



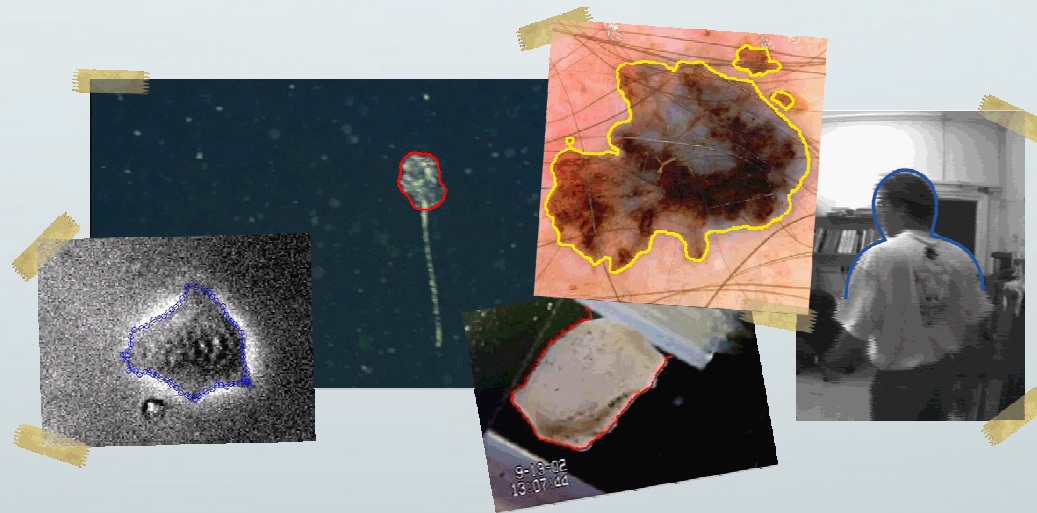
Dynamic shape detection and analysis of deformable structures in biomedical imaging

*Dip. di Ing. Dell'Informazione
Dottorato in Scienza e Tecnologia dell'Informazione – XXII Ciclo*

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Alberto Silletti - 13th April 2010

Outline

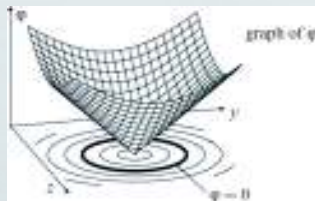
- Introduction: what is a shape?
- The shape detection problem
 - generalized active contours
 - reticular shape detection
- Dynamic shape detection
- Analysis
- Synthesis

Introduction: What is a shape?

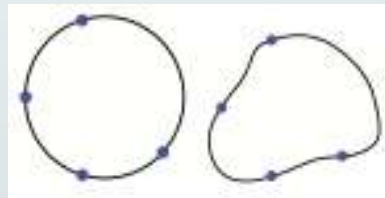
A fuzzy concept, many definitions occurs over scientific and common literature:

- *The spatial arrangement of something as distinct from its substance; “geometry is the mathematical science of shape” (wordReference)*
- *A shape is all the geometrical information that remains when location, scale and rotational effects are filtered out from an object (George Kendall, mathematician)*

$$X \xrightarrow{\text{sensors}} I \xrightarrow{\text{intuition}} \Phi(X) \xrightarrow{\text{representation}} \mathcal{R}(\Phi)$$



Level sets



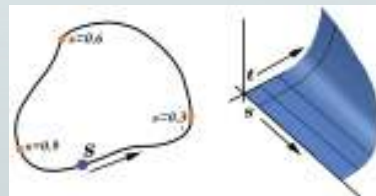
Landmark



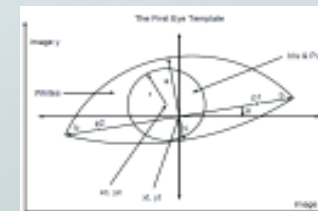
Polygons



Voxels or pixels



Spline



Deformable model

The shape detection problem

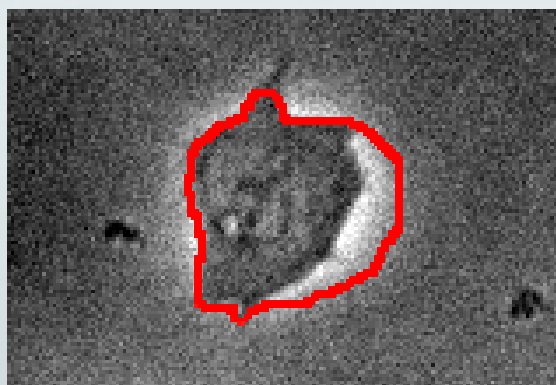
Given an image I (data), and chosen a shape representation model $\mathcal{R}(\Phi)$ the shape detection problem consist of:

$$\mathcal{R}(\Phi) = \arg \min_{\mathcal{R}(\Phi_c) \in \mathcal{S}} \mathcal{E}(I, \Phi_c)$$

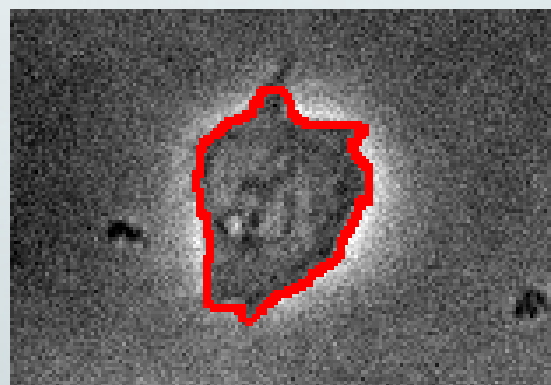
where \mathcal{E} is a suitable energy function and \mathcal{S} is the space of the possible representations.

Shape detection involves:

- *the design of \mathcal{E}*
- *the procedure to minimize over \mathcal{S}*



non optimal \mathcal{E}



optimal (minimum) \mathcal{E}

Generalized Active Contours 1/4

An Active Contour is a curve $C(s) \in \mathbb{R}^d$, $s \in [0, 1]^d$

evolving in pseudo-time according to an associated energy:

$$\mathcal{E}(C) = \mathcal{S}(C) + \mathcal{P}(C)$$

$$\mathcal{S}(C) = \int_{\partial C} \alpha(s) \left| \frac{\partial C}{\partial s} \right|^2 + \beta(s) \left| \frac{\partial^2 C}{\partial s^2} \right|^2 ds$$

$$\mathcal{P}(C) = \int_{\partial C} \mathcal{D}[I] ds$$

Generalized Active Contours extends the framework:

$$\mathcal{E}_g(C) = \mathcal{S}_g(C) + \mathcal{P}_g(C)$$

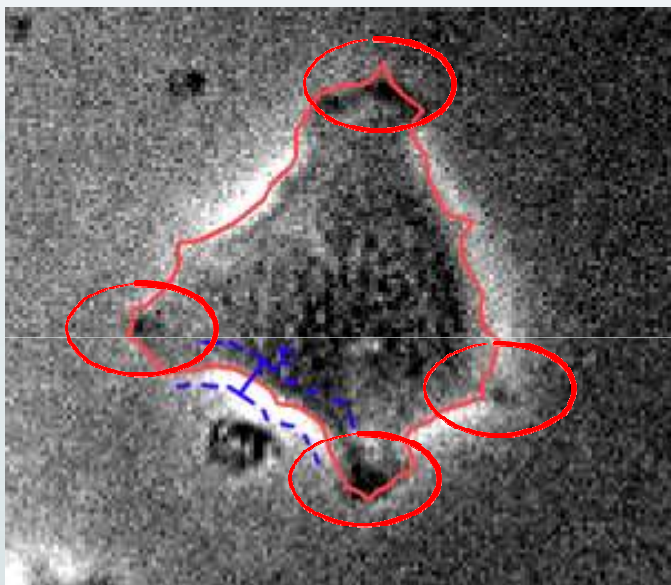
$$\int_{\partial C} \alpha(s) \left| \frac{\partial C}{\partial s} \right|^2 + \beta(s) \left| \frac{\partial^2 C}{\partial s^2} \right|^2 ds + \sum_{i=1}^{n_f} \gamma_i \mathcal{F}_i$$

$$\int_{\partial C} \mathcal{D}[I] ds + \sum_{j=1}^{n_g} \lambda_j \mathcal{G}_j$$

where \mathcal{F}_i and \mathcal{G}_j are suitable energy terms, each of those addressing a peculiar aspect of the shape (such as color, texture or central moments).

Generalized Active Contours 2/4

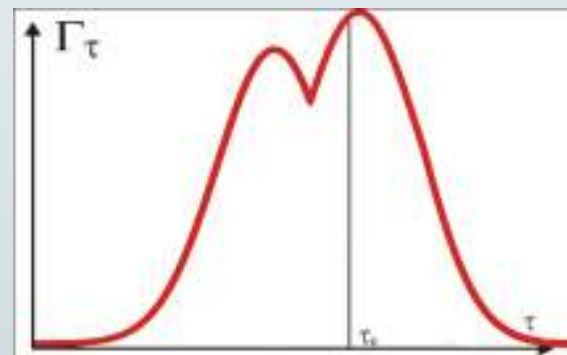
Generalized Active Contours build \mathcal{F}_i and \mathcal{G}_j energies *as probability density functions*.
Let's pretend you want to capture the dark protein appendices:



1. Choose a metric to characterize them, such as color at boundary:

$$\tau(C) = \int_{\partial C_{\pm r}} I(s) ds$$

2. Infer the probability density Γ_{τ} - analytically, empirically or by training -



3. Build an energy term as:

$$\mathcal{E}_{\tau} = -\Gamma_{\tau}(\tau_c)$$

Generalized Active Contours 3/4

The work provides a Toolbox with several build-in energy terms

Energy Toolbox II/II		
Name	Figure	Definition
Area()	<p>Active Contour C(s) Active Contour's interior region ϕ</p> <p>Area energy drives the Contour towards an area size c</p>	$E = -\frac{1}{\sigma\sqrt{2\pi}} \cdot e^{-\frac{c^2}{2\sigma^2}}$ $\mu = \int_{C^-} area(C) - c d\omega$ $c = \text{desired area size}$
Inflate() Deflate()	<p>growing Active Contour C(s)</p> <p>Inflate (or Deflate) the Active Contours forcing a larger (smaller) area. μ_{max} is an upper bound, possibly unreachable</p>	$E = -\frac{\min(\mu, \mu_{max})}{\mathcal{N}}$ $\mu = \int_{C^-} ds$ $\mu_{max} = \text{max value for } \mu$ $\mathcal{N} = \text{normalizing constant}$
Inter Class Color Variance()	<p>external color variance Active Contour C(s) Active Contour's internal color variance</p> <p>Inter Class Color Variance enhances the color variance between the interior and the exterior</p>	$E = -\frac{1}{\sigma\sqrt{2\pi}} \cdot e^{-\frac{c^2}{2\sigma^2}}$ $\mu = \Sigma(I(C^+)) - \Sigma(I(C^-)) $
Internal Minimal (Maximal) Color Variance()	<p>Active Contour C(s) Active Contour's internal color variance</p> <p>Internal Minimal (or Maximal) Variance favours a small (large) internal color variance</p>	$E = -\frac{\min(\mu, \mu_{max})}{\mathcal{N}}$ $\mu = \Sigma(I(C^-))$ $\mu_{max} = \text{max value for } \mu$ $\mathcal{N} = \text{normalizing constant}$
Inter Class Color()	<p>external color Active Contour C(s) Active Contour's internal color</p> <p>Inter Class Color enhances the color difference between the interior and the exterior</p>	$E = -\frac{1}{\sigma\sqrt{2\pi}} \cdot e^{-\frac{c^2}{2\sigma^2}}$ $\mu = \left \int_{C^-} I(C) d\omega + \int_{C^+} I(C) d\omega \right $

Energy Toolbox I/II		
Name	Figure	Definition
Inner Color()	<p>non matching color Active Contour C(s) Active Contour's interior region ϕ</p> <p>Inner Color energy drives the Contour to have an internal color c</p>	$E = -\frac{1}{\sigma\sqrt{2\pi}} \cdot e^{-\frac{c^2}{2\sigma^2}}$ $\mu = \int_{C^-} I(\omega) - c d\omega$ $c = \text{desired inner color}$ $C^- = \text{interior of } C$
Outer Color()	<p>desired external color Active Contour C(s) Active Contour's interior region ϕ</p> <p>Outer Color energy drives the Contour towards an external color c The areas of c color remain out of the Contour</p>	$E = -\frac{1}{\sigma\sqrt{2\pi}} \cdot e^{-\frac{c^2}{2\sigma^2}}$ $\mu = \int_{C^+} I(\omega) - c d\omega$ $c = \text{desired inner color}$ $C^+ = \text{exterior of } C$
Surrounding Inner Color()	<p>outer color Active Contour C(s) Active Contour's neighborhood</p> <p>Surrounding Inner Color forces the Contour to have a small internal neighbor of color c</p>	$E = -\frac{1}{\sigma\sqrt{2\pi}} \cdot e^{-\frac{c^2}{2\sigma^2}}$ $\mu = \int_{C^-} I(\omega) - c d\omega$ $c = \text{desired outer color}$
Surrounding Outer Color()	<p>outer color Active Contour C(s) Active Contour's neighborhood</p> <p>Surrounding Outer Color forces the Contour to have a small external neighbor of color c The areas of c color remain out of the Contour</p>	$E = -\frac{1}{\sigma\sqrt{2\pi}} \cdot e^{-\frac{c^2}{2\sigma^2}}$ $\mu = \int_{C^+} I(\omega) - c d\omega$ $c = \text{desired outer color}$
Spikeness()	<p>Spikeness()</p> <p>Depending on the sign, force a spiky appearance or a cloudy one c</p>	$E = -\frac{\min(\mu, \mu_{max})}{\mathcal{N}}$ $\mu = \int_C \kappa(s) ds$ $\kappa(s) = \frac{\partial^2 c(s)}{\partial s^2}$ $\mu_{max} = \text{max value for } \mu$ $\mathcal{N} = \text{normalizing constant}$

Generalized Active Contours 4/4

Assuming all the density function are Gaussian, we rewrite the energy terms as:

$$\Gamma = \mathcal{N}(\vec{\mu}, \vec{\Sigma}) = \mathcal{N} \left(\begin{bmatrix} \mu_1 \\ \mu_2 \\ \vdots \\ \mu_i \\ \vdots \\ \mu_{n_f+n_g} \end{bmatrix}, \begin{bmatrix} \Sigma_{11} & \Sigma_{12} & \Sigma_{13} & \cdots \\ \Sigma_{21} & \Sigma_{22} & \Sigma_{23} & \cdots \\ & \Sigma_{31} & & \ddots \\ \vdots & & & \Sigma_i \\ & & & & \ddots \\ & & & & & \Sigma_{n_f+n_g, n_f+n_g} \end{bmatrix} \right)$$

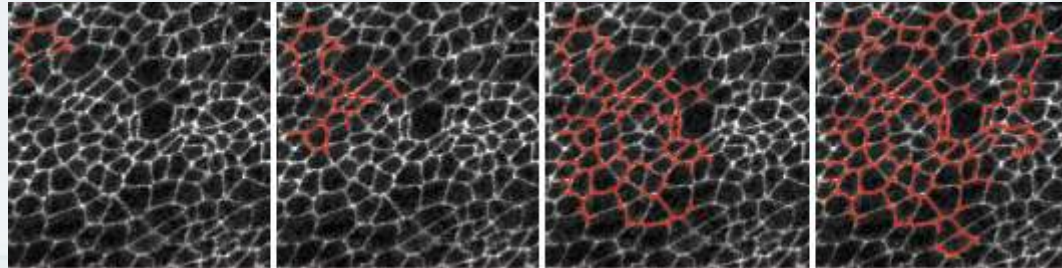
We can assess the quality of our Snaking algorithm looking at the eigenvectors: if they are close to zero, then we have a robust algorithm.

We can also simplify the matrix over certain thresholds

$$T \cdot \begin{bmatrix} \sigma_1 & 0 & 0 & \cdots \\ 0 & \sigma_2 & 0 & \cdots \\ 0 & 0 & \ddots & \\ \vdots & & & \sigma_i \\ & & & & \ddots \\ & & & & & \sigma_k \end{bmatrix} \cdot T^T$$

Reticular shape detection

-The random walk agents approach-

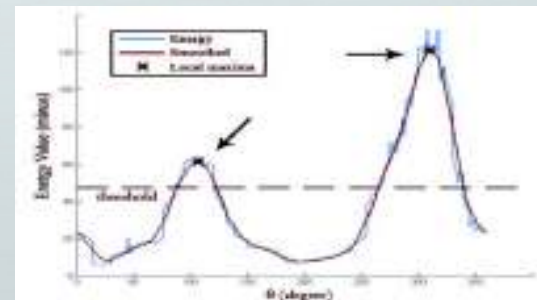
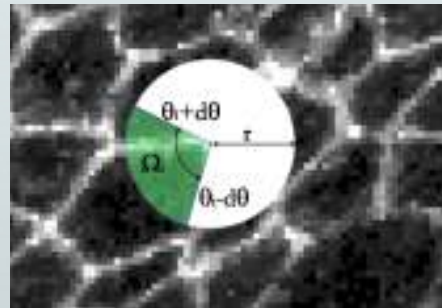
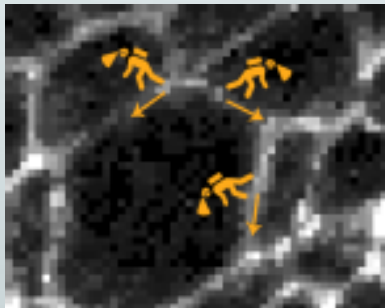


Idea: many Random Walk Agents \mathcal{A}_i flood the frame, each of them finding a path. Each agent has a position \mathbf{p} and an energy $\hat{\mathcal{E}}$.

Agents locally minimize an energy term $\hat{\mathcal{E}}$, a local approximation of \mathcal{E} . \mathcal{E} is unknown and difficult to design, whereas $\hat{\mathcal{E}}$ is

$$\hat{\mathcal{E}}(\theta_i) = \frac{\int_{\Omega_i} \sqrt{(I(\omega) - 1)^2} d\omega}{\int_{\Omega_i} d\omega}$$

$$\mathbf{p}(t+1) = \mathbf{p}(t) + k \cdot \mathbf{g}(\hat{\mathcal{E}})$$



Reticular shape detection

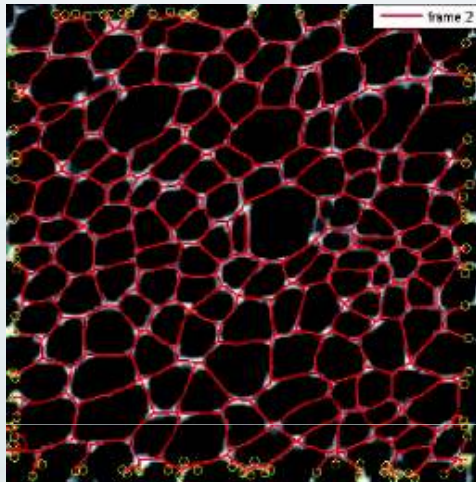
-The random walk agents approach-

Code snippet 4 - Random Walk Agents main loop

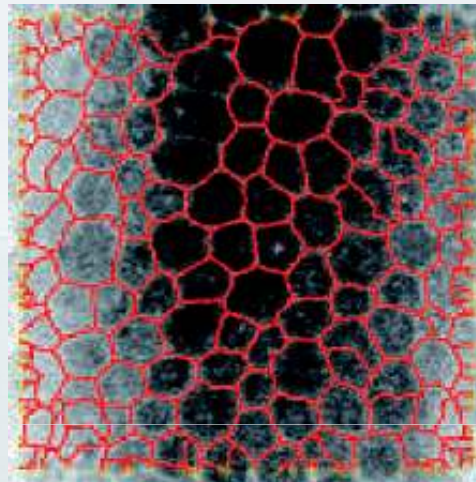
```
1  PQ ← initPriorityQueue() //create a priority queue
2  G ← initGraph() //create an empty graph
3  while PQ not empty() {
4      A ← dequeue(PQ) //extract the best agent
5      valid, border ← validateAgent(A, I, G)
6      if valid
7          G ← add2Graph(G, A) //add to the graph
8      if (valid and not(border)) {
9          E ← computeEnergyFunction(I, Ai)
10         D ← pickDirections(E)
11         for k=1 to |D| {
12             Ak ← moveAgent(A, Dk)
13             PQ ← enqueue(Ak, PQ) }
14     }
15 }
```

Reticular shape detection

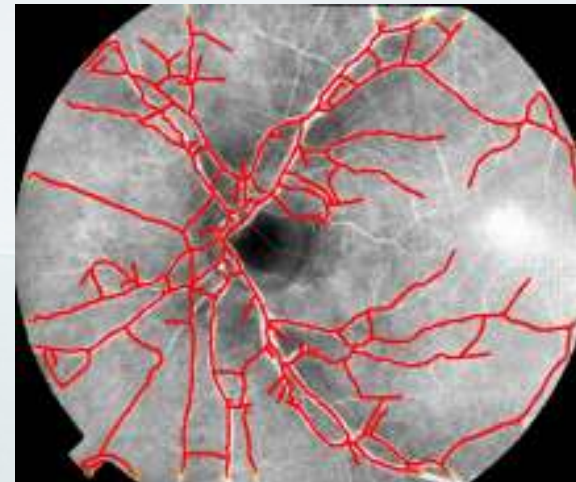
-The random walk agents approach-



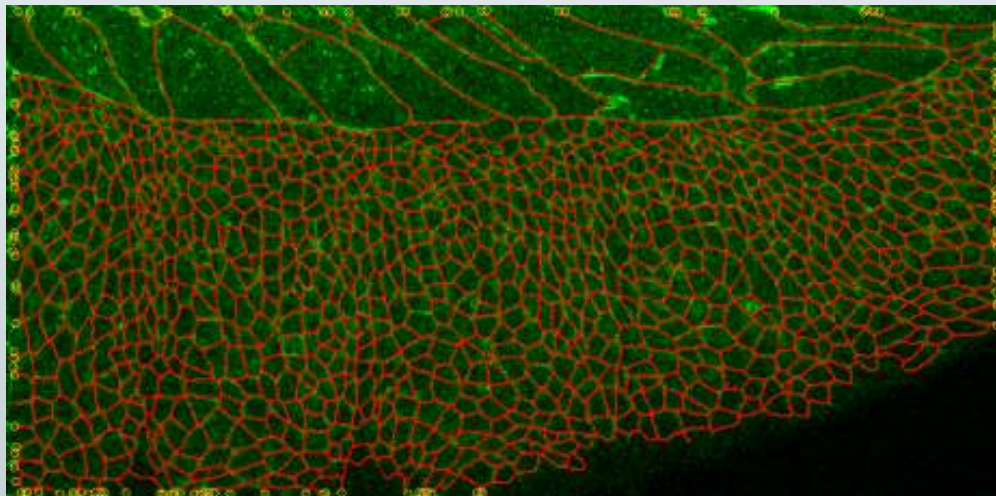
Drosophila wing



Corneal fundus



Retina



Drosophila developmental stage

Dynamic shape detection



Given a sequence of temporal coherent images $\{I_1, \dots, I_t, \dots, I_N\}$, Dynamic shape detection is the problem of detecting the shape in each frame:

$$\begin{aligned} \Phi_1 &= \arg \min_{\Phi_c \in \mathcal{S}} \mathcal{E}(I_1, \Phi_c) \\ &\quad \dots \\ \Phi_N &= \arg \min_{\Phi_c \in \mathcal{S}} \mathcal{E}(I_N, \Phi_c) \end{aligned}$$

Images temporal coherence: $I_{t+1} = I_t + \Delta_t^{t+1}$

Shape temporal coherence: $\Phi_{t+1} = \Phi_t + \delta_t^{t+1}$

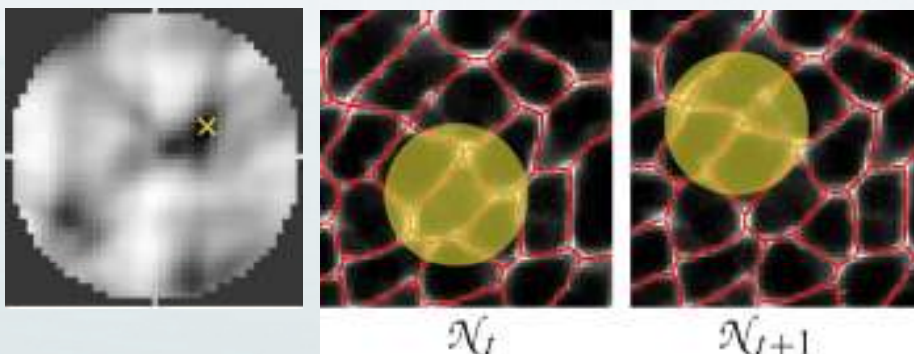
The optimum δ_t^{t+1} satisfies: $\Phi_{t+1} = \Phi_t + \delta_t^{t+1} = \arg \min_{\Phi_c \in \mathcal{S}} \mathcal{E}(I_{t+1}, \Phi_c)$

We compute δ_t^{t+1} using the *J-maps* approach

Dynamic shape detection

-the J-maps approach-

Chosen a location $p_t = [x, y] \in I_t$, a $J_{p_t}(\Delta x, \Delta y)$ map is scalar function whose values are (inversely) related to the probability of p_t being translated into $p_{t+1} = [x + \Delta x, y + \Delta y] \in I_{t+1}$



$$J_{p_t}(\Delta x, \Delta y) = \int_P |\mathcal{N}_t(p_t) - \mathcal{N}_{t+1}(p_{t+1})| dp$$

$$\delta_t^{t+1} = \arg \min_{v_x, v_y} J_{p_t}$$

where \mathcal{N}_t is around $p_t = [x, y] \in I_t$,
and \mathcal{N}_{t+1} is around $p_{t+1} = [x + \Delta x, y + \Delta y] \in I_{t+1}$

- The J-maps describe the deformation of I_t into I_{t+1}
- The args -min of the J-maps is the Optical Flow Field
- The args-min computed in the location of points of $\Phi(X)$ is δ_t^{t+1}

Dynamic shape detection

-the J-maps approach-

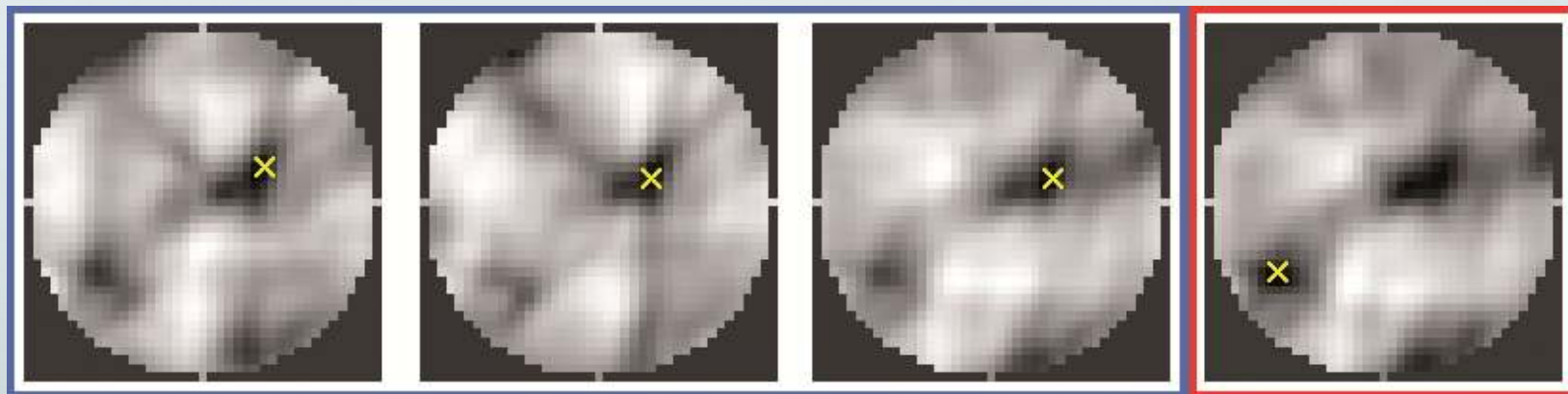
Correction: we impose $\frac{\partial \delta_t^{t+1}}{\partial p} \rightarrow 0$

The J-maps correction is an iterative steps. For each map J_p we consider a set of neighbor maps $\{J_{p_1}, J_{p_2} \dots J_{p_n}\}$, and compute the global minima for each of them:

$$\{(v_{x1}, v_{y1}), (v_{x2}, v_{y2}), \dots (v_{xn}, v_{yn})\}$$

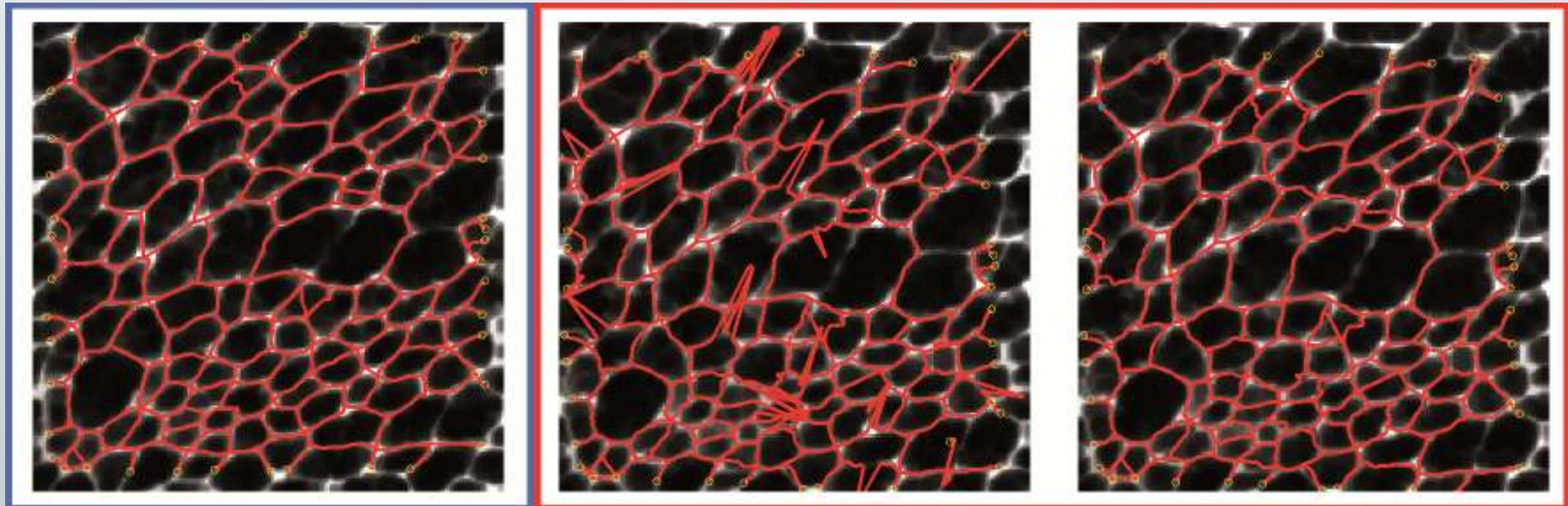
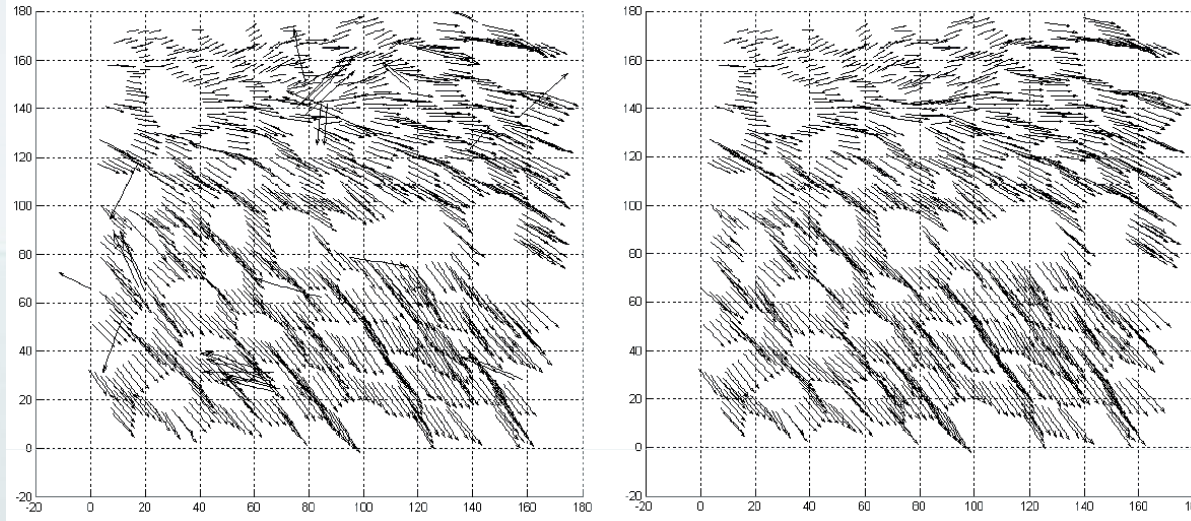
We correct J_p according to: $\bar{J}_p = J_p \cdot (1 - \mathcal{G}([\bar{v}_x, \bar{v}_y], \Sigma_{xy}))$

where \mathcal{G} is a Gaussian build on $\{(v_{x1}, v_{y1}), (v_{x2}, v_{y2}), \dots (v_{xn}, v_{yn})\}$



Dynamic shape detection

-the J-maps approach-

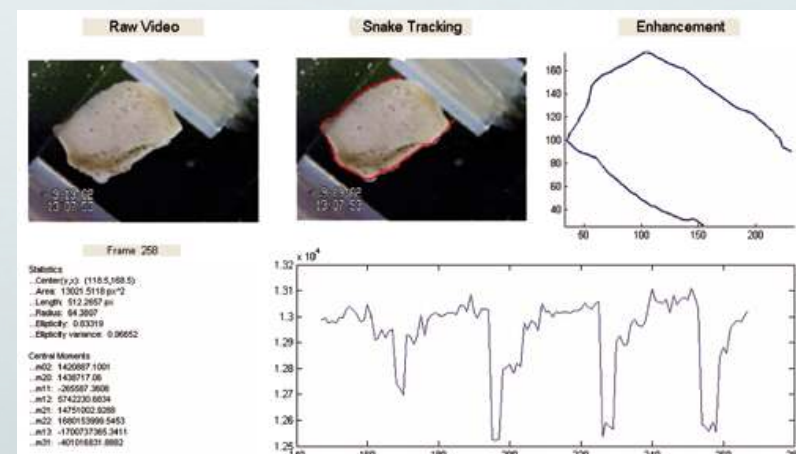
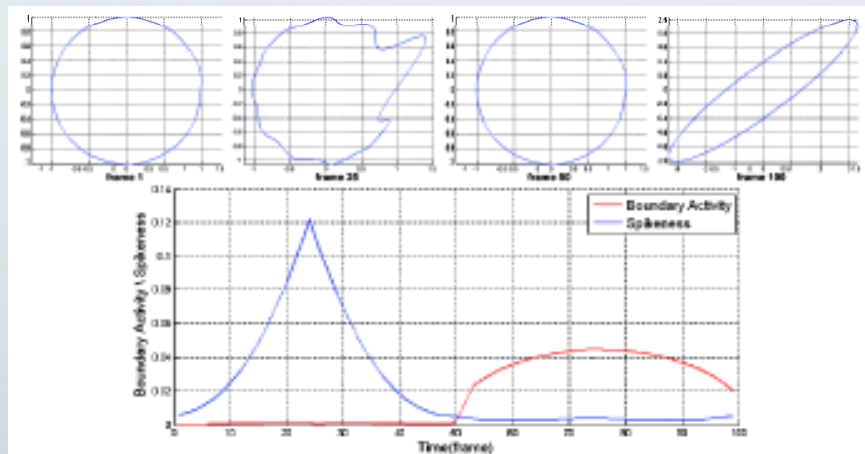


Shape analysis

Shape analysis is the process of extracting metrics and descriptors from shape or sequence of shapes.

$$\{\mathcal{R}(\Phi_0), \dots, \mathcal{R}(\Phi_t), \dots, \mathcal{R}(\Phi_N)\} \xrightarrow{\text{analysis}} \mathbb{R}^k$$

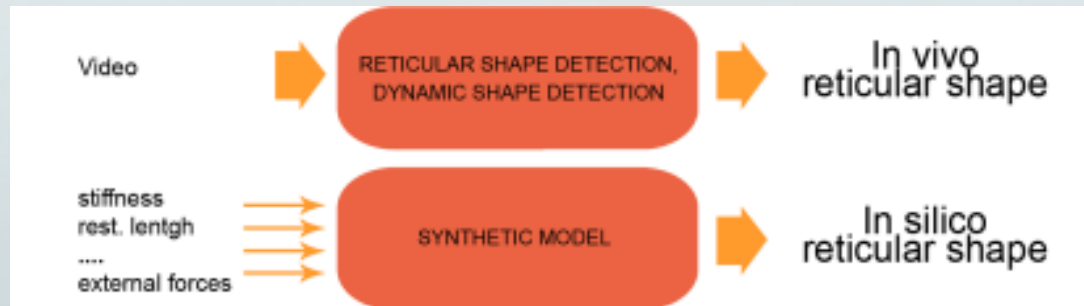
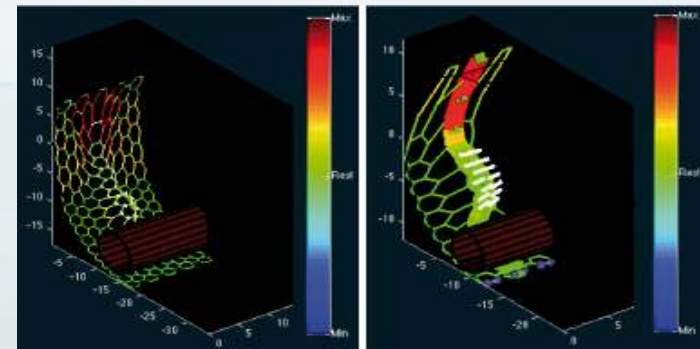
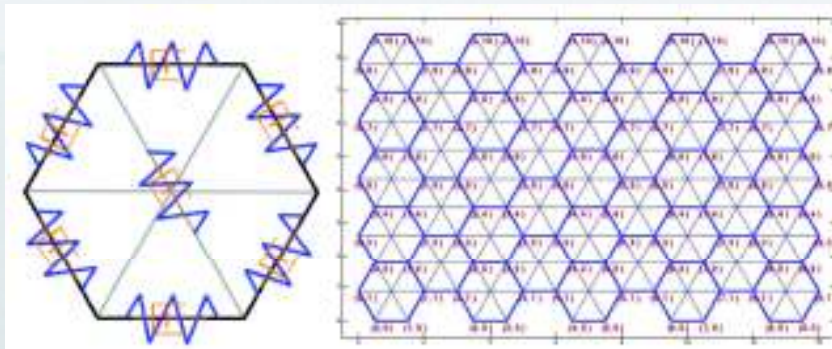
The idea is to forget about the Representation Model and use only a reduced set of “numbers” to capture properties of the shape. These values are forces, lengths, labels, descriptors, medical diagnosis, classifications.



Shape synthesis

The synthesis step involves the creation of a model of the structure of interest.,
mechanical, chemical, stastical...

We build a *mechanical model* for the Drosophila epithelium.



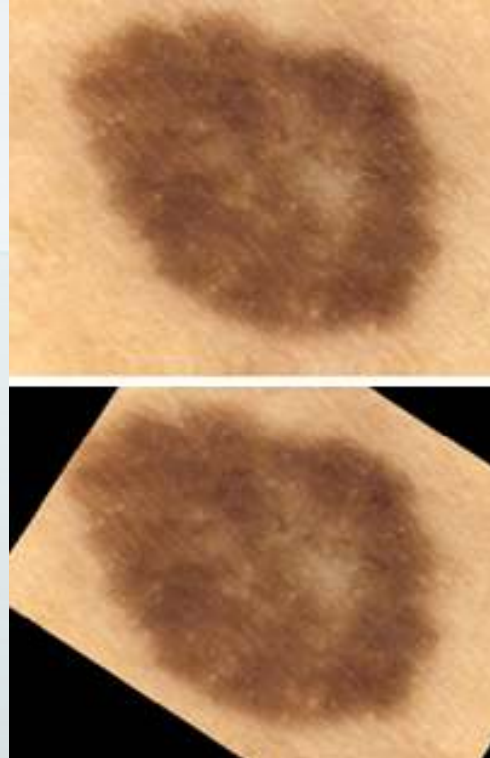
The idea is to simulate what we see in the microscopes in “silico” (Matlab in our case).

Here we use a “FEM” simulator (work in progress)

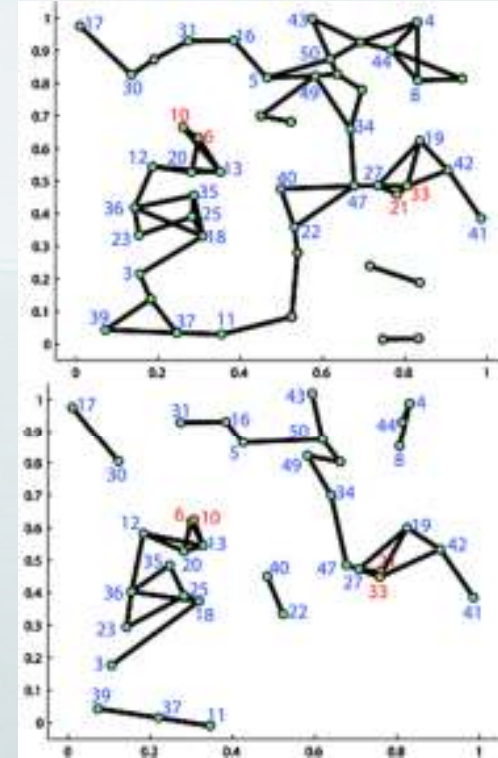
-not mentioned in the slides-



Full Body Scanner project



Robust Registration of dermatoscopic images



Spectral methods for point set matching

Questions?