

http://www.aimspress.com/

AIMS Medical Science, 2(4): 295–302. DOI: 10.3934/medsci.2015.4.295 Received date 25 May 2015, Accepted date 5 September 2015, Published date 14 September 2015

Research article

Segmentation of Left Ventricle with a Coupled Length Regularization and Sparse Composite Shape Prior: a Variational Approach

Wenyang Liu¹ and Dan Ruan^{1,2,*}

¹ Department of Bioengineering, University of California, Los Angeles, CA 90095, USA

² Department of Radiation Oncology, University of California, Los Angeles, CA 90095, USA

* Correspondence: Email: DRuan@mednet.ucla.edu; Tel: 310-267-8947

Abstract: Segmentation of left ventricles in Cine MR images plays an important role in analyzing cardiac functions. In this study, we propose a variational method that incorporates both prior knowledge on geometrical coupling and shapes of the endo- and epicardium. Specifically, we dynamically maintain and update a smoothly varying distance between the endo- and epicardial contours, represented by a pair of level set functions, with a novel coupling energy embedded in the length regularization. We encode the shape prior with a sparse composite model based on a set of training templates. A robust fidelity with Gaussian mixture models is employed to provide robust intensity estimates in each subregion under insufficient local gradient information. Quantitative evaluation of the proposed method demonstrates competitive/better DSC and APD accuracy compared to other state-of-the-art approaches.

Keywords: left ventricle segmentation; the level set method; variational method; sparse shape prior

1. Introduction

Cine MRI is a suitable modality in analyzing cardiac functions by segmenting the endo- and epicardial contours of the left ventricle (LV) [1]. Given the unclear boundaries between these structures and against the background, prior geometric knowledge needs to be incorporated. A typical idea is to maintain a close-to-constant distance between the endo- and epicardial contours (a "ring" structure) with either hard values [2] or by a soft constraint [3]. However, such static/pre-defined distance may fail to capture large LV shape variations across slices and phases among different subjects. More recently, it has been proposed to use a two-level contours of a single level set function to represent the endo- and epicardium [4], with their enclosing relations maintained by a smoothly varying distance. A local intensity model has been utilized to handle the overlapped distributions. However, representing the endo- and epicardium with a single level set function may hinder flexible

incorporation of shape prior specific to individual structures [5, 6, 7]. The local intensity model could also be overly sensitive to contour initializations and kernel parameter settings [8].

In this study, we propose to represent the endo- and epicardium with two separate level set functions, but introduce a novel coupling setup embedded in the length regularization, with a symmetric "dilation/erosion" morphological relation imposed, to achieve a smoothly varying distance between the endo- and epicardial contours. The coupled level set representations further allows us to incorporate a sparse composite shape descriptor to boost the performance. Gaussian mixture models are utilized to represent overlapped intensity distributions in each subregion. Compared to local intensity models, it is more robust to contour initializations, noise, and selection of setup parameters.

2. Method

2.1. Variational segmentation formulation with a coupled length regularization and sparse composite shape prior

Figure 1 illustrates the proposed cardiac region separation scheme. Specifically, given an image $I: \Omega \to \mathbb{R}$ and two dependent level set functions $\phi_1, \phi_2: \Omega \to \mathbb{R}$, we partition Ω into three subdomains: $\Omega_1 \triangleq \{\mathbf{x} | \phi_1(\mathbf{x}) \le 0\}, \Omega_2 \triangleq \{\mathbf{x} | \phi_1(\mathbf{x}) > 0, \phi_2(\mathbf{x}) \le 0\}, \Omega_3 \triangleq \{\mathbf{x} | \phi_2(\mathbf{x}) > 0\}$, representing the left ventricular cavity, myocardium and background, respectively. A typical variational segmentation formulation consists of two components: $E(\Phi) = E_{fidelity} + E_{reg}$, a data fidelity and a regularization, which we will elaborate bellow.



Figure 1. The proposed cardiac region separation scheme: left ventricular cavity Ω_1 , myocardium Ω_2 , background Ω_3 .

2.1.1. $E_{fidelity}$

We choose to model intensity distributions with Gaussian mixtures for each Ω_i [9]:

$$P(I(\mathbf{x})|\Omega_i, \theta_i) = \sum_{j=1}^{n_i} \frac{w_{j,i}}{\sqrt{2\pi\sigma_{j,i}^2}} \exp(-\frac{(I(\mathbf{x}) - \mu_{j,i})^2}{2\sigma_{j,i}^2}),$$
(1)

where n_i represents the number of Gaussian components for each Ω_i , $\theta_i = \{ w_{j,i}, \mu_{j,i}, \sigma_{j,i} \}$ represents the weights and parameters for the *j*th Gaussian component. $E_{fidelity}$ is then constructed as

AIMS Medical Science

Volume 2, Issue 4, 295–302.

a weighted log-likelihood as:

$$E_{fidelity}(\Phi, \Theta) = -\int_{\Omega} (1 - H(\phi_1)) \log(c_1 P(I(\mathbf{x})|\theta_1)) d\mathbf{x}$$

$$-\int_{\Omega} H(\phi_1) (1 - H(\phi_2)) \log(c_2 P(I(\mathbf{x})|\theta_2)) d\mathbf{x}$$

$$-\int_{\Omega} H(\phi_2) \log(c_3 P(I(\mathbf{x})|\theta_3)) d\mathbf{x},$$
 (2)

where H represents the heaviside function, c_1 , c_2 , c_3 weights the importance/gains of correctly segmenting each corresponding regions.

2.1.2. E_{reg}

Our design of regularization consists of two components: E_{length} and E_{shape} .

 E_{length} : For a smoothly varying distance between the endo- and epicardial contours at each location, we introduce a novel coupling in the length regularization with

$$E_{length} = \int_{\Omega} g_1 \delta(\phi_1) |\nabla \phi_1| d\mathbf{x} + \int_{\Omega} g_2 \delta(\phi_2) |\nabla \phi_2| d\mathbf{x} + \frac{\beta}{2} \int_{\Omega} |\nabla d|^2 d\mathbf{x},$$
(3)

where δ represents the Dirac delta function, $g_1 = (\phi_2(\mathbf{x}) + d(\mathbf{x}))^2$ and $g_2 = (\phi_1(\mathbf{x}) - d(\mathbf{x}))^2$ are the dilated/eroded geometry indicator functions that *traps* ϕ_1 to be the distance *d* from ϕ_2 and vice versa.

 E_{shape} : Inspired by a recently developed shape regularization [7], we construct two shape library $D_{endo} = [\psi_1^{endo}, \psi_2^{endo}, ..., \psi_m^{endo}]$ and $D_{epi} = [\psi_1^{epi}, \psi_2^{epi}, ..., \psi_m^{epi}]$, representing the training shapes. The shape regularization is designed as:

$$E_{shape} = \int_{\Omega} \{ (\phi_1 - D_{endo} \mathbf{w})^2 + (\phi_2 - D_{epi} \mathbf{w})^2 \} d\mathbf{x} + \gamma ||\mathbf{w}||_1,$$
(4)

which dynamically regularizes the current estimates of ϕ_1 , ϕ_2 towards a sparse linear combinations of the training shapes weighted by **w**. In this study, we utilize the single weight **w** to ensure the coupling of training shape selections, since the corresponding column/shape in D_{endo} and D_{epi} comes from the same patient. However, our regularization also permits a more general form that has different library size with different weights (\mathbf{w}_{endo} and \mathbf{w}_{epi}) under the situation when training shapes do not pair up or come from heterogeneous sources.

With the introduced E_{length} and E_{shape} , our E_{reg} is designed as the following:

$$E_{reg}(\Phi, d, \mathbf{w}) = \frac{\lambda_1}{2} \int_{\Omega} (\phi_2 + d)^2 \delta(\phi_1) |\nabla \phi_1| d\mathbf{x} + \frac{\lambda_2}{2} \int_{\Omega} (\phi_1 - d)^2 \delta(\phi_2) |\nabla \phi_2| d\mathbf{x} + \frac{\beta}{2} \int_{\Omega} |\nabla d|^2 d\mathbf{x} + \frac{\alpha}{2} \int_{\Omega} \{(\phi_1 - D_{endo} \mathbf{w})^2 + (\phi_2 - D_{epi} \mathbf{w})^2\} d\mathbf{x} + \gamma ||\mathbf{w}||_1 + \frac{\mu}{2} \sum_{i=1}^2 \int_{\Omega} (\nabla \phi_i - 1)^2 d\mathbf{x},$$
(5)

where $\sum_{i=1}^{2} \int_{\Omega} (\nabla \phi_i - 1)^2 d\mathbf{x}$ is designed to preserve the signed distance property of each level set functions [10]. Smooth approximations of the heaviside and Delta functions are used for numerical

AIMS Medical Science

implementation [11]. Algorithm 1 presents the block coordinate descent/minimization scheme used to minimize the variational energy.

ALGORITHM 1 Minimization scheme

1: while $|E^k - E^{k-1}| < tol$ do minimization w.r.t. $\theta_1, \theta_2, \theta_3$ with Expectation-Maximization (EM) algorithm [12]: 2: $\theta_1^{k+1}, \theta_2^{k+1}, \theta_3^{k+1} \leftarrow EM(\phi_1^k, \phi_2^k, I, \mathbf{w}^k, d^k)$ minimization w.r.t. w with alternating direction method of multipliers (ADMM) [13]: 3: $\mathbf{w}^{k+1} \leftarrow ADMM(\frac{\alpha}{2}(\|\boldsymbol{\phi}_1^k - \boldsymbol{D}_{endo}\mathbf{w}\|_2^2 + \|\boldsymbol{\phi}_2^k - \boldsymbol{D}_{epi}\mathbf{w}\|_2^2) + \gamma \|\mathbf{w}\|_1)$ gradient descending w.r.t. d 4: $d_1^{k+1} \leftarrow d_{iter_num}^k$ **while** *m* < *iter_num* **do** $d_{m+1}^{k+1} = d_m^{k+1} + \Delta t \{ -\lambda_1(\phi_2^k + d_m^{k+1})\delta(\phi_1^k) | \nabla \phi_1^k | - \lambda_2(\phi_1^k - d_m^{k+1})\delta(\phi_2^k) | \nabla \phi_2^k | + \beta \Delta d_m^{k+1} \}$ end while gradient descending w.r.t. ϕ_1, ϕ_2 $\phi_{1,1}^{k+1} \leftarrow \phi_{1,iter_num}^k, \phi_{2,1}^{k+1} \leftarrow \phi_{2,iter_num}^k$ 5: $\phi_{1,i+1}^{k+1} = \phi_{1,i}^{k+1} + \Delta t \{ -\delta(\phi_{1,i}^{k+1}) \log(c_1 P(I(\mathbf{x}) | \theta_1^{k+1})) + \delta(\phi_{1,i}^{k+1}) (1 - H(\phi_{2,i}^{k+1})) \log(c_2 P(I(\mathbf{x}) | \theta_2^{k+1})) + \delta(\phi_{1,i}^{k+1}) (1 - H(\phi_{2,i}^{k+1})) \log(c_2 P(I(\mathbf{x}) | \theta_2^{k+1})) + \delta(\phi_{1,i}^{k+1}) (1 - H(\phi_{2,i}^{k+1})) \log(c_2 P(I(\mathbf{x}) | \theta_2^{k+1})) + \delta(\phi_{1,i}^{k+1}) (1 - H(\phi_{2,i}^{k+1})) \log(c_2 P(I(\mathbf{x}) | \theta_2^{k+1})) + \delta(\phi_{1,i}^{k+1}) (1 - H(\phi_{2,i}^{k+1})) \log(c_2 P(I(\mathbf{x}) | \theta_2^{k+1})) + \delta(\phi_{1,i}^{k+1}) (1 - H(\phi_{2,i}^{k+1})) \log(c_2 P(I(\mathbf{x}) | \theta_2^{k+1})) + \delta(\phi_{1,i}^{k+1}) (1 - H(\phi_{2,i}^{k+1})) \log(c_2 P(I(\mathbf{x}) | \theta_2^{k+1})) + \delta(\phi_{1,i}^{k+1}) (1 - H(\phi_{2,i}^{k+1})) \log(c_2 P(I(\mathbf{x}) | \theta_2^{k+1})) + \delta(\phi_{1,i}^{k+1}) (1 - H(\phi_{2,i}^{k+1})) \log(c_2 P(I(\mathbf{x}) | \theta_2^{k+1})) + \delta(\phi_{1,i}^{k+1}) (1 - H(\phi_{2,i}^{k+1})) \log(c_2 P(I(\mathbf{x}) | \theta_2^{k+1})) + \delta(\phi_{1,i}^{k+1}) (1 - H(\phi_{2,i}^{k+1})) \log(c_2 P(I(\mathbf{x}) | \theta_2^{k+1})) + \delta(\phi_{1,i}^{k+1}) (1 - H(\phi_{2,i}^{k+1})) \log(c_2 P(I(\mathbf{x}) | \theta_2^{k+1}))$ $+\frac{\lambda_1}{2}\delta(\phi_{1,i}^{k+1})\nabla\cdot((\phi_{2,i}^{k+1}+d_{m+1}^{k+1})^2\frac{\nabla\phi_{1,i}^{k+1}}{|\nabla\phi_{\cdot}^{k+1}|})-\lambda_2(\phi_{1,i}^{k+1}-d_{m+1}^{k+1})\delta(\phi_{2,i}^{k+1})|\nabla\phi_{2,i}^{k+1}|$ $-\alpha(\phi_{1,i}^{k+1} - D_{endo}\mathbf{w}^{k+1}) + \mu(\Delta\phi_{1,i}^{k+1} - \nabla \cdot (\frac{\nabla\phi_{1,i}^{k+1}}{|\nabla\phi_{\cdot}^{k+1}|}))\}$ $\phi_{2,i+1}^{k+1} = \phi_{2,i}^{k+1} + \Delta t \{ -\delta(\phi_{2,i}^{k+1}) H(\phi_{1,i+1}^{k+1}) \log(c_2 P(I(\mathbf{x})|\theta_2^{k+1})) + \delta(\phi_{2,i}^{k+1}) \log((c_3 P(I(\mathbf{x})|\theta_3^{k+1})) + \delta(\phi_{2,i}^{k+1}) \log(c_3 P(I(\mathbf{x})|\theta_3^{k+1})) + \delta(\phi_{2,i}$ $-\lambda_{1}(\phi_{2,i}^{k+1}+d_{m+1}^{k+1})\delta(\phi_{1,i+1}^{k+1})|\nabla\phi_{1,i+1}^{k+1}| + \frac{\lambda_{2}}{2}\delta(\phi_{2,i}^{k+1})\nabla\cdot((\phi_{1,i+1}^{k+1}-d_{m+1}^{k+1})^{2}\frac{\nabla\phi_{2,i}^{\kappa+1}}{|\nabla\phi_{2,i}^{\kappa+1}|})$ $-\alpha(\phi_{2,i}^{k+1} - D_{epi}\mathbf{w}^{k+1}) + \mu(\Delta\phi_{2,i}^{k+1} - \nabla \cdot (\frac{\nabla\phi_{2,i}^{k+1}}{|\nabla\phi_{2,i}^{k+1}|}))\}$

end while

6: end while

2.2. Parameter settings

For all experiments in this study, we manually chose and fixed the parameters as: $c_1 = c_2 = 1$, $c_3 = 0.1$, $\lambda_1 = \lambda_2 = 2$, $\beta = 2$, $\alpha = \gamma = 0.02$, $\mu = 1$ and time step $\Delta t = 0.1$. We set the Gaussian component number to be $n_1 = n_2 = 1$ for the cavity and myocardium, and $n_3 = 2$ for the

background. Our algorithm was implemented with matlab 2012b. It took about one minute to segment one image volume on a macbook pro laptop with quad-core i7 2.3GHz and 8GB RAM. No specific code optimization and/or paralellization schemes were used.

3. Experimental results

We evaluated the proposed method on datasets from MICCAI LV segmentation challenge (http: //smial.sri.utoronto.ca/LV_Challenge/Home.html), which consists of 15 training datasets and 15 validation datasets from patients of different pathologies. The Cine steady state free precession MR short axis images were obtained with a 1.5T GE Signa MRI, with FOV = $320 \times 320 \text{ mm}^2$, image size = 256×256 and slice-thickness = 8 mm. The manual segmented endo- and epicardial contours were provided in all slices at end-diastolic (ED) and end-systolic (ES) phases. Our training endo- and epicardium libraries were built upon 15 training datasets, and the evaluation was performed on the 15 validation datasets.

The segmentation accuracy was evaluated based on both Dice similarity coefficient (DSC) metric and average perpendicular distance (APD). DSC is defined as $DSC \triangleq \frac{2|C_{seg} \cap C_{truth}|}{|C_{seg}|+|C_{truth}|}$, where C_{seg} and C_{truth} are the segmented foreground from the obtained and manual segmentation respectively. APD measures the perpendicular distance from the segmented contour to the corresponding manually contour, averaged over all contour points.

Figure 2 compares segmented contours from our method with manual segmented contours on one typical LV example. Figure 3 demonstrates the flexibility and robustness of the proposed method on segmenting one LV example with various thickness of its myocardium within and across different slices. Table 1 reports and compares the DSC and APD statistics of the proposed method with results from other state-of-the-art methods. The proposed method had slightly lower APD compared to [4]. It achieved competitive/better DSC and APD accuracy compared to the other state-of-the-art methods.



Figure 2. Segmentation results on one typical LV example: contours from our method (upper row), manual contours (lower row).



Figure 3. Segmentation results on one LV example with various thickness of the myocardium within and across different slices: contours from our method (upper row), manual contours (lower row).

Table 1.	DSC and	APD	statistics	of seg	nentation	results	on	validation	datasets	from
MICCAI	[challenge									

	DSC				APD (mm)				
	endocardium		epicardium		endocardium		epicardium		
	Mean	S.D.	Mean	S.D.	Mean	S.D.	Mean	S.D.	
proposed	0.89	0.04	0.94	0.02	2.10	0.53	2.09	0.54	
[14]	0.88	0.03	0.93	0.02	2.44	0.62	2.05	0.59	
[4]	0.89	0.04	0.94	0.02	1.93	0.37	1.64	0.42	
[15]	0.89	0.04	0.92	0.02	2.04	0.47	2.35	0.57	

4. Discussion and conclusion

Our proposed coupled length/distance regularization dynamically updates and encourages a smoothly varying distance between the endo- and epicardium through the optimization process. It is more flexible and accurate to capture the large endocardium-epicardium variations across different slices and among different patients, as compared to conventional approaches where such distance is usually static and pre-determined by heuristics [2, 3, 16].

The proposed method is similar to the dual-layer single level set approach [4] in regularizing a smoothly varying distance between the endo- and epicardium, but with added flexibility of incorporating individual priors for each structure separately [5, 6, 7], such as via coupled or decoupled sparse composite shape models depends on the training source. In this study, we utilize a single weight \mathbf{w} to ensure the coupling of epi- and endocardium selections, since those shapes come from a single patient in each shape library. However, under situations that training shapes do not pair up, our formulation would be flexible enough to provide an asymmetric and decoupled regularization. For example, when endo- and epicardial training shapes come from different groups of patients, our

model is capable of providing an asymmetric regularization by enforcing different weights (\mathbf{w}_{endo} and \mathbf{w}_{epi}) from such different training sources to leverage all training instances. Such flexibility also permits further extension of the current model to a joint left and right ventricle (RV) segmentation framework, where training LV and RV shapes often come from heterogeneous sources. In the specific context where differentiating intensity gradient is often absent at the structure interface, the more global Gaussian mixture model we use has the advantage of being more stable, more robust to contour initialization and noise, and does not require the tuning of (heterogeneous) local kernel parameters.

In summary, we have proposed a novel variational segmentation method by introducing geometrical coupling into the length regularization and a flexible coupled/decoupled sparse composite shape prior. Our method has been evaluated on the validation datasets from MICCAI challenge and achieved comparable/better DSC and APD accuracy compared to other state-of-the-art methods. A joint left and right ventricle segmentation method is under development.

Conflict of Interest

All authors declare no conflicts of interest in this paper.

References

- 1. Lelieveldt B, Geest R, Lamb H, et al. (2001) Automated observer-independent acquisition of cardiac short-axis MR images: A pilot study 1. *Radiology* 221: 537–542.
- 2. Lynch M, Ghita O, Whelan PF (2006) Left-ventricle myocardium segmentation using a coupled level-set with a priori knowledge. *Comput Mem Imag Grap* 30: 255–262.
- 3. Kohlberger T, Funka-Lea G, Desh V (2007) Soft level set coupling for LV segmentation in gated perfusion SPECT. In *Medical Image Computing and Computer-Assisted Intervention (MICCAI)* 4791: 327–334.
- 4. Feng C, Li C, Zhao D, et al. (2013) Segmentation of the left ventricle using distance regularized two-layer level set approach. In *Medical Image Computing and Computer-Assisted Intervention* (*MICCAI*) 8149: 477–484.
- 5. Tsai A, Yezzi A, Wells W, et al. (2003) A shape-based approach to the segmentation of medical imagery using level sets. *IEEE Trans Med Imaging* 22: 137–154.
- 6. Cremer D, Osher S, Soatto S (2006) Kernel density estimation and intrinsic alignment for shape priors in level set segmentation. *Int J Comput Vision* 69: 335–351.
- 7. Liu W, Ruan D (2014) Segmentation with a shape dictionary. In *Biomedical Imaging (ISBI)*, *IEEE International Symposium on*: 357–360.
- Wang L, Li C, Sun Q, et al. (2008) Brain MR image segmentation using local and global intensity fitting active contours/surfaces. In *Medical Image Computing and Computer-Assisted Intervention* (*MICCAI*) 5241: 384–392.
- 9. Verma N, Muralidhar GS, Bovik AC, et al. (2011) Model-driven, probabilistic level set based segmentation of magnetic resonance images of the brain. In *Engineering in Medicine and Biology Society (EMBC), IEEE Annual International Conference on:* 2821–2824.

- 10. Li C, Xu C, Gui C, et al. (2005) Level set evolution without re-initialization: a new variational formulation. In *Computer Vision and Pattern Recognition (CVPR), IEEE Computer Society Conference on* 1: 430–436.
- 11. Osher S, Fedkiw R (2003) Level set methods and dynamic implicit surfaces. Springer.
- 12. Dempster AP, Laird NM, Rubin DB (1977) Maximum likelihood from incomplete data via the EM algorithm. *J R Stat Soc Series B (Methodological)* 39: 1–38.
- 13. Boyd S, Parikh N, Chu E, et al. (2011) Distributed optimization and statistical learning via the alternating direction method of multipliers. *Foundations and Trends in Machine Learning* 3: 1–122.
- 14. Jolly MP, Xue H, Grady L, et al. (2009) Combining registration and minimum surfaces for the segmentation of the left ventricle in cardiac Cine MR images. In *Medical Image Computing and Computer-Assisted Intervention (MICCAI)* 5762: 910–918.
- 15. Constantinides C, Chenoune Y, Kachenoura N, et al. (2009) Semi-automated cardiac segmentation on Cine magnetic resonance images using GVF-snake deformable models. *The MIDAS Journal-Cardiac MR Left Ventricle Segmentation Challenge*.
- 16. Folkesson J, Samset E, Kwong R, et al. (2008) Unifying statistical classification and geodesic active regions for segmentation of cardiac MRI. *IEEE Trans Inf Technol Biomed* 12: 328–334.



© 2015, Wenyang Liu, et al., licensee AIMS Press. This is an open access article distributed under the terms of the Creative Commons Attribution License (http://creativecommons.org/licenses/by/4.0)