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Skeletal muscle segmentation from MRI dataset using a model-based approach

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Magnetic resonance imaging (MRI) and computed tomography scans are used to assess muscle volume, but the manual segmentation, slice by slice, is long and tedious. We proposed an improvement in the deformation of a parametric-specific object method using image processing. The 3D subject-specific geometry was reconstructed based on a few selected number of MRI slices by fast rough contouring using polygons. These polygons were matched to the muscle shape by an optimisation method using an original cost function. Then, parametric-specific object was constructed and deformed. The shape was improved using a loop and the cost function in all MRI slices. The 11 main muscles of the thigh were considered, and the time required to get the shape of all muscles was 21 min, with a volume error inferior to 5% and a point-surface distance error (2RMS) inferior to 5 mm. This method provides a good compromise between segmentation time and an accurate representation of the muscles shape.

Keywords: MRI; tomodensitometry; muscle; segmentation; volume

1. Introduction

Muscle volume decrease is an important factor related to the muscle weakness observed with age (Narici et al. 2003), neuromuscular pathology (Malaiya et al. 2007) or injury (Williams et al. 2005). *In vivo* muscle volume estimation is then of great interest for clinical patient follow-up as for biomechanical analysis, which requires patient-specific muscle geometry (Arnold et al. 2000; Blemker et al. 2007). However, muscle volume estimation remains a great challenge when seeking for accuracy, reproducibility and speed.

Muscle volume has been evaluated using magnetic resonance imaging (MRI) axial slices (Tracy et al. 1999, 2003; Fukunaga et al. 2001; Lund et al. 2002; Eng et al. 2007; Morse et al. 2007; Sudhoff et al. 2009) or computed tomography (CT)-scan axial slices (Rice et al. 1989; Overend et al. 1992; Jolivet et al. 2008). Muscle 3D reconstruction was assessed by manual contouring of many images in which muscle of interest was present (Tracy et al. 1999; Fukunaga et al. 2001; Eng et al. 2007), but the process is long and tedious, and such studies are only conducted on a limited number of subjects. Muscle volume was also estimated using only a given number of cross-sectional muscle area (Lund et al. 2002; Tracy et al. 2003; Morse et al. 2007), but in such cases, the muscle shape is not accessible. Very few studies use automatic muscle segmentation because of the noise included in the images. To our knowledge, only two studies focus on muscle segmentation on MRI using model-based approach

applied for the first one on masseter muscles and for the second one (Gilles and Magnenat-Thalmann 2010) on the lower limb muscles. The first method implies the construction of a first coarse segmentation based on a priori knowledge of the specific muscle geometry followed by an optimisation of the muscle boundary, whereas the second method uses a scalable simplex surface equipped with reversible medial representations which could be less adapted for pathological patient.

Our group recently proposed a method, based on the deformation of a parametric-specific object (DPSO), to assess muscle geometry using a reduced number of axial images (Jolivet et al. 2008): a manual contouring is performed on this reduced set of images, then a parametric shape-based interpolation combined with a kriging technique allows to get a surface model without using the intermediate CT slices and without any a priori knowledge related to the muscle shape. The method has been validated (Jolivet et al. 2008), and it has been shown that the accuracy of muscle volume estimation using DPSO method (i.e. a function of the reduced number of slices used for muscles) converges very quickly (Sudhoff et al. 2009). Although this method provides a drastic improvement in reconstruction time, it still requires a thorough digitisation of the reduced set of slices. Moreover, from the obtained solution, image-processing techniques could allow automatic improvement by taking into account the intermediate images.

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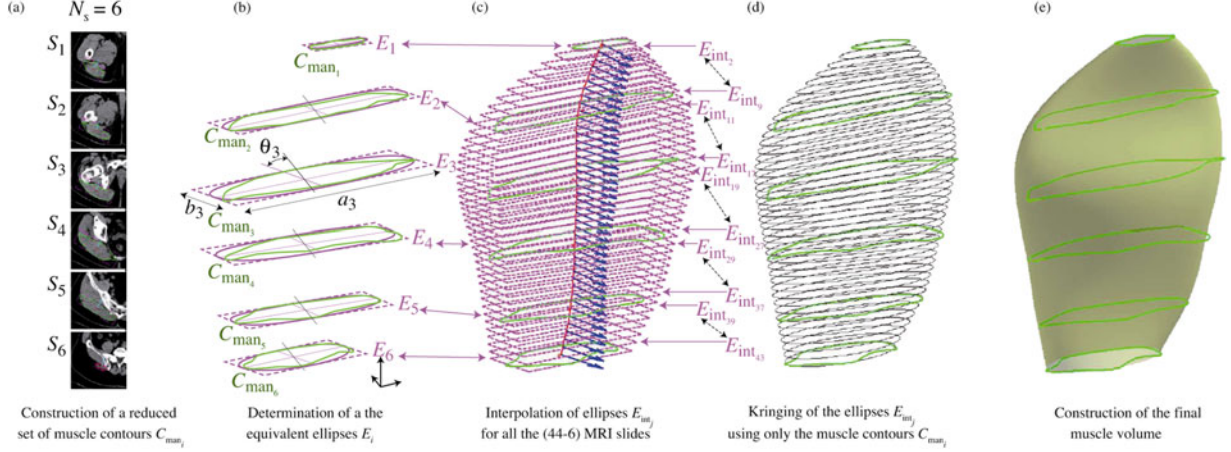


Figure 1. DPSO method presented for a reduced set of six slices.

The objective of this study was to propose and evaluate an improvement in the DPSO muscle segmentation method based on specific image analysis.

2. Materials and methods

The proposed method is based on the DPSO method, which is briefly described hereafter and illustrated in Figure 1.

First, from all the slices associated with a dedicated muscle, a reduced number N_s of slices are considered (Figure 1(a)): two at the extremities (S_1 and S_{N_s}) and the others at regularly spaced levels.

On each slice S_i of the reduced set (i is the slice number), a fine manual contouring C_{man_i} is performed. Then, a best-fit ellipse E_i is associated with the contour, described by three parameters: length a_i , width b_i and orientation θ_i with regard to a global coordinate system (Figure 1(b)). A set of n equidistant points is defined both on the contour and on the ellipse, respectively, named PC_{ij} and PE_{ij} (j is the point number).

For each intermediate slice (k is the intermediate slice number), which is not contoured, an intermediate ellipse E_{int_k} is defined using cubic spline interpolation of the descriptive parameters (Figure 1(c)). n equally distributed points PE_{kj} are also considered.

The whole set of points on the ellipses (PE_{ij} and PE_{kj}) describe a simplified parametric model of the muscle (Figure 1(c)). The points on the contours of the reduced set (PC_{ij}) are used as control points to deform this simplified model using a kringing technique (Trochu 1993), yielding the solution for muscle parametric subject-specific model (Figure 1(d)) allowing for muscle volume computation (Figure 1(e)). In this study, image processing was used to improve the process at two levels:

- Semi-automatic contouring of the reduced set of images, with a fast manual contouring followed by automatic adjustment,

- Automatic adjustment of the intermediate contours resulting from the kringing technique in order to take advantage of the information on the image and to increase the accuracy.

In both cases, automatic adjustment of an initial contour is done using an optimisation method with a specific cost criterion that is detailed hereafter.

2.1 Simplified contour generation and its optimisation

In this new approach, the fine manual contouring C_{man_i} that was performed on the reduced set of slices is substituted by a rough polygon C_{poly_i} (Figure 2(a)) in order to gain segmentation time. Then, an image-based optimisation process is carried out to improve this polygonal contour automatically, leading to a robust optimised contour C_{opt_i} (Figure 2(c)).

For that, p points $PC_{poly_{ij}}$ are equally distributed on the polygon contour as presented in Figure 2(b) (j is the number of the point on the contour). A cost function G is then used to identify the optimal position X_{ijk} of these points along the local normal of the polygonal contour in their close neighbourhood (k is the number of the point along the normal). $k = 2/3 \times p$ was arbitrarily chosen and equally distributed, respectively, along \vec{n} and $-\vec{n}$ directions.

This cost function G is the product of three terms: G_{Grad} (image gradient), G_{Int} (pixel intensity) and G_{Dist} (distance to the initial contour). They are defined as follows:

- Gradient cost function $G_{\text{Grad}}(X_{ijk})$: a Sobel filter in the two orthogonal directions in the image yielded the gradient value $\text{Grad}(X_{ijk})$ for each point X_{ijk} . Then, a gradient cost was computed using a Laplacian–Gaussian function (Sobel and Feldman 1973):

$$G_{\text{Grad}}(X_{ijk}) = \frac{1}{\sigma\sqrt{2\pi}} e^{-\frac{1}{2}(\text{Grad}(X_{ijk}) - \mu)/\sigma} \quad (1)$$

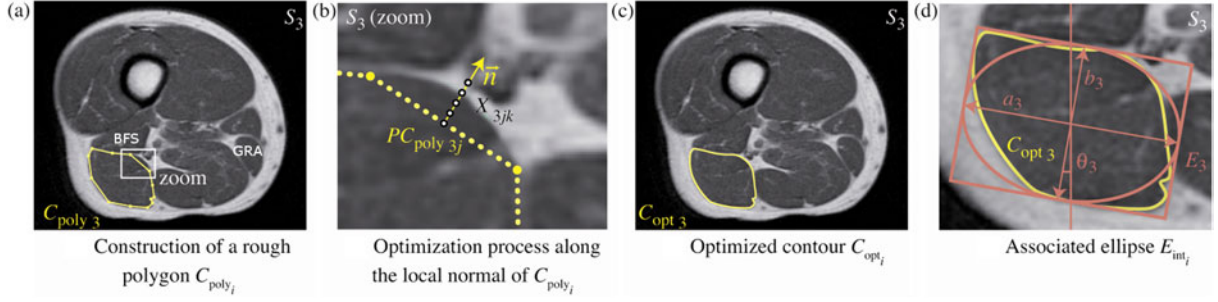


Figure 2. Optimisation process of the rough contouring.

with $\mu = \max_k(\text{Grad}(X_{ijk}))$ and $\sigma = \mu/2$

This function allowed to detect the variation in pixel intensity. Specifically in the example of Figure 2, the function was used to detect the boundaries between the muscular tissue (in black) and the fat tissue (in white). The Gaussian function grants a higher importance to the high value of the gradient.

- Pixel intensity cost function $G_{\text{Int}}(X_{ijk})$: R_{poly} and R_X were regions defined by the contour C_{poly_i} and the modified contour C'_{poly_i} considering the location of the point X_{ijk} . $P(\cdot)$ denotes the number of pixel in a region, $\text{mean}(\cdot)$ the mean intensity of pixel in a region and $\text{var}(\cdot)$ the variance of pixel intensity:

If $P(R_X) \geq P(R_{\text{poly}})$:

$$G_{\text{Int}}(X_{ijk}) = 1 - \frac{|\text{mean}(R_X - R_{\text{poly}}) - \text{mean}(R_{\text{poly}})|}{2\sqrt{\text{var}(R_{\text{poly}})}}. \quad (2)$$

The added region should have the same pixel intensity as the initial region. The more the $G_{\text{Int}}(X_{ijk})$ is close to 1, the more similar the pixel intensities are.

If $P(R_X) < P(R_{\text{poly}})$:

$$G_{\text{Int}}(X_{ijk}) = \frac{|\text{mean}(R_{\text{poly}} - R_X) - \text{mean}(R_{\text{poly}})|}{2\sqrt{\text{var}(R_{\text{poly}})}}. \quad (3)$$

The removed region should have a different pixel intensity from the initial area. In this case, the larger the $G_{\text{Int}}(X_{ijk})$ is close to 1, the more different the pixel intensities are.

This function allowed to choose the point among those which present a high value of G_{Grad} and to keep a global coherence of the pixel intensity if no points are highlighted with the G_{Grad} function.

- Distance cost function $G_{\text{Dist}}(X_{ijk})$:

$$G_{\text{Dist}}(X_{ijk}) = \frac{\max_p(\|X_{ijp} - PC_{\text{poly}_{ij}}\|) - \|X_{ijk} - PC_{\text{poly}_{ij}}\|}{\max_p(\|X_{ijp} - PC_{\text{poly}_{ij}}\|)}. \quad (4)$$

This function allowed to limit the deviation of the contour if the boundaries between two muscles were not distinct. For example, in Figure 2, the biceps femoris long head muscle (segmented muscle) and the semitendinosus muscle (on the right of the segmented muscle) were difficult to distinguish.

- Cost function definition G :

$$G(X_{ijk}) = G_{\text{Int}}(X_{ijk}) \cdot G_{\text{Int}}(X_{ijk}) \cdot G_{\text{Dist}}(X_{ijk}). \quad (5)$$

- All the positions of p points of the contour C_{poly_i} are optimised at the minimum of the cost function $G(X_{ijk})$, and C_{poly_i} became C_{opt_i} (Figure 2(c)):

$$C_{\text{opt}_i} = \{\max_{j,k} G(X_{ijk})\} \quad (6)$$

2.1.1 Interpolation process

The two steps of the interpolation process are derived from the classical DPSO method, briefly described in the following.

- (1) For each optimised contour C_{opt_i} , an equivalent ellipse E_i is determined (Jolivet et al. 2008). This ellipse is described by its centre, axis orientations, width and length parameters as previously described (Figure 2(d)). Then, intermediate ellipses E_{int_i} are defined on each slice that is not incorporated into the reduced set of slices. A cubic spline interpolation is used to evaluate the values of parameters for the location of these slices. On all the ellipses E_i and E_{int_i} , p points are equally distributed in the 3D coordinate system related to the image acquisition device. Thus, a 3D surface object is defined, based on these 3D points of the whole set of ellipses.
- (2) The second step consisted of a nonlinear deformation on this surface, using kriging algorithm (Trochu 1993; Jolivet et al. 2008). Points that described E_i are used as control points to globally deform the 3D surface object so that each of those points fits exactly description points of the C_{opt_i} contours. At the end of

this interpolation process, the first estimate of muscle contour C_{int_k} is obtained for each intermediate slice. These contours are progressively used as an initial solution for the muscle segmentation of intermediate slices using an iterative process described in the following paragraph.

2.1.2 Iterative process

The iterative process is illustrated in Figure 3. The intermediate contours C_{int_k} that are adjacent to C_{opt_i} contours are denoted C_{adj_k} . They are optimised using the method described previously and become muscle optimised contours denoted C_{opt,adj_k} . Therefore, the reduced set of optimised contours includes both the previous C_{opt_i} and the new C_{opt,adj_k} contours. The interpolation process is then applied for slices in which contours have not yet been optimised. This iterative procedure is applied until an optimised muscle contour is defined for each slice between S_1 and S_{N_s} .

2.2 Muscle segmentation evaluation

MRI acquisitions of the hip and the thigh were acquired for four subjects (Table 1).

Table 1. Population characteristics: age (years), height (cm) and weight (kg).

	Sex	Age (years)	Weight (kg)	Height (cm)
Subject 1	F	27	53	162
Subject 2	F	32	81	181
Subject 3	F	27	82	182
Subject 4	F	26	56	174
Mean		28	68	175
STD		2.7	15.6	9.2

The local ethical committee granted its approval for this study (CPP Ile de France VI, Hôpital Pitié- Salpêtrière, Paris, France, 119-08). A Philips Panorama HFO 1.5 T MRI was used to obtain a series of axial images from the superior iliac crest to the ankles in decubitus position of the subject with feet taped together. Acquisitions were realised with spin echo T1 sequence (TR 680 ms and TE 50 ms) and the Q-body coil. Slice thickness was 10 mm with a pixel resolution of $0.78 \times 0.78 \text{ mm}^2$.

Muscles of the thigh were considered. The anterior muscle group was composed of rectus femoris (RF), vastus lateralis, vastus medialis and vastus intermedialis that were grouped together in the vastus muscle group (VMG), sartorius (S) and tensor fasciae latae muscle (TF). Biceps longus (BL), biceps brevis (BB), semitendinous (ST) and semimembranous (SM) of the posterior muscle group were also reconstructed for all subjects as gracilis muscle (G).

First, to evaluate the segmentation error induced by the automatic contour optimisation of the rough polygon, the point-to-curve distance between the points of the proposed method and the reference contour was computed. For seven slices, 50 points per slice, per muscles were projected along the normal of the reference contour, and the distance between the pair of points was calculated. The 2RMS point-to-curve distance error was studied for each muscle.

Second, muscles included in this study were manually outlined in all images of acquisition. For each muscle, using all contours, a 3D surface object was generated, named M_{Ref} , and reference muscle volume ($V_{m,Ref}$) was computed for each muscle.

Finally, to analyse the sensitivity of the muscle segmentation to the size of the reduced set of MRI images N_s , the convergence of the muscle volume V_{m,N_s} and of the 3D muscle shape M_{N_s} was evaluated with N_s varied from 2 to 7, as seven slices are sufficient to obtain convergence (Sudhoff et al. 2009). To estimate shape differences between M_{Ref} and M_{N_s} , point-to-surface distance was determined between each point of M_{N_s} and surface of M_{Ref} . Mean distance and 2 root-mean-square (2RMS) that corresponded to 95% confidence interval were computed for each M_{N_s} .

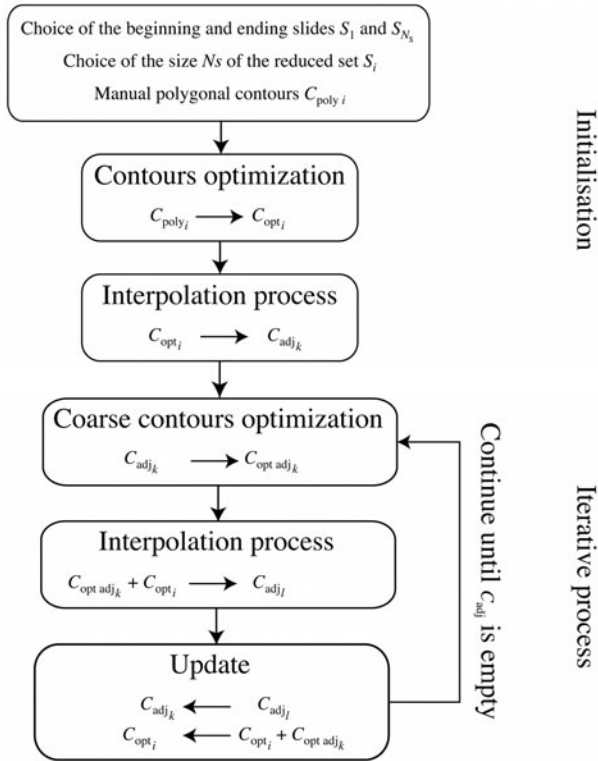


Figure 3. Illustration of muscle segmentation procedure. C_{poly} , rough manual polygon; C_{opt_i} , optimised contour; C_{adj} , adjacent to an optimised contour (before optimisation); and $C_{opt,adj}$, adjacent to an optimised contour (after optimisation).

Table 2. Number of slices for each of the nine visible muscles included in the study (mean, maximum and minimum) and reference muscle volume computed from the segmentation generated from muscle contours (mean, maximum and minimum).

Muscle	Abbreviation	Number of slices Mean (min–max)	Reference Volume (cm ³) Mean (min–max)
Biceps femoris short head	BFS	17 (14–20)	91 (66–121)
Biceps femoris long head	BFL	22 (17–27)	159 (112–203)
Rectus femoris	RF	25 (22–26)	236 (193–288)
Gracilis	GRA	26 (24–29)	92 (57–121)
Sartorius	SAR	40 (38–44)	123 (91–157)
Semimembranosus	SM	13 (14–19)	165 (97–236)
Semitendinosus	ST	24 (15–28)	160 (104–225)
Tensor fasciae latae	TFL	12 (10–13)	59 (40–83)
Vastus intermedialis	VMG	24 (21–28)	1394 (1048–1774)
Vastus lateralis			
Vastus medialis			

3. Results

The mean number of image on which each muscle was visible is indicated in Table 2. The global 2RMS point-to-curve distance error for the contour segmentation was 1.29 mm, with a maximum of 2.26 mm for the BFS muscle and a minimum for GRA muscle with 0.63 mm (Table 3).

The volume difference (Figure 4) decreases when the number of manual contours N_s increases. If we consider a difference lower than 5%, six manual muscle contours would be required for ST muscle, five for BFS, RF and SAR and four for BFL, GRA, SM, TFL and VMG, instead of seven for RF and SAR, six for BFL, five for BFS, GRA, SM, TFL and ST and four for VMG with the non-optimised method.

Point-to-surface distance between surface M_{N_s} and surface of M_{Ref} (Figure 5) decreases when the number of manual contours N_s increases. If we consider a difference lower than 5 mm, five manual muscle contours would be required for RF, SM, ST and VMG muscles, four for BFS, BFL, SAR and TFL and three for GRA.

4. Discussion

This study proposed a method to obtain a fast and accurate segmentation of muscle and was applied on the muscles of the thigh. This method required manual muscle contours on a reduced number of images, and the segmentation process was realised in two steps: the first coarse segmentation followed by an optimisation iterative procedure. With respect to the already published techniques, this new development regards the automatic contour optimisation (ACO) based on a cost function containing an original coherence criterion. ACO allowed

time reduction in the manual processing of the reduced set of contours. It also allowed automatic adjustment of the intermediate slices that were previously unprocessed using DPSO method. The cost function appeared efficient because the DPSO method provided an initial solution, which was already close to the adjusted one. Table 4 summarises the improvements in the proposed method.

The sensitivity analysis of the number of manual muscle contours showed that the volume difference between the segmented muscle and the reference muscle volume, obtained by manually contouring all the images in which muscle of interest was present, lowered to 5% with only 4–6 manual muscle contours, depending on the muscle. While the previous DPSO method needed a reduced set of seven slices and 26 min to build the whole model, this new approach needs only five slices and 21 min, which is of great interest regarding the classical manual segmentation time that was up to 80 min. A significant part of the enhancement is due to the iterative interpolation process that is very efficient on long-curved path muscle such as the SAR.

Some limitations had to be dealt with. A difficulty was to identify slices where considered muscles appeared or disappeared because the muscle–tendon transition is not really clear. A previous study has shown that this uncertainty of these extreme slice locations leads to an error estimation lower than 2.1%, depending on the muscle (Sudhoff et al. 2009). However, this requires well-trained operators, and further improvements are necessary to reduce operator dependence, particularly in longitudinal studies in which the muscles change and the operators are not necessarily the same.

Table 3. Point-to-curve distance error between the points of the proposed method and the reference contour (7 slices, 50 points per slice).

Muscle	BFL	BFS	GRA	RF	SAR	ST	SM	TFL	VMG	Global
Mean	0.65	0.88	0.42	0.75	0.62	0.59	0.69	0.56	0.73	0.70
2*RMS	1.08	2.26	0.63	1.24	1.35	0.88	1.09	0.98	1.06	1.29

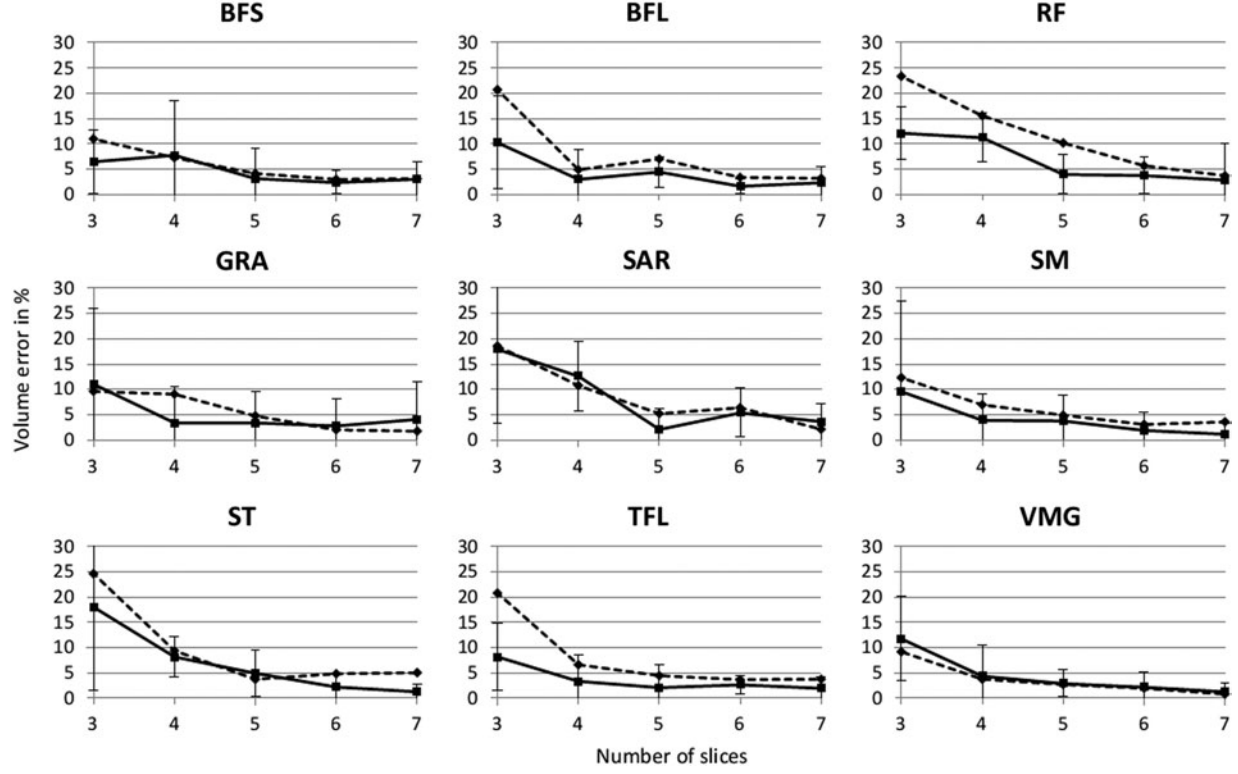


Figure 4. Volume difference in per cent between reference volume $V_{m,Ref}$ and segmented volume V_{m,N_s} for each muscle included in the study (dotted line represents the DPSO method Jolivet et al. (2008) and continuous line represents the presented method).

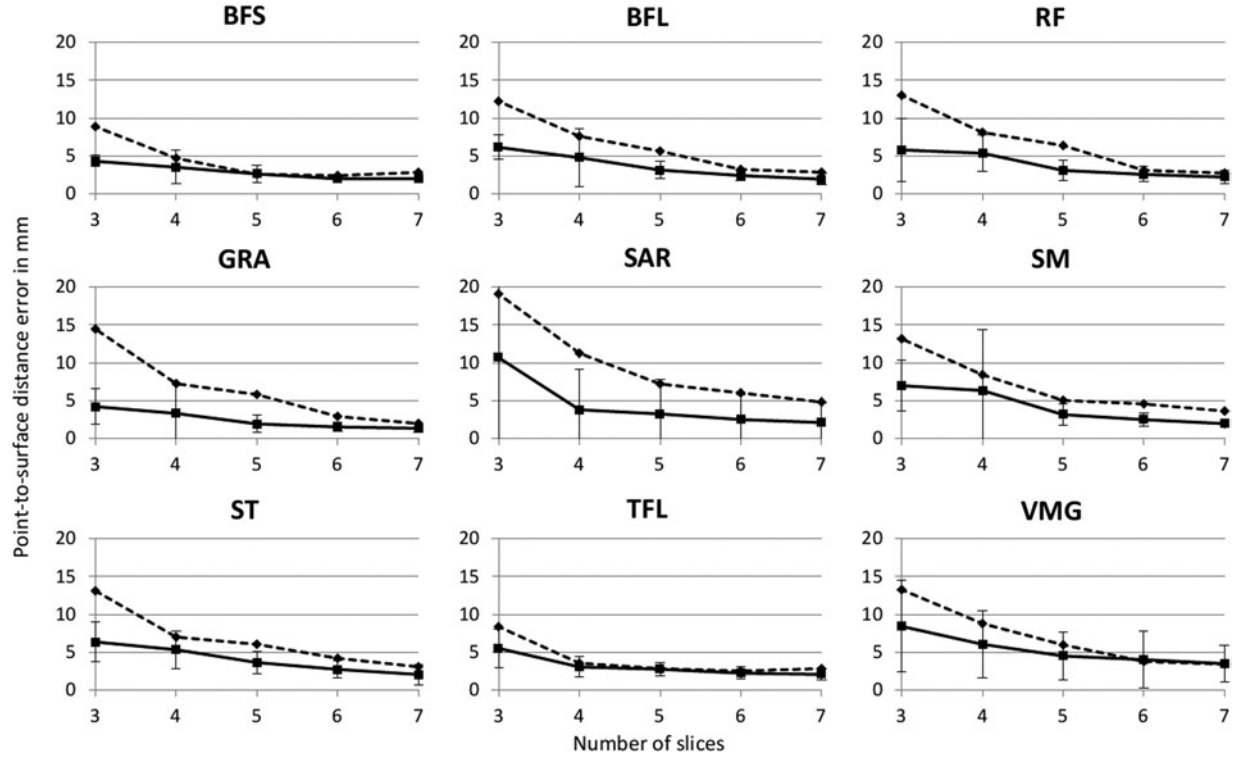


Figure 5. Point-to-surface distance between surface M_{N_s} and surface of M_{Ref} for each muscle included in the study (dotted line represents the DPSO method Jolivet et al. (2008) and continuous line represents the presented method).

Table 4. Main differences between the DPSO method and the presented method.

	DPSO method	Proposed method
Automatic image processing	No	Automatic contour optimisation (ACO) using an original cost function
Muscle contour segmentation in a reduced set of slices	Manual contouring	Rough polygon + ACO
Automatic estimation of intermediate muscle contours	Yes (DPSO)	Yes (DPSO)
Automatic adjustment of the estimated intermediate muscle contours	No	Yes (ACO)

As for various medical images, segmentation is an uneasy task because of the noise included in the images. Especially the boundary of two muscles (e.g. whole vastus or BFS and BFS) was not easy to identify. The cost function $G_{\text{Dist}}(X_{ijk})$ limited the aberrant solution, especially during the *iterative process*. A manual correction might be necessary to improve the estimated shape. For this study, no manual corrections were made.

The location of the slices S_i included in the reduced set could affect the definition of the first coarse segmentation; consequently, refinement step would also be affected. In this study, these slices were equally spaced. The cost function that was used for the optimisation process combined the classical gradient cost function with two functions related to coherence criteria: a distance criterion, based on the fact that the initial contour is already close to the real one, and a pixel intensity criterion, based on the fact that the global signal intensity for the muscle is clearly differentiated from the surrounding. The distance criterion can be less relevant for muscles that undergo wide and sudden change in shape and cross-sectional area, because in this case the initial contour for the optimisation process remains quite different from the real contour, and the combined criteria for segmentation did not fully succeed in retrieving this solution: particularly, the 10-mm slice thickness used in this study, corresponding to routine clinical protocol, is a limitation, while thinner slices would probably allow to take progressively into account this shape variation. For such routine protocols, it could be interesting to use training data-set to determine the optimal location of the slices for manual contouring (Ng et al. 2009). Nevertheless, in general, the iterative process proposed in this study with optimisation of muscle contour gradually ensured to optimise an estimated muscle contour that was close to muscle contour solution that would be obtained with full manual reconstruction.

Some muscles were inconstantly distinguishable because the fascia between those two muscles is not visible in most of the cases. This was the case of the vastus medialis muscle and the vastus lateralis muscle, and their volume was evaluated together (Sudhoff et al. 2009).

Another limitation was the difficulty in evaluating the reference muscle volume *in vivo*. In many studies as in this study, the reference standard measure of muscle volume

was the volume computed using manual muscle contour in all images in which muscle was present. Thus, accuracy of this reference volume estimation was not clear. Manual segmentation of *in vitro* MRI acquisition of forearm followed by dissection and volume measurement demonstrated that volume estimation accuracy was related to area-to-volume ratios and related error was $> 10\%$ for high value of this ratio (Eng et al. 2007). Moreover, inter-operator reproducibility study, including two operators, focused on muscles crossing the knee and showed that volume error estimation between the operators with entire manual procedure on MRI acquisition of the lower limb was between 2.7% and 13.9% depending on the muscles (Sudhoff et al. 2009).

The image processing with the use of the cost function allowed an easier way to segment the muscle contours. However, this method induced a global 2RMS point-to-curve distance error of 1.29 mm. This error was due to the difficulty in distinguishing the boundaries between the muscles, especially for the BFS muscle, which had an error of 2.26 mm. GRA appeared as the easier muscle to segment, as it is along the fat tissue and has a thick fascia and had an error of 0.63 mm. Those muscles are visible in Figure 2(a).

In conclusion, the method of muscle segmentation presented allowed estimating the muscle volume *in vivo* by minimising human intervention from MRI acquisition. Using the results of sensitivity analysis, clinicians could define the number of manual contours to perform muscle segmentation for a given expected precision on volume estimation.

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