

Manifold-enhanced Segmentation through Random Walks on Linear Subspace Priors

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Segmentation of skeletal muscles in 3D Magnetic Resonance Imaging, which poses some very specific issues: simultaneous multi-object segmentation, non-discriminative appearance of the muscles, partial contours between them, large inter-subject variations, spurious contours due to fat infiltrations. For these reasons, it is necessary to impose knowledge-based shape priors into segmentation methods. In [2, 3, 4], segmentation is achieved with multi-object elastical deformable models, using either a reference atlas or a hierarchical statistical prior model. A discrete optimization procedure on a graph framework is used in [7], where a higher-order pose invariant shape prior is imposed on surface landmarks nodes. A pixel-wise region-based approach was proposed in [1] where prior knowledge is enforced by embedding the model into a statistical low-dimensional non-linear manifold through PCA in the Isometric Log-Ratio space. Random Walks for image segmentation, presented in [6], is a numerical method for computing globally large discrete regions-based segmentation methods, and is notoriously robust to partial contours. Prior knowledge on intensity within the RW formulation was introduced in [5].

In this paper, we propose a segmentation method based on RW, in which shape deformation is constrained to remain close to a PCA shape space built from training examples. Using the PCA allows us to model complex non-rigid shape variations relying on a few eigen-modes. Our method also benefits from the high performances of the RW optimization process.

The RW method amounts to computing the probability x_i^s that the node v_i is assigned to the label s . It was shown ([6]) that this probabilities minimize the functional:

$$E_{\text{RW}}^s(x^s) = x^{sT} L x^s \quad (1)$$

where L is the combinatorial Laplacian matrix of the graph, defined as:

$$\forall (i, j) \in \mathcal{E}, L_{ii} = \sum_k w_{ik}, L_{ij} = -w_{ij}. \quad (2)$$

It is common practice to use as a Gaussian weighting function for w_{ij} . Once computed x^s for each label s , the segmentation is obtained by retaining the label of maximum probability: $\hat{s}_i = \arg \max_s x_i^s$.

Since minimizing (1) is an independent process for each label s , the whole RW process can be equivalently synthesized in one equation, via concatenation of the x^s and diagonal concatenation of L :

$$E_{\text{RW}}(x) = x^T \bar{L} x. \quad (3)$$

The principle of a shape space is to design a low dimensional affine space approximating this implicit space. Assume we possess T co-registered segmented training volumes. We perform the PCA on vectors $\{\hat{x}^i\}_{i=1\dots T}$, which are ground truth segmentations represented as probability vectors. Retaining the n eigen-modes of greatest variance, the projection of any segmentation in the shape space is represented as:

$$\tilde{x} = \bar{x} + \Gamma \gamma.$$

Thus, any segmentation x can be expressed as:

$$x = dx + \Gamma \gamma + \bar{x} \quad (4)$$

where $dx \in \mathbb{R}^{KN \times 1}$ is the deviation of x from the shape space.

In order to obtain a segmentation which remains close to the shape space, we want to minimize the objective function (3) with respect to

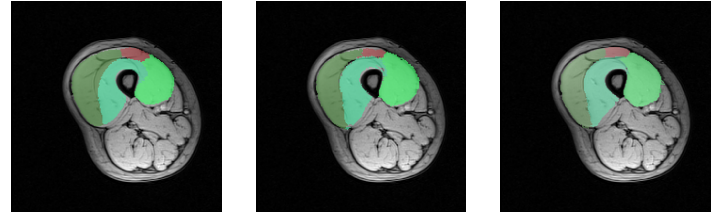


Figure 1: Effect of the PCA shape prior: (left) mean segmentation using $x = \bar{x}$, (middle) shape space segmentation using $x = \Gamma \gamma + \bar{x}$, (right) segmentation with shape prior using $x = dx + \Gamma \gamma + \bar{x}$. The shape space segmentation fits the boundaries better than the mean segmentation, but has fuzzy contours due to the approximation of projecting complex shapes into a linear subspace.

both dx and γ , while keeping dx small. This leads us to the following functional:

$$E_1(dx, \gamma) = (dx + \Gamma \gamma + \bar{x})^T \bar{L} (dx + \Gamma \gamma + \bar{x}) + \lambda \|dx\|^2. \quad (5)$$

The minimum of (5) verifies:

$$(A^T \bar{L} A + \lambda B) y = A^T \bar{L} \bar{x} \quad (6)$$

with:

$$y = \begin{bmatrix} dx \\ \gamma \end{bmatrix}, A = [I_{KN} \ \Gamma], B = \begin{bmatrix} I_{KN} & 0 \\ 0 & 0 \end{bmatrix}. \quad (7)$$

The system of equations (6) can be solved with iterative methods like Bi-Conjugate Gradient. Computation time is approximately 15 min per segmentation on a 2.8 GHz Intel process with 4GB of RAM. We present results obtained with our method on a set of 3D MR volumes of muscles. Our data set is comprised of 30 volumes of the right thigh of healthy subjects, covering a wide range of ages and morphologies. On figure 1, we show the effect of the PCA shape prior on one example of our dataset.

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