# **SEGMENTATION OF MEDICAL DATA USING EVOLVING PLANE CURVES**

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(10)

We present mathematical models, computational methods and software solution [6] for computing the plane curve evolution with applications to biological and medical image segmentation. Our solution is based on Lagrangian approach [3, 4, 5]. The plane curve  $\Gamma_t$ ,  $t \ge 0$  is driven by a suitable velocity vector field [1, 2] projected to the normal of the curve. Furthermore, the motion is regularized by the local curvature of the evolving curve and optimized with respect to discrete curve representation by the asymptotically uniform tangential redistribution [4, 5].

#### Velocity vector field





Controlled nontrivial tangential velocity If we consider general curve evolution in normal and tangential directions

> $\partial_t \mathbf{r} = \beta \mathbf{N} + \alpha \mathbf{T}$ . (8)

with

$$\beta = \mu v_{\mathbf{N}} + \epsilon k \qquad \partial_s \alpha = k\beta - \langle k\beta \rangle_{\Gamma} + \left(\frac{L}{g} - 1\right)\omega, \qquad (9)$$

where L is global and g local length of the moving curve,  $\langle k\beta \rangle_{\Gamma}$ is an average of  $k\beta$  along the curve and  $\omega$  is a relaxation parameter, the gridpoints representing numerically the curve are distributed asymtotically uniformly as time evolves [4]. For this model we get intrinsic advection-diffusion equation

 $\partial_t \mathbf{r} = \mu \, v_{\mathbf{N}} \mathbf{N} + \epsilon \, \partial_{ss} \mathbf{r} + \alpha \partial_s \mathbf{r}.$ 

Segmented structures in zebrafish embryogenesis







which is discretized either by explicit (in curve building steps) or



### Curve in velocity vector field

The position vector  $\mathbf{r}$  of evolving curve  $\Gamma_t$  is driven by differential equation (2)

 $\partial_t \mathbf{r} = \mathbf{v} \qquad \mathbf{r}_i^{m+1} = \mathbf{r}_i^m + \tau \mathbf{v}$ 

which is discretized in time and space using gridpoints  $\mathbf{r}_i^m$ , i = 11, ..., N, m denotes time step number and  $\tau$  time step size. The motion starts interactively by user defined uniformly divided line segment which is moved automatically towards the edge.



semi-implicit (in curve post-processing step) flowing finite volume method [3, 5].

Distribution of grid point distances in the final segmentation curve by the various models. The starting uniform discretization of line segment is given by 2 pixels.





## Stopping criterion

The curve is stopped at the m-th time step, if

 $|(v_{\mathbf{N}})_{i}^{m}| < 0.01$  or  $(v_{\mathbf{N}})_{i}^{m} (v_{\mathbf{N}})_{i}^{m-1} < 0, \forall i.$ (11)

## Medical image segmentation







There can be a problem with distribution of curve representing gridpoints due to a discrete character of the vector field.



### Modifications of vector field

 $v_{\mathbf{N}} = \mathbf{v} \cdot \mathbf{N}$  $\partial_t \mathbf{r} = v_{\mathbf{N}} \mathbf{N}$ (3)

The nonlinear term  $v_N$  represents projection of v to the normal N of the moving curve. Removing non-controlled tangential part of velocity is reasonable for uniformly discretized initial curve [6].





(4)

#### Regularization by curvature

 $\partial_t \mathbf{r} = \mu v_{\mathbf{N}} \mathbf{N} + \epsilon k \mathbf{N}.$ 

By the influence of local curvature the motion of grid points is more





#### Segmented evolving structure in embryogenesis









regular, they are tied together in the numerical scheme,  $\mu$  and  $\epsilon$  are parameters weighting advection by vector field and regularization by curvature. Due to Frenet's formula

> $k\mathbf{N} = \partial_s \mathbf{T} = \partial_{ss} \mathbf{r} \qquad \partial_t \mathbf{r} = \mu v_{\mathbf{N}} \mathbf{N} + \epsilon \partial_{ss} \mathbf{r}.$ (5)

where  $\partial_s$  denotes derivative with respect to arclength parametrisation s, we obtain intrinsic diffusion equation (5), which is discretized by the explicit flowing finite volume method [3, 5]

 $\mathbf{r}_{i}^{m+1} = \mathbf{r}_{i}^{m} + \tau \mu \, v_{\mathbf{N}} \, \mathbf{N}_{i}^{m} + \tau \epsilon \frac{2}{h_{i+1}^{m} + h_{i}^{m}} \left( \frac{\mathbf{r}_{i+1}^{m} - \mathbf{r}_{i}^{m}}{h_{i+1}^{m}} - \frac{\mathbf{r}_{i}^{m} - \mathbf{r}_{i-1}^{m}}{h_{i}^{m}} \right)$ (6)  $h_i^m = \sqrt{(x_i^m - x_{i-1}^m)^2 + (y_i^m - y_{i-1}^m)^2}, \quad \mathbf{r}_i^m = (x_i^m, y_i^m).$ (7)



Zebrafish embryogenesis image segmentation

View of **Evo** application with segmented cells



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