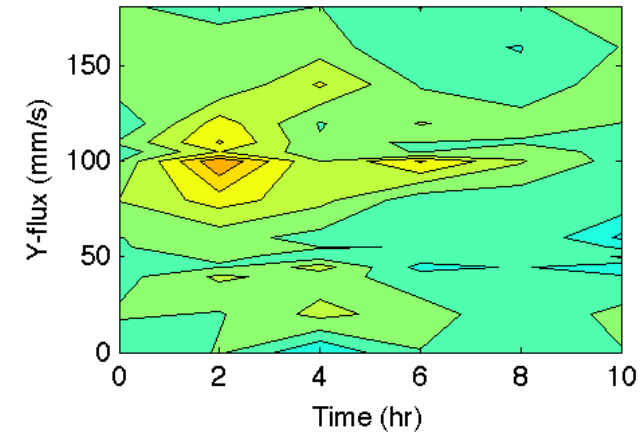
size range = 20 50



size range = 20 50



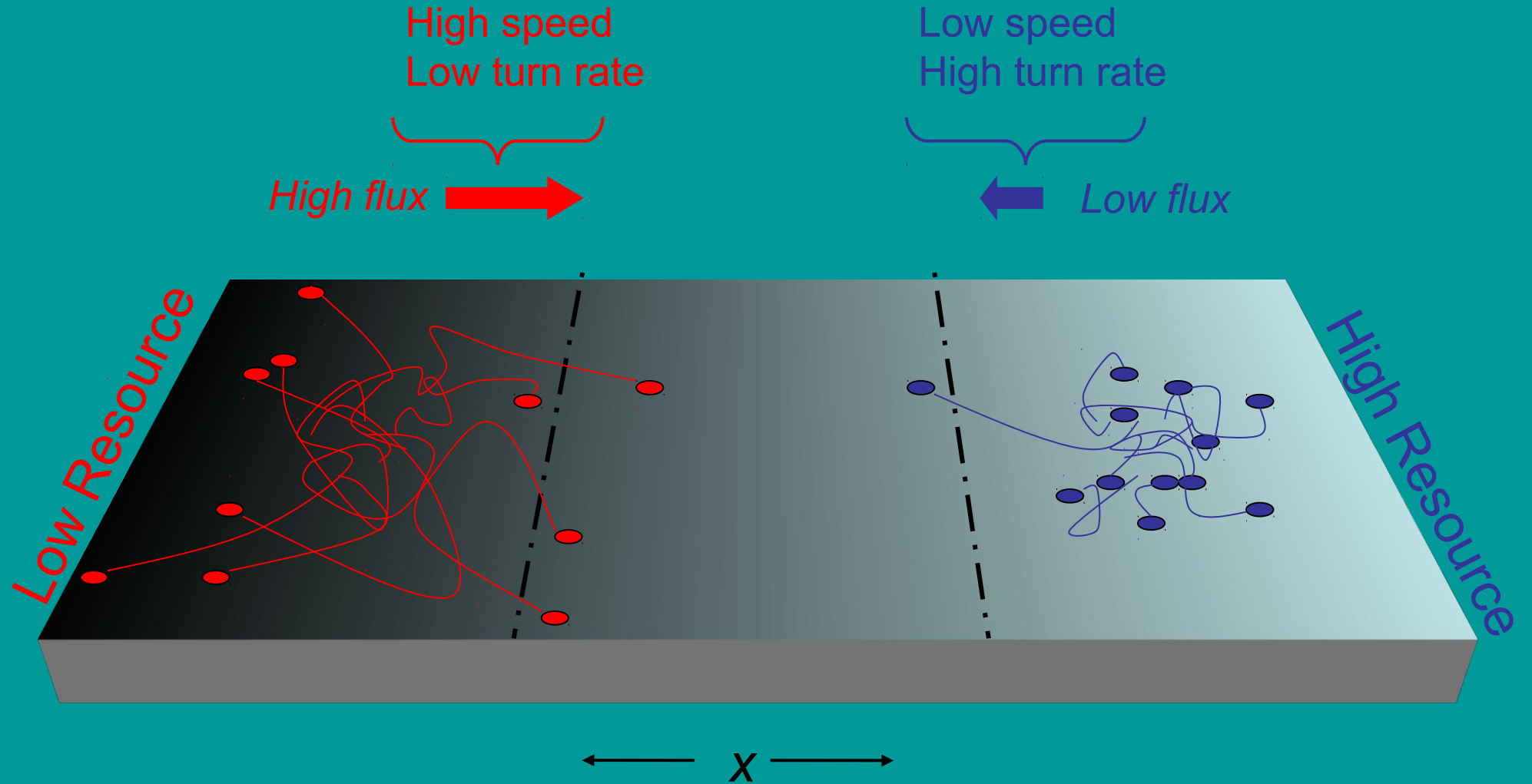
size range = 20 50



Upward fluxes (# mm/s)

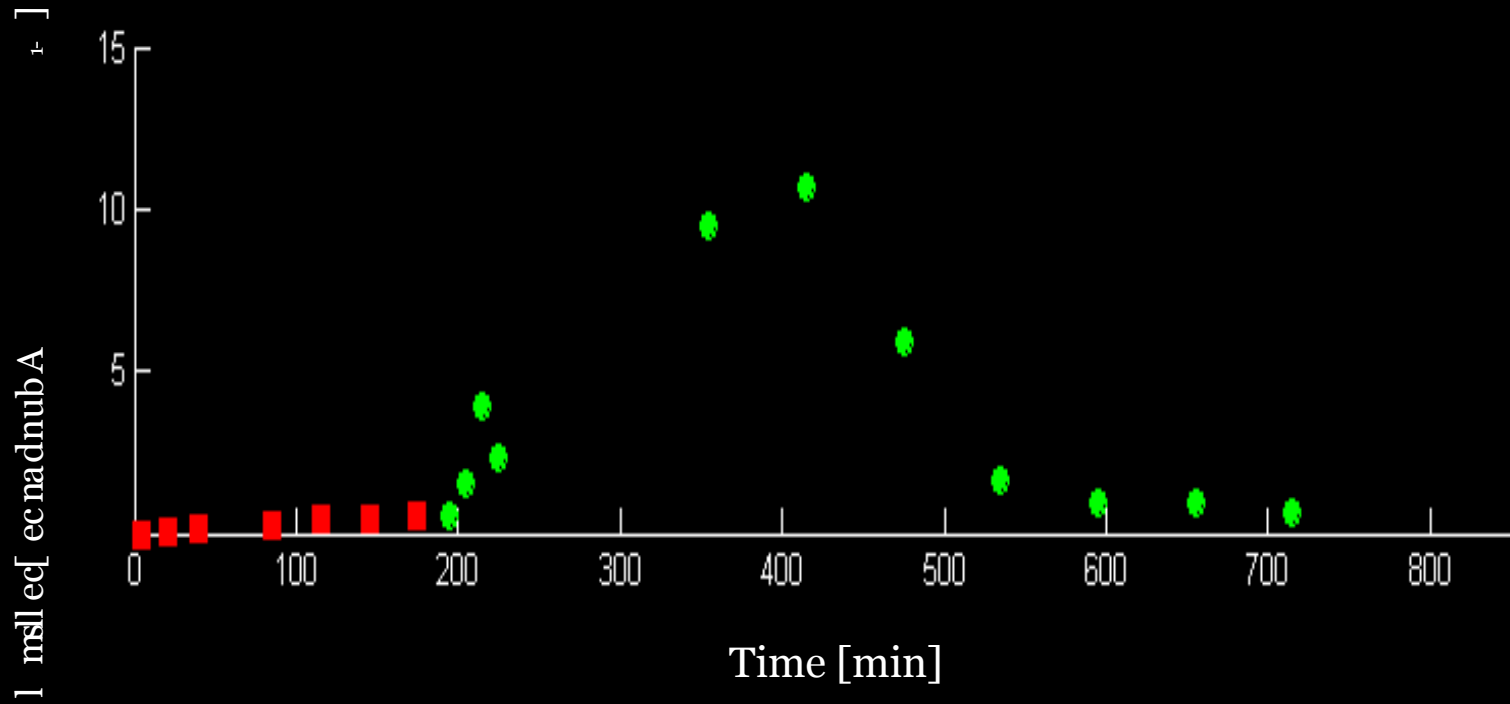
Foraging with sensory constraints

Biased Random Walks Mediated by Internal States



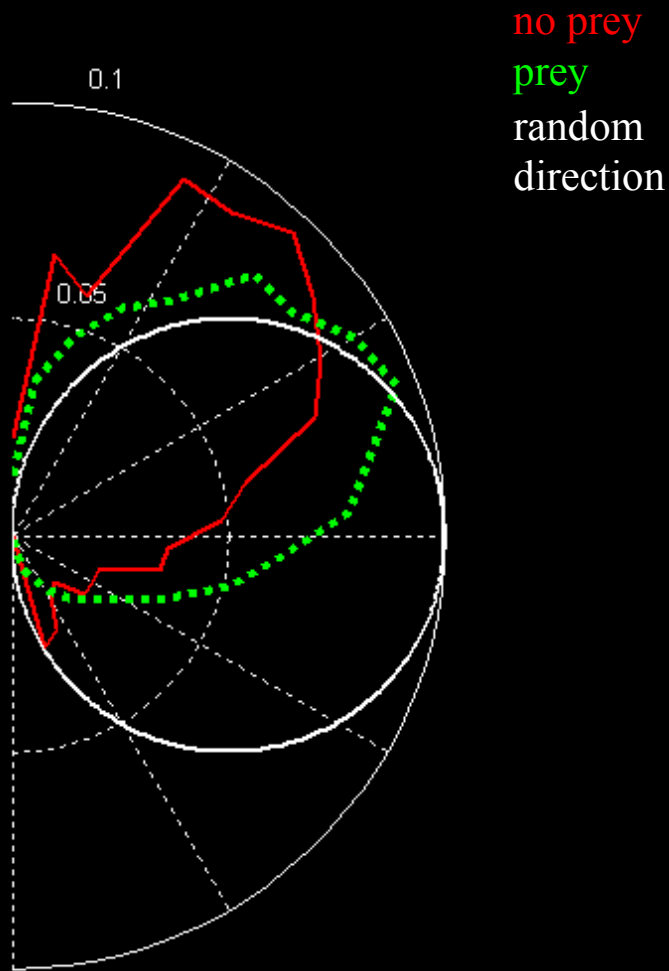
Heterotrophic protists aggregate to thin layer of phytoplankton prey cells

Susanne Menden-Deuer

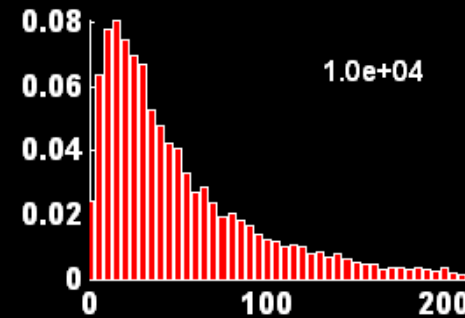


Mechanisms of aggregation: behavioral responses to food

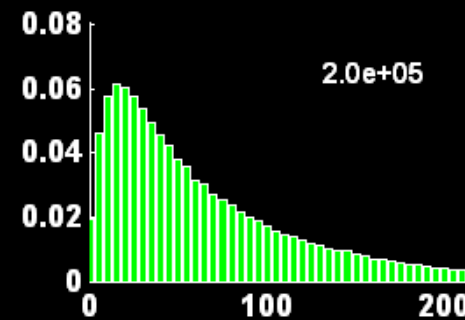
Direction



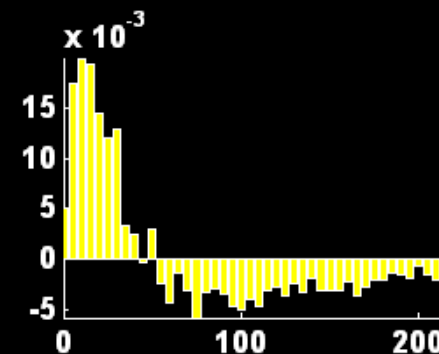
Turning rate [deg. s⁻¹]



no prey

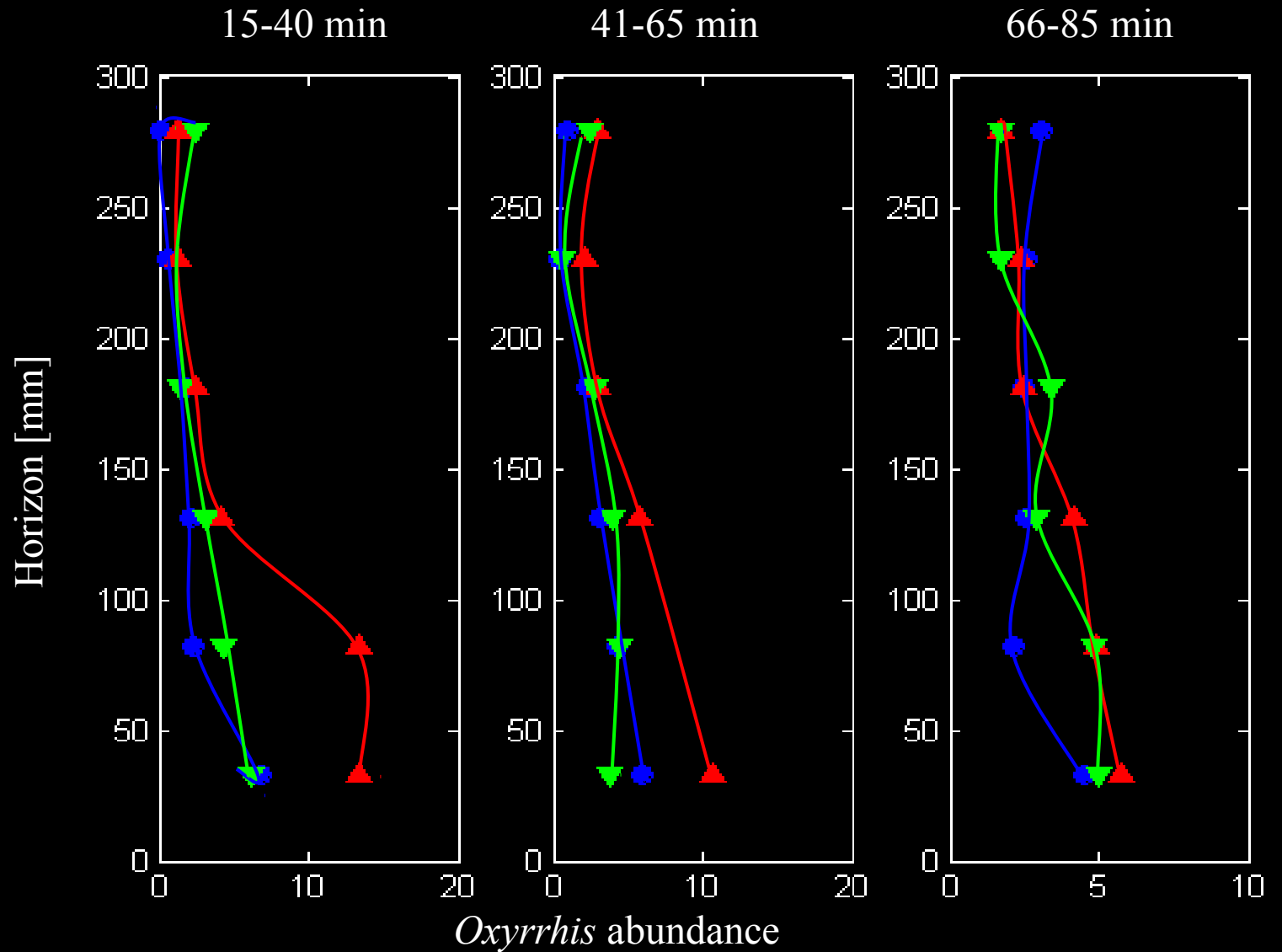
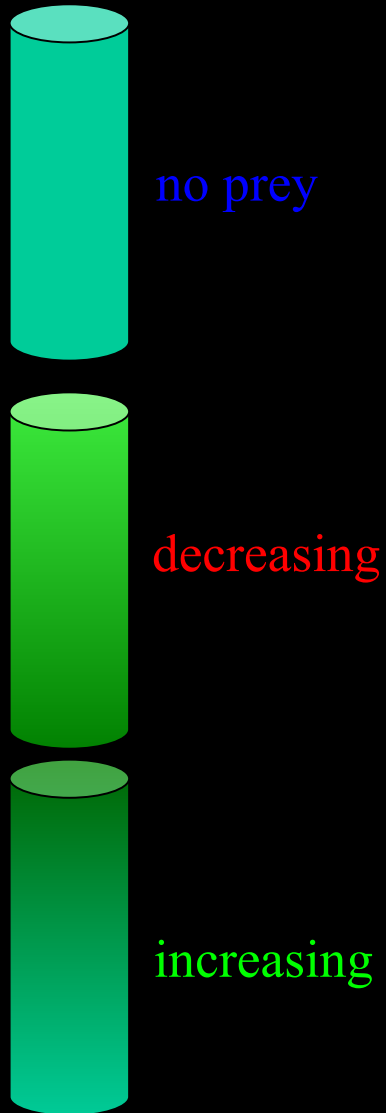


with prey

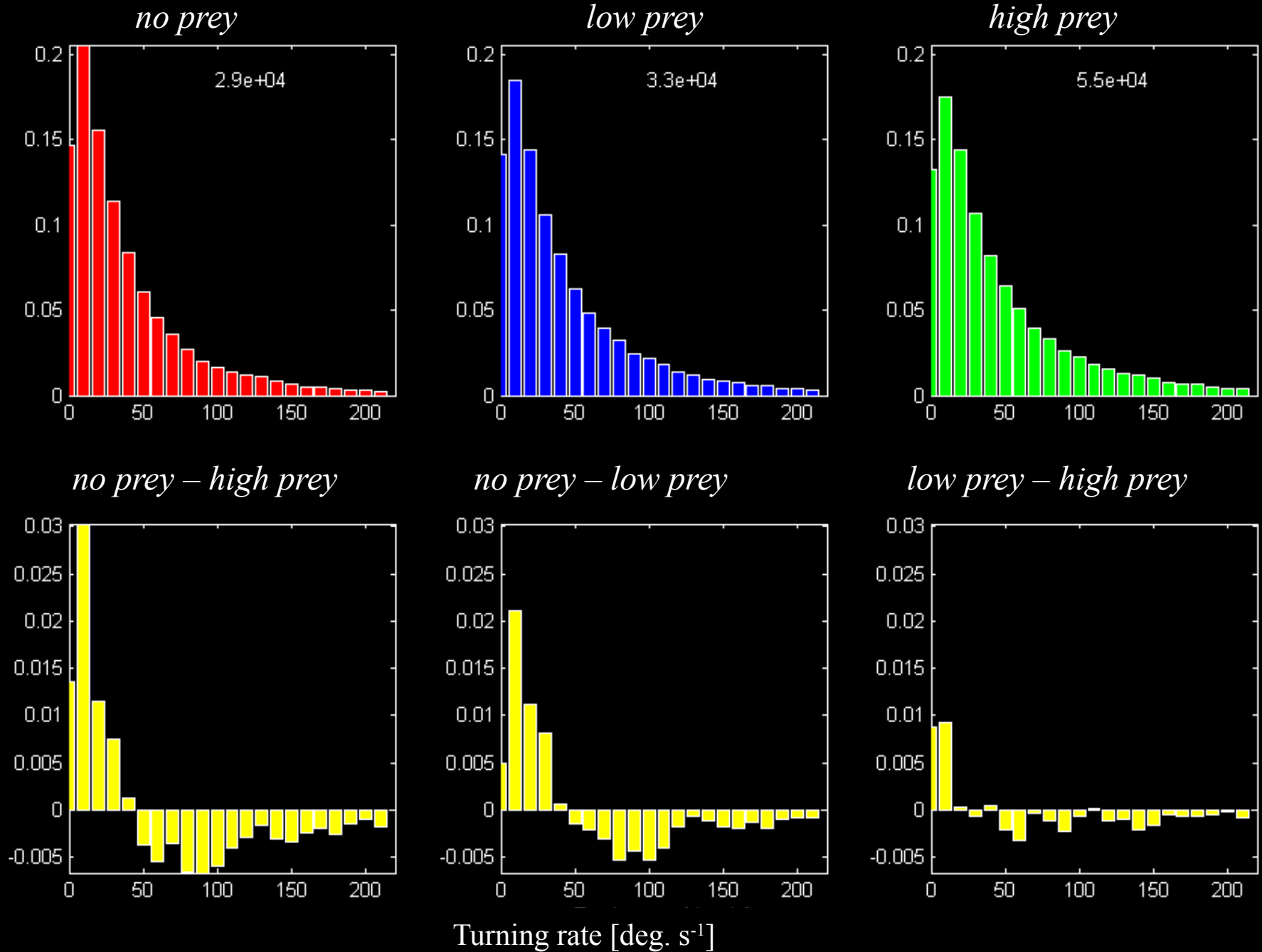


behavioral
response

Population redistribution in prey gradients

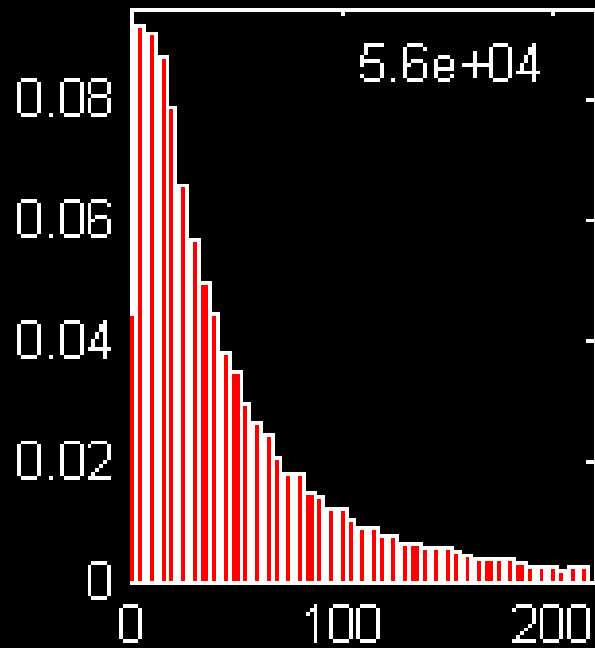


Frequency of high turning rates increases with ambient prey concentration

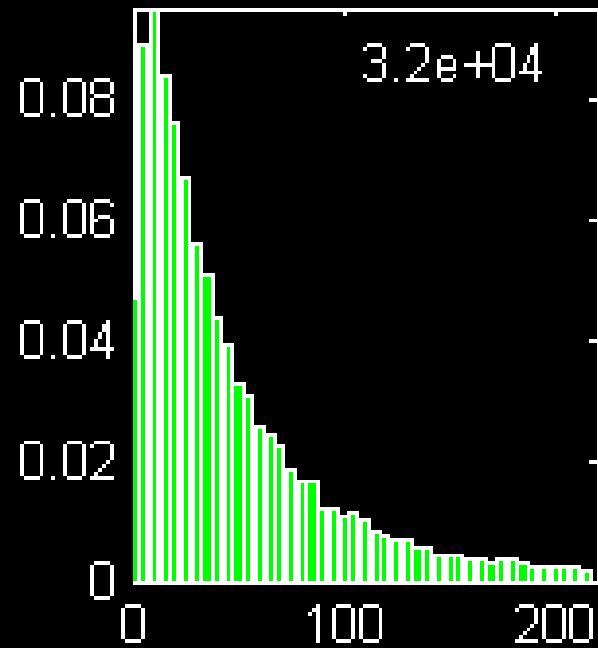


Distributions of turning rates do not differ depending on gradient direction

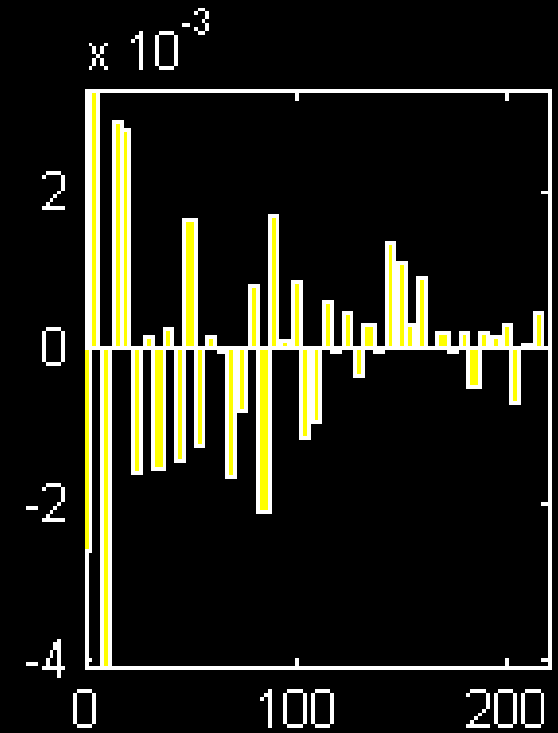
decreasing



increasing



gradient response



Turning rate [deg. s⁻¹]