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## Online atlasing using an iterative centroid \*

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Abstract. Online atlasing, i.e. incrementing an atlas with new images as they are acquired, is key when performing studies on databases very large or still being gathered. We propose to this end a new diffeomorphic online atlasing method without having to perform again the atlasing process from scratch. New subjects are integrated following an iterative procedure gradually shifting the centroid of the images to its final position, making it computationally cheap to update regularly an atlas as new images are acquired (only needing a number of registrations equal to the number of new subjects). We evaluate this iterative centroid approach through the analysis of the sharpness and variance of the resulting atlases, and the transformations of images, comparing their deviations from a conventional method. We demonstrate that the transformations divergence between the two approaches is small and stable and that both atlases reach equivalent levels of image quality.

## 1 Introduction

An atlas is a generic term covering several uses. Overall, it corresponds to an average model of the brain both in terms of shape and intensity. It is therefore a powerful tool to understand brain variability, to compute statistics on populations [14] or to segment regions [3]. Ideally, atlases are made from a large number of subjects by combining their images through registration and transformation composition which can be computationally quite costly.

Among the many available atlas construction methods, Guimond et al. [8] proposed to register the subjects onto an arbitrary image, followed by an unbiasing and averaging step, leading to an atlas that becomes the new reference image. By iterating the process, the atlas becomes less and less biased by the initial reference image. Joshi et al. and Rohlfing et al. [9,12] among others focused on another atlas construction class, called template free approaches, working with groupwise registration. Although those methods have the advantage of being unbiased by construction, their computational cost can be prohibitive.

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Historically databases used for atlas construction were small (about 50 to 100 subjects at most). However, there is a recent trend for the creation of large databases, out of which great atlases depicting the variability of the population across the life span could be inferred. Among these databases, the human connectome project (HCP, now completed) is made of around 1200 subjects of the same age range [7]. Even bigger databases are currently being constituted from more subjects (up to 100,000 in the UK biobank [11]) or across the life span (HCP lifespan) on which researchers already wish to start studies. This causes a problem as the databases are still being gathered and thus the atlas needs to be updated. With current methods, the atlas needs to be recomputed each time new acquisitions are added which is very costly. New methods are therefore needed to allow online atlasing, i.e. the capacity of incrementing atlases with new images as they come without having to perform again the whole atlas construction, as opposed to the above-mentioned direct atlasing.

We propose a new method that considers atlasing from a different angle. Inspired by the notion of centroid for a set of points in a Euclidean space and the way it can be constructed following an iterative procedure, Cury et al. [6] derived a diffeomorphic atlas construction method working on surfaces. We propose here a generalization of this iterative centroid method for image atlasing where the centroid (atlas) is updated gradually as new images are incorporated. The online aspect of this technique has several advantages: when a new images arrives, its integration does not necessitate to restart the atlasing process from scratch. Also, it only requires one registration per new image making it computationally cheap. Benefitting from the log-Euclidean framework for diffeomorphisms from Arsigny et al. [1], the method allows the production of a diffeomorphic atlas unbiased up to a linear transformation (rigid or affine at the choosing of the user).

We first introduce the iterative centroid method and its specificities for image atlasing in Section 2. Then in Section 3, we present experiments comparing how the atlases built using direct atlasing (here a modified Guimond et al. method [10]) and with our method differ from each other as they include more and more subjects. We demonstrate with three different metrics that the iterative centroid and direct atlases diverge only by a small amount in terms of transformations and that their intensity based features do not vary.

## 2 Method

## 2.1 Theoretical background

Considering the registration of an image J onto an image I, we denote by  $\tilde{J}$  the registered image and T the transformation operating on coordinates x used to resample J onto I:  $I(x) \sim \tilde{J}(x) = J \circ T(x)$ . Assuming that T lives in a connected Riemannian manifold, we consider  $\gamma(t)$  the geodesic between Id and T such that  $\gamma(0) = Id$  and  $\gamma(1) = T$ . In this framework (named large deformation diffeomorphic metric mapping - LDDMM), the deformation T is parameterized by a time varying vector field integrated over time (the geodesic  $\gamma$ ).

Diffeomorphisms are widely used in non linear registration because of their interesting properties: differentiability, bijectivity and differentiability of the inverse map. As the LDDMM is a computationally intensive framework, Arsigny et al. [1] presented a log-Euclidean framework to compute statistics on diffeomorphisms parameterized by a time independent i.e. stationary velocity field (SVF). For such diffeomorphisms, operations are defined as their Euclidean counterparts on the SVFs. One of interest is the power of a diffeomorphism defined as  $T^{\alpha} = \gamma(\alpha) = \exp{(\alpha \log(T))}$ . In the following, all non linear transformations are assumed to be diffeomorphisms parameterized with SVFs.

Additionally, given 2 SVFs v and w, it is possible to compute the logarithm of their composition while avoiding a very costly logarithm computation. This is made possible by the Baker-Campbell-Hausdorff (BCH) formula stating that in a BCH-Lie group, for v and w small enough,  $\log(\exp(v) \circ \exp(w))$  can be approximated as a series of Lie brackets:

$$\log(\exp(v) \circ \exp(w)) \approx \mathrm{BCH}(v, w) = v + w + \frac{1}{2}[v, w] + \dots \tag{1}$$

Where [v, w](x) = Jac(v)(x).w(x) - Jac(w)(x).v(x). It has been shown in [2,13] that the use of the BCH formula is well suited for diffeomorphisms parameterized as SVFs. To simplify notations, unless specified otherwise, the composition of two transformations associated to two SVFs v and w is performed via the BCH:  $\exp(v) \circ \exp(w) = \exp(\text{BCH}(v, w))$ .

#### 2.2 Iterative centroid atlas construction

In a Euclidean space  $\Omega$ , the centroid  $b_n$  of a set of points  $\{x_1,\ldots,x_n\}$  is defined as  $b_n = \operatorname{argmin}_{y \in \Omega} \sum_{i=1}^n \|y - x_i\|^2$ . A direct solution is  $\frac{1}{n} \sum_{i=1}^n x_i$  but an iterative approach is also imaginable. Starting from  $b_1 = x_1$ , we can use the recursion:

$$b_{k+1} = \frac{k}{k+1}b_k + \frac{1}{k+1}x_{k+1} \tag{2}$$

As depicted in [6], one can extend this formulation to the Riemannian case for surfaces by using the generalization of the notion of straight line through geodesics. The updated centroid  $b_{k+1}$ , in that case, is on the geodesic linking  $b_k$  and  $x_{k+1}$ , at  $\frac{1}{k+1}$  times the distance from  $b_k$ .

We define our iterative centroid atlasing, following the same spirit, on images and diffeomorphisms parameterized by SVFs. Thereafter we assume that the registration of an image I onto an atlas A provides two transformations L and T, where L is linear and T is a diffeomorphism, so that  $A \sim I \circ L \circ T$ . Generalizing Eq. 2 to a set of images  $\{I_1, \ldots, I_n\}$ , we start from  $A_1 = I_1$  and get:

$$A_{k+1} = \frac{k}{k+1} A_k \circ T_{k+1}^{-\frac{1}{k+1}} + \frac{1}{k+1} I_{k+1} \circ L_{k+1} \circ T_{k+1}^{\frac{k}{k+1}}$$
 (3)

The atlasing scheme is illustrated Fig. 1. We apply to  $A_k$  a transformation situated at  $\frac{1}{k+1}$  of the distance from the identity to  $T_{k+1}^{-1}$  along the geodesic. We

also apply to  $I_{k+1}$ , in addition to the linear transformation, a transformation situated at  $\frac{k}{k+1}$  of the distance from the identity to  $T_{k+1}$  along the geodesic. Also,  $A_k$  is made from k images so a weight  $\frac{k}{k+1}$  is affected to its intensities while  $I_{k+1}$  intensities get a weight  $\frac{1}{k+1}$ . At the end, those two images are transformed and weighted according to their contribution to the atlas.

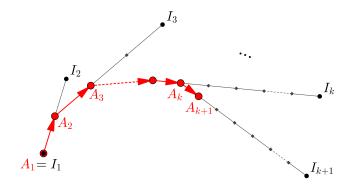


Fig. 1. Shift of the centroid as new images are incorporated.

Unlike [6], we are dealing with images and therefore interpolations on a voxel grid to reconstruct a new image after a transformation. This resampling step has to be avoided as much as possible to keep a maximum of details. To this end it is preferable not to use  $A_k$  (already a resampled image) when constructing  $A_{k+1}$ , but instead to operate on the initial images using transformation compositions such that each image undergoes only one resampling. This is achieved by expressing the sequence  $(A_k)_{k\in\mathbb{N}}$  using only the initial images  $\{I_j, j=1,\ldots,k\}$ . Rearranging Eq. 3, the following formulation emerges:

$$A_{k+1} = \frac{1}{k+1} \left( \sum_{j=1}^{k} \left( I_j \circ L_j \circ \Theta_{j,k} \circ T_{k+1}^{-\frac{1}{k+1}} \right) + I_{k+1} \circ L_{k+1} \circ T_{k+1}^{\frac{k}{k+1}} \right)$$
(4)

Where  $\theta_{j,k}$  is the non linear part of the transformation bringing  $I_j$  onto  $A_k$ . There are k compositions using BCH to perform at iteration k which correspond to the update of the transformations  $\theta_{j,k}$  for  $j \leq k$ .

**Initialization:** One can start at iteration 1 by initiating  $A_1 = I_1$  and  $\theta_{1,1} = Id$ . However, a very interesting point of the method is the possibility to complement an already existing atlas  $A_p$  constituted of p images as long as it can be written as  $A_p = \frac{1}{p} \sum_{j=1}^p I_j \circ L_j \circ T_j$ . To do so, we simply assign  $\theta_{j,p} = T_j$ ,  $\forall j \leq p$ .

**Iterative procedure:** At the end of iteration k, we have, for each  $j \leq k$ , transformations  $\theta_{j,k}$  that map images  $I_j$  onto the atlas  $A_k$ :  $A_k \sim I_j \circ L_j \circ \theta_{j,k}$ .

To build  $A_{k+1}$  from these, we do the following: 1- we register  $I_{k+1}$  onto  $A_k$ :  $A_k \sim I_{k+1} \circ L_{k+1} \circ T_{k+1}$  (one registration); 2- for  $j \leq k$ , we compute the updated  $\theta_{j,k+1}$  as  $\theta_{j,k+1} = \theta_{j,k} \circ T_{k+1}^{-\frac{1}{k+1}}$  (k BCH); 3- we assign  $\theta_{k+1,k+1} = T_{k+1}^{\frac{k}{k+1}}$ ; and finally 4- we build  $A_{k+1} = \frac{1}{k+1} \sum_{j=1}^{k+1} I_j \circ L_j \circ \theta_{j,k+1}$ .

#### 2.3 Unbiased atlas construction

So far, we have only assumed that the registration produced a linear and a non linear part. Going in more details, let us now assume that this registration can be written  $A \sim I \circ R \circ S \circ D$  where:  $R \circ S$  is an affine transformation decomposed into a rigid part R and a stretching part S, and D is a diffeomorphism. Herebefore, by taking  $L = R \circ S$  and T = D i.e. L encapsulating all the linear transformations and T only local displacements, the method will lead - similarly to [8] - to an atlas unbiased up to an affine transformation. On the other hand, if we take L = R and  $T = S \circ D$  i.e. L only considering rigid motion while T encapsulating both growth and local displacements, the method will lead to an atlas unbiased up to a rigid transformation (useful typically for longitudinal studies).

## 3 Experiments and results

### 3.1 MRI database

For our experiments, we have used T1 weighted preprocessed images (brain extracted) from 100 subjects randomly chosen among the Human Connectome Project (HCP) database (for more details, see Van Essen et al. [7]). The size of the images is  $260 \times 311 \times 260$  with a voxel size of 0.7 mm isotropic.

## 3.2 Implementation details

**Registration** Our method is agnostic to the registration used. We performed registrations in two steps using symmetric block matching algorithms [5,4] available in Anima <sup>1</sup>:

- 1. An affine registration that outputs a transformation matrix B which is decomposed through polar decomposition into a rigid part R and a stretching part S as depicted in Legouhy et al. [10].
- 2. A diffeomorphic registration that directly outputs the SVF of the diffeomorphism log(D) linking the two images. This allows to take advantage of the log-Euclidean framework without logarithm computation [13,2].

We chose to compute at lases up to a rigid transformation. Therefore, the linear transformation L is rigid: L = R and the non linear transformation is the combination of stretching and local displacements:  $T = S \circ D$ .

<sup>&</sup>lt;sup>1</sup> Anima: Open source software for medical image processing from the Empenn team. https://github.com/Inria-Visages/Anima-Public - RRID:SCR\_017017

**Transformation composition** Compositions of two diffeomorphisms are performed on their SVFs using order 2 BCH. To compose a linear transformation L and a SVF, we first apply  $\log(L)$  to the real coordinates of an image grid to transform it into an SVF and then compose using order 2 BCH.

## 3.3 Comparison between iterative centroid and direct atlasing

In the following experiments, we evaluate the quality of the iterative centroid atlas and its divergence with respect to direct atlasing, here a method developed in Legouhy et al. [10] available in Anima-Scripts <sup>2</sup> based on Guimond et al. [8]. For both methods, we used the same registration and composition tools with the same parameters. Starting from an atlas built with 50 images using direct atlasing, we added one by one the 50 other images using our online atlasing. We then compared the direct atlases computed at several steps with the online ones using three metrics.

**Divergence** Both methods output linear and non linear transformations (denoted  $T_j$  for online atlasing and  $\tilde{T}_j$  for direct atlasing) mapping each image onto the atlas. We first propose to evaluate the divergence between the two atlases A and  $\tilde{A}$  by an image  $\delta$  defined from the transformations:  $\delta\left(A,\tilde{A}\right)=$ 

 $\frac{1}{n}\sum_{j=1}^{n}\sqrt{\sum_{d=1}^{3}\left(T_{j,d}-\tilde{T}_{j,d}\right)^{2}}$  where d indexes the three components in space. This measure has the advantage of being agnostic to edge issues that can happen in intensities comparisons.

Fig. 2 presents a qualitative view of the evolution of the differences between the direct and online atlases with respect to the number of subjects. We observe that most of the differences increase with the number of added subjects and occur in the cortical areas while central structures of the brain are spared. Fig. 3 gives quantitative plots over the brain of the divergence measures. It suggests that the divergence tends to grow (from a median of 0.777 mm for 60 images, to 1.095 mm for 100) but at a slow pace that decreases (+0.153 mm between 60 and 70 images but only +0.025 mm between 90 and 100).

Iconic evaluations The second part of the evaluation assesses the quality of the atlases based on image intensities. We evaluate first the atlas standard deviation across subject images after transformation on the atlas, the transformations being constructed differently upon the atlasing method. This measure allows to quantify how well the images are aligned after atlas construction (the lower the standard deviation, the better):  $\operatorname{sdis}(A) = \sqrt{\frac{1}{n} \sum_{i=j}^{n} (I_j \circ L_j \circ T_j - A)^2}$ . For better interpretation, we normalized sdis by the average of the intensities of the atlases from the two methods to create  $\operatorname{sdis}_n$ .

<sup>&</sup>lt;sup>2</sup> Anima-Scripts: Open source scripts using Anima software for medical image processing from the Empenn team. https://github.com/Inria-Visages/Anima-Scripts-Public - RRID:SCR\_017072

In addition, we propose a measure of the sharpness of the atlases. Given N(i), a patch around a voxel i of an image A, we define a local measure of the sharpness at voxel i,  $\operatorname{sha}(A(i))$ , as the standard deviation of the patch normalized by its average:  $\operatorname{sha}(A(i)) = \operatorname{sd}_{N(i)}(A)/\operatorname{mean}_{N(i)}(A)$ . In our experiments, we chose a patch of diameter 5 voxels. We present in Fig. 4 a quantitative view over voxels inside the brain mask of these two iconic measures with respect to the number of images in the atlas. These results show that both methods output atlases that are very similar in terms of image quality. The sharpness remains equivalent for both methods independently of the number of images added. sdis<sub>n</sub> increases only between 60 and 70 images and is then stable, but the evolution and values are the same for both atlas construction methods.

### 4 Conclusion

We have presented an online atlasing method that allows the incorporation of new images into an existing atlas without having to restart the atlasing process from the beginning. Based on an iterative centroid process, this approach only necessitates one registration per new image. We observed that the divergence between our method and direct atlasing using the same tools is localized in cortical areas and tends to grow but at a slow and decreasing pace. Also, the obtained atlases from both approaches have shown no differences in terms of image quality. The trend being at large, growing databases, the proposed online atlasing

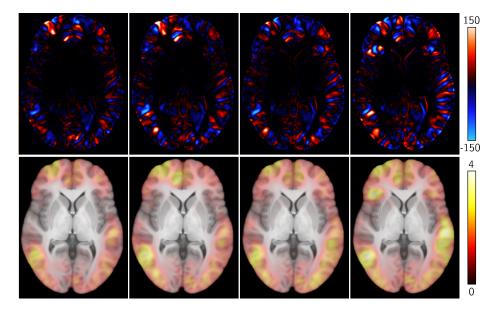


Fig. 2. From left to right at lases with 60, 70, 80 and 100 images. Top: intensity differences between iterative centroid and direct at lases. Bottom: at lases from iterative centroid method overlaid with divergence measure  $\delta$ .

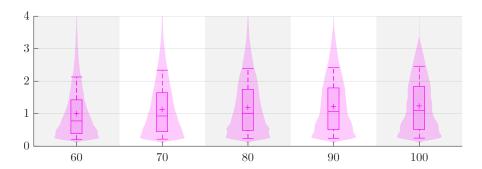
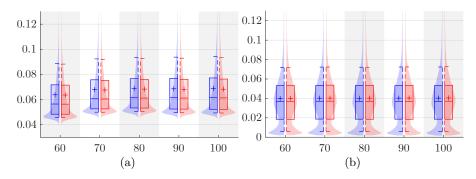


Fig. 3. Boxplots and violin plots of the divergence  $\delta$  between the atlases (restrained to the union of the masks) with 60, 70, 80, 90 and 100 images.



**Fig. 4.** Boxplots and violin plots of inter-subjects normalized standard deviations  $sdis_n$  (a) and sharpness sha (b) for atlases (restrained to the union of the masks) with 60, 70, 80, 90 and 100 images. Blue: iterative centroid. Red: direct atlasing.

method offers an interesting tool to update an atlas at reasonable computational cost as new images arrive. Finally, we derived a construction method up to either an affine or a rigid transformation. This makes the method especially eligible for longitudinal atlasing as a future work, simply by adapting the weight of each image in Eq. 4.

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