Intrinsic 2D/3D registration based on a hybrid approach: use in the radiosurgical imaging process.
Maximilien Vermandel, Nacim Betrouni, Jean-Yves Gauvrit, David Pasquier, Christrian Vasseur, Jean Rousseau

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INTRODUCTION

Recent decades have seen the emergence of new medical imaging techniques, such as the Computerized Tomography (CT) scanner, Nuclear Medicine (NM), Magnetic Resonance Imaging (MRI) and Positron Emission Tomography (PET). All these modalities have favoured the progressive use of digital imaging to such an extent that even the oldest imaging modality, radiology, now uses digital sensors such as brightness amplifiers and flat CCD sensors. The increasing acquisition speed, digital storage and the variety of modalities are gradually changing the daily working practices of physician, who have access to a large amount of data to refine their diagnoses. This profusion of images has given rise to several disciplines such as the management of image flows (storage, access, etc.) and post-treatment. Post-treatment ranges from image processing, improving image quality, quantification, classification, to registration and image fusion. The registration firstly became necessary in order to fuse the images coming from different modalities and whose information was complementary (e.g. morphological and functional images). Then, the field of application became open to more surgical or interventionist practise. For example, neurosurgery required the combined use of morphological images, defining the pathological areas, and activation images, defining the brain’s functional areas. The work described in this paper concerns the planning for the treatment of Arteriovenous Malformations (AVM) in radiosurgery. We present an intrinsic registration methodology that does not use any external reference and can be used to fuse radiological images of the arteriovenous system (the 2D X-ray Angiography (XRA) image) and Magnetic Resonance Angiography images (MRA, 3D). We first define the terms and definitions used in this paper. Then, after a brief description of imaging and radiosurgery methods, we present the various existing registration techniques and those developed by the authors to fuse XRA and MRA images. Finally, we describe the validation protocol that has been developed and some preliminary results.

Terms and definitions

Registration – As a general rule, registration (1,2) can be defined as the search for a mathematical relation between two data sets in order to be observed in a common space. For example, rigid image registration is the search for the
transformation $T$, combining rotation and translation, allowing the transition from a floating image $F$ to a reference image $R$. In this case, each image represents the same object but, when the acquisition conditions are different, a transformation has to be applied to image $F$ so that it can be superimposed on image $R$.

The image registration can be performed using two procedures:

- using an external reference whose geometry is known (i.e. stereotactic frame, fiducial markers).
- using the similarities existing between the images, only the anatomical content is used for registration, this is a so-called intrinsic registration. Two methods of computation must be highlighted:
  - feature-based approach which uses geometrical features extracted from image data such as surface, contour...
  - intensity-based approach which uses a statistical dependency between images according to grey-level.

Image fusion — Image fusion is often used with the aim to view two images simultaneously. Once the registration has been performed, the two images can be superimposed. They can be viewed simultaneously in the same space by using various algorithms (3) (transparency, pixel based fusion, segmented based fusion, etc.).

MATERIALS AND METHODS

Radiosurgery

AVM consists of a disease in the vascular network, one of the commonest malformations being the angioma (Fig. 1).

An angioma is a network of fine secondary and parasitic blood vessels (nidus), connecting the arterial and venous systems and thereby causing disruption to cerebral blood supply. Intracranial AVMs are usually diagnosed by a diagnostic XRA visualizing the cerebral blood vessels. Depending on its size, its treatment may require a one shot irradiation delivered to the patient by radiosurgery (4,5). This treatment requires the combined use of vascular and anatomical images such as XRA and MRI. These two exams are acquired using a stereotactic frame fixed to the patient's head. The frame is required to locate the lesion within the space of the Treatment Planning System (TPS). Once the AVM has been located within the TPS, the position of the irradiation beams are calculated taking into account sensitive organs such as the optic chiasma, eyes and brain stem. The aim of this paper is to present a new methodology for locating AVM in the TPS space using diagnostic XRA instead of framed XRA.

Our method is illustrated here through the Gamma Knife radiosurgery but the methodology is also valid for other radiosurgical techniques. The Gamma Knife mainly consists of a radiation unit (Fig. 2a) containing 201 $^{60}$Cobalt sources (30Ci at delivery, radioactive half-life 5.27 years) and a moveable helmet with a choice of four collimators of 4, 8, 14 or 18mm diameter. The 201 orientations that are available on the Gamma Knife, when no collimator is obstructed by a plug, are shown in Fig. 2b.

The aim of planning treatment is to irradiate the target volume with a more suitable dose, while sparing the healthy surrounding structures. It must be achieved with the accuracy of about one millimetre. The coordinates of the isocentres are determined on the basis of the dimensions, the shape and the location of the lesion identified on the images. Normally, several target centres are needed to cover the entire volume, which involves repositioning the patient by an electromechanical system.

Fig. 2  (a) the Gamma Knife and principle of irradiation with the gamma knife using multiple photon beam, (b) shows all the available beams, for a treatment only few of them are kept, all the other are “plugged”.

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Fig. 1b  From MRI view.

Fig. 1a  Arteriovenous Malformation, from X-Ray view.
The Gamma Knife requires the use of a stereotactic system (6) (Fig. 3 and 4) to locate the target during irradiation. The stereotactic frame is screwed onto the external surface of the patient's head and remains fixed to the patient from the imaging exams to the treatment.

The bone structures are eliminated by subtracting the first image of the series (in which only the bone structures appear) from the images with the contrast medium (Fig. 6), hence the term Digital Subtracted Angiography (DSA).

Radiosurgery of AVMs requires planning using anatomical and vascular images. The anatomical images are acquired using CT and MRI. The vascular images are acquired by X-ray angiography (XRA), which provides images of the cerebral vascularisation with an excellent spatial and temporal resolution. XRA is used as the reference for locating the target. Angiography technique is particularly invasive since it requires the use of a catheter that is guided from the femoral artery to the carotid and a large amount of contrast agent has to be injected. The acquisition (up to 10 images/s) provides an indication of the arterial and venous blood flow times (Fig. 5).

The standard sets of images needed for the treatment of the disease are shown in Fig. 7. The disease is diagnosed by a standard DSA (without stereotactical frame), but, for the planning, a second DSA is acquired with the locating stereotactic frame in position as well as MRI and CT. We propose here to ease the imaging procedure by suppressing the framed DSA acquisition (7).

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reference space). Note that the MRA acquisition is not a constraint of the proposed method because it is increasingly requested by neuro-radiologists for post-treatment follow-up. Generally, the acquisition by Magnetic Resonance Angiography relies on two techniques. The first requires the injection of gadolinium, a substance that locally modifies magnetic susceptibility. The second is based on the blood kinetics in the axial plane; this technique is called Time Of Flight (TOF).

Image acquisition

MRA images were acquired on a Siemens Magnetom VisionMRI scanner. (see table 1) and DSA images were acquired on a Philips Allura V5000 angiograph (see table 2).

Table 1  MRA images parameters.

<table>
<thead>
<tr>
<th>Sequence Type</th>
<th>Gadolinium MRA</th>
<th>TOF</th>
</tr>
</thead>
<tbody>
<tr>
<td>Magnetic Field</td>
<td>1.5T</td>
<td>1.5T</td>
</tr>
<tr>
<td>Slice thickness</td>
<td>0.5mm</td>
<td>0.7mm</td>
</tr>
<tr>
<td>Plan</td>
<td>Face</td>
<td>Transverse</td>
</tr>
<tr>
<td>TR</td>
<td>6.8ms</td>
<td>37ms</td>
</tr>
<tr>
<td>TE</td>
<td>2.3ms</td>
<td>7.2ms</td>
</tr>
<tr>
<td>Row and Col</td>
<td>0.5mm</td>
<td>0.5mm</td>
</tr>
<tr>
<td>PixelSize</td>
<td></td>
<td></td>
</tr>
<tr>
<td>Number of Slices per volume</td>
<td>130</td>
<td>80</td>
</tr>
<tr>
<td>Resolution</td>
<td>512x512</td>
<td>512x512</td>
</tr>
</tbody>
</table>

Table 2  DSA images parameters.

<table>
<thead>
<tr>
<th>Sequence Type</th>
<th>Multiframe acquisition</th>
</tr>
</thead>
<tbody>
<tr>
<td>Source Intensifier Distance</td>
<td>variable</td>
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<tr>
<td>Paired projection</td>
<td>no</td>
</tr>
<tr>
<td>Row and Col PixelSize</td>
<td>variable</td>
</tr>
<tr>
<td>Number of projections per exam</td>
<td>24</td>
</tr>
<tr>
<td>Resolution</td>
<td>1024x1024</td>
</tr>
</tbody>
</table>

Registration techniques

The aim of registration methods between volume and projections is to determine the best position of the tomographic slices in the space of the planar modality (Fig. 9).

The 2D/3D registration may appear to be relatively simple if there is an external reference frame, such as Leksell frame, that is firmly fixed to the object and readily identifiable in each imaging modality. The problem is much more difficult if there is no frame. The only possible relation becomes the object itself, which does not have the same grey level from one modality to another.

2D/3D registration has been the subject of innovative developments. Two classes of approaches can be distinguished: intensity-based methods and geometric or feature-based methods. We propose here a third class. Hybrid approach is a compromise between intensity-based and feature-based methods. Overall, the methods are all based on the same principle, where:

(A) From its position in the projection space, the volume is virtually projected onto a plane so as to obtain a new 2D floating image. This is then compared to the 2D reference image (DSA).

(B) The difference between the two images is estimated using a similarity measure.

(C) The difference observed is then used to calculate a new position of the volume.

Intensity-based approaches – Intensity-based approaches (8,9,10) rely on statistical dependence between planar images and volume. The principle is the same whatever the application environment. The registration is done iteratively, by projecting the volume and comparing the result with the planar image. The strategy consists in creating digital radiographs from tomographic slices (Digitally Reconstructed Radiography, DRR). DRRs are compared to the native projection image. DRRs are generated using a pinhole model projection, identical to radiography (11), so that the corresponding result is an image similar to an X-ray image. The projection conditions are shown in Fig. 10.

From the virtual source, projected points result from the integration of values encountered along the corresponding ray (see eq. 1).
Intrinsic 2D/3D registration based on a hybrid approach: Use in the radiosurgical imaging process.

\[ I_{\text{out}} = \frac{\text{det vector}}{\text{source}} \int I(x, y, z) dx \, dy \, dz \]  

(1)

Where \( I_{\text{out}} \) the intensity after attenuation at the exit of the volume, \( I(x, y, z) \) is the intensity at the position \((x, y, z)\) such as the voxel grey level.

In some case it could be interesting to use a model closer to the actual photon beam attenuation as shown in eq. 2. The relation between the input and output intensities is therefore the integral of the linear attenuation coefficients along a ray.

\[ I_{\text{out}} = I_{\text{in}} \cdot e^{- \left( \int \text{det vector} \text{source} \beta(x, y, z) dx \, dy \, dz \right)} \]  

(2)

Where \( I_{\text{in}} \) designates the intensity at entry of the volume, \( I_{\text{out}} \) the intensity after attenuation at the exit of the volume, \( \beta(x, y, z) \) the attenuation coefficient at the position \((x, y, z)\) in the volume. Theoretically \( \beta(x, y, z) \) is different for each type of tissue crossed by the beam, so in practice a constant value multiply by the grey level could be used to ease the computation.

The comparison between the DRRs and the original radiographs uses a similarity measure. Penney (8) provided a comparison of several similarity measures applied to 2D/3D intensity-based registration.

**Feature-based approaches** - Feature-based approaches (12, 13) rely on the registration of geometric 2D and 3D primitives extracted from the images segmentation. The registration is then performed by an iterative process where the projected 3D structures are compared to the 2D structures through a least mean square distance for example. An illustration of feature-based approach is given Fig. 11.

![Fig. 11 Illustration of the feature-based registration using skeleton, MRA/CT slices are segmented, centre lines are extracted and projected to the projection plan, the 2D skeleton obtained is compared to the centre lines extracted from X-ray angiography.](image1)

**The hybrid approach** The hybrid registration we have developed (7, 14, 15) is a compromise between these two techniques and is applied on non paired DSA images. A part of segmentation of the feature-based approach is retained for the MRA (Fig. 12 a), which is subjected to a semi-automatic processing to extract a part of the tree-structure. On the other hand, the DSA is not subjected to any processing other than the subtraction that is routinely used for removing the bone structures. This configuration is adapted from intensity-based registration.

The segmented MRA volume can be placed in the DSA space and virtually projected onto the imaging system sensor. For the projection, a fast solution fitting a photon attenuation model has been developed. This virtual X-ray projection allows a statistical dependency to be used, whatever the tomographic imaging method (Fig. 12 b).

![Fig. 12 Hybrid registration, (a) reconstructed volume, (b) virtual DRR using fast projection of the vessel surface.](image2)

The part below describes point by point our methodology:

- **Segmentation of MRA volume**
  First, we generate a Maximum of Intensity Projection (MIP) as described below (Fig. 13 a). A specific artery is selected on the MIP image (Fig. 13 b) using a region growing algorithm supervised by the physician.

![Fig. 13 Maximum of Intensity Projection algorithm, (a) the maxima of grey levels among a ray are projected to a plan, which gives the image shown in (b).](image3)

The resulting segmented MIP image is used to detect the vascular structure through the entire MRA dataset, applying fuzzy set theory and data fusion. For each voxel a vessel membership degree is computed, its value is between 0 and 1, 1 being the highest membership degree to a vessel. This degree (14) takes into account the Contrast to Noise Ratio, the neighborhood of each voxel. Fig. 14 shows an example of automatic detection performed by the algorithm where a threshold at 0.5 has been applied on the membership degrees. Once the contours of the vessel structure have been found, we use the method of 3D reconstruction proposed by Vial et al. (16).

![Fig. 14 Resulting contour obtained with a threshold of 0.5 applied to the membership degree of the fuzzy image of the MRA during slices segmentation.](image4)
Projection

Although DRR and virtual X-ray projection are quite similar, it must be pointed out that DRR is based on the projection of voxels (discrete), and therefore a complete tomographic volume, whereas our X-Ray projection is computed using a surface (continuous) defined by a series of vertices and polygons. This solution produces a reduced data set that provides a much faster projection while taking into account the attenuation through the matter.

Our projection algorithm relies on the projection of polygons (modeling the surface of the reconstructed volume) using the Z-Buffer algorithm to compute the thickness of matter crossed by the virtual photon beam. The grey level is then computed using the equation (2). The Fig. 15 shows the principle in 2D.

![Image 1](https://example.com/image1)

**Fig 15** Illustration of the projection algorithm used to simulate X-ray beam attenuation: this algorithm uses a "distance- buffer" and a "Z-buffer" to calculate the thickness of vessel which has been crossed.

At the end of the process, each pixel of the projection has a corresponding thickness of crossed matter stocked in the "distance buffer", thus finally each pixel is set to:

\[
\text{pixel} = \text{GLMAX} \cdot e^{(-\alpha \cdot \Delta x_{\text{int}})}
\]

(3)

Where GLMAX is the maximum of grey level (according to the DSA), \(\Delta x_{\text{int}}\) is the thickness stocked in the distance buffer and \(\alpha\) represents the attenuation coefficient of the crossed matter (taken at 0.15).

The short computing time for each projection enables to achieve complete registration in about 2-3 min per DSA view on a standard computer (Pentium IV, 2Ghz, 1 Go of RAM).

Similitude measure

The similarity measure calculated here to compare projection of the MRA volume and DSA is a basic quadratic distance between virtual projection and DSA, as shown on eq. 4.

\[
\text{Energy} = \sum_{i=1}^{I} \sum_{j=1}^{J} \left( P_{i,j} - P'_{i,j} \right)^2
\]

(4)

Where, \(P_{i,j}\) and \(P'_{i,j}\) are respectively the \(i\) and \(j\) coordinates pixels of the original image \(I\) and the image \(I'\) computed from projection. This measure has been used because it is very fast to compute and it is very sensitive to small translation which may exist between DSA and virtual X-ray projection.

Initialisation

The initialisation is driven interactively by the physician who roughly disposes the volume projection on the DSA, initializing in this way the 6 degrees of freedom.

Optimization

To optimize the volume position, we apply an algorithm developed by Salazar (17) and based on the Hybrid Monte Carlo algorithm (HMC). This method, known as the Hybrid Simulated Annealing (HSA), has the property to modify simultaneously the parameters set using a simulated annealing scheme. Thus, HSA is almost as fast as a conjugate gradient but avoiding local minima. In its standard approach, the HMC introduces a set of auxiliary momenta variables \(p_1, p_2, \ldots, p_N\) and a related Hamiltonian function \(H(x,p)\), where \(x\) is the \(N\) parameters vector to optimize (here the six degree of freedom), given \(E\) the energy introduced by (4).

\[
H(x,p) = E(x) + \frac{1}{2} \sum_{i=1}^{N} p_i^2 = E(x) + \frac{p^2}{2}
\]

(5)

Vector \(p\) is a set of independent, Gaussian distributed, random variables of zero mean. New configurations are computed using the Hamilton's equation of motion, described by (4).

\[
\dot{x}_i = p_i, \quad \dot{p}_i = -\frac{\partial E}{\partial x_i}
\]

(6)

The Hamilton's equation of motion are numerically integrated using a "leap-frog" scheme with a time step \(\delta t\). In this way each new configuration, defined by \(x'\) and \(p'\), are computed by a set of relations (7). Where, \(F(x) = \frac{\partial E(x)}{\partial x}\), is the force acting on the variable \(x_i\).

\[
x' = x_i + \delta t \cdot p_i \cdot \frac{\delta E}{\partial x_i}, \quad \text{and} \quad p' = p_i - \delta t \cdot \frac{\partial E(x)}{\partial x'_i}, \quad i = 1, \ldots, N
\]

(7)

As in the standard simulated annealing, the new configuration is accepted using the Metropolis test, with a probability described by (8).

\[
P(\Delta H) = e^{(-\Delta H / T)}
\]

(8)

The acceptance probability is computed from Hamiltonian variation \(\Delta H = H(x', p') - H(x, p)\) where \(T\) is the step temperature. When, stability is observed during a step, the temperature is cooled. The process is stopped when the global system is stable or when the temperature is equal to zero.

Our hybrid solution that is evaluated here for registration in radiosurgery has therefore proved to be very efficient in terms of accuracy and speed thanks to the fusion of conventional methods. From 2D coordinates to 3D coordinates – Thanks to registration, the DSA views are paired using the MRA volume as a common reference. In such a configuration we can use the principle of epipolar geometry. Once projections are paired, the use of epipolar principle makes a point on one view corresponding to a point on specific line in the second view (Fig. 16).

**Fig 16** Illustration of epipolar geometry where p1 and p2 are the projection of a given target. Epipolar geometry enables to get the coordinate of the target when p1 and p2 are known and views paired.
In this way it is possible to navigate in the MRA and DSA projections (and vice-versa) by using the shared reference system defined by the volume (or the frame, in Leksell configuration).

RESULTS

The possible use of 2D/3D registration in the irradiation planning in radiosurgery involves a stage of evaluating the interchangeability of the anatomical registration solution with the gold standard, which is registration with the Leksell frame.

Therefore two stages were achieved: an evaluation of the accuracy of the hybrid solution, on phantom and clinical data and a ground truth validation. It should be noted that, prior to this study, the robustness and re-projection error (18) were already evaluated in a previous work (14). The results obtained therefore enabled us to initiate this study.

Evaluation of the accuracy on a phantom

A phantom of the vascular network (Fig. 17) was used during the stage of evaluating the accuracy. Its known geometry enabled us to determine the accuracy of the stereoscopic pairing of DSA views. MRA and DSA of the phantom were acquired. We obtained coronal, sagittal and axial image sets from MRA and four DSA projections. The volume reconstructed from MRI sets was registered with each of the four projections. The distances between the different "branches" of the vascular network were measured each couple of projections and compared to the known physical distances.

The measurement of the distances used the principles of epipolar geometry (Fig. 16). A point was chosen on a first DSA view and its homologue on the other view. The intersection of the two corresponding epipolar straight lines gave us the coordinates of the 3D point in the DSA space. The same operation repeated for each point (Fig. 18) of a segment finally drives us to determine 3D distances.

During the evaluation stage the extremity of the segment appeared with much more contrast than the rest of the phantom, which assumes to have a good robustness and reproducibility of measurement.
This method of validation by epipolar geometry is particularly interesting since it tests both the validity of the registration, which is in our case the only means of pairing DSA projections, and it provides the quantitative accuracy of the registration. The 3D distance errors obtained are given in Table 3.

Table 3 Results of measurements on the phantom where 3D distances were measured using the interactive navigation tools (Fig. 18).

<table>
<thead>
<tr>
<th>Root Mean Square Error</th>
<th>0.7mm</th>
</tr>
</thead>
<tbody>
<tr>
<td>Standard Deviation error</td>
<td>0.6mm</td>
</tr>
<tr>
<td>Maximum error</td>
<td>0.9mm</td>
</tr>
</tbody>
</table>

The error presented is based on the difference between the real distance and the measured distance.

Evaluation of the accuracy under clinical conditions

For the patients study, a good way to estimate the accuracy is to use non-invasive fiducial marker. Unfortunately, MRA and DSA could not be acquired on the same day. Using markers in both modalities cannot be reproducible because of the lack of accuracy when placing markers for MRA and then for DSA. In this configuration it is impossible to measure reprojection error or 3D distance error. Actually, in our methodology the DSA views are not initially paired, only the registration with a single volume can pair the views. Thus we have chosen to use only markers with DSA and to estimate the epipolar error which is the deviation between two epipolar straight lines. Since epipolar error gives the accuracy of pairing projections, it constitutes an indirect but absolute measure of registration accuracy.

Images were acquired using a standard clinical protocol defined by physician and used for aneurysm treatment. In accordance with the clinical protocol, three non invasive markers were placed for 4 patients during their DSA exams. No particular orientation constraint nor additional image was required. Each marker is selected on a view and its homologue on the other, and the error is computed. Using small markers assumes to have a good robustness and reproducibility of measurement. The epipolar errors obtained during the measurement on patients are given in Table 4.

Table 4 Results of measurements obtained on clinical data, 3 markers are selected on the DSA views, the error presented is the epipolar error.

Maximum error | 0.8mm |
Standard deviation error | 0.3mm |

"Ground truth" validation

In this study, the term "ground truth" validation means the comparison of our method with the real radiosurgery conditions practised using the Leksell frame. For the validation, as the MRI-Leksell transformation is known, we have to compare the coordinates of the same points obtained by Leksell localization and the 2D/3D registration.

For this protocol, 3 patients treated by radiosurgery were given DSA and MRA localization exams. These exams were processed firstly by standard localization using the Leksell reference system and secondly using 2D/3D registration. It was therefore possible to validate the registration by means of an objective comparison of the two techniques, the Leksell localization being the so-called ground truth.

The validation (Fig. 19) was performed using the DSA.

Fig. 19 Detail of transformations needed to obtain Leksell coordinates from the DSA space depending on initial conditions: intrinsic registration or Leksell registration.

On each of the views, 4 points, designating in each case the same 3D element (elements of the Leksell frame to avoid bias during the measures), was selected. In the localization configuration using the stereotactic frame, the 3D coordinates of the point can be deduced directly within the Leksell space. In the second case, where the localization is deduced from registration, the 3D coordinates of the point are calculated in the MRA reference system from the same series of DSA coordinates.

To compare the measurements, the points obtained in the MRA space have to be converted to the Leksell reference system by means of the MRA → Leksell transformation. In this way we obtain two series of coordinates in the Leksell space for the same 3D point. The deviation measured illustrates the error between the two localization methods. The results are given in Tables 5 and 6.

Table 5 Results obtained for measurements achieved with and without Leksell localization.

<table>
<thead>
<tr>
<th>Patient</th>
<th>Coordinates in Leksell space from frame</th>
<th>Coordinates in Leksell space from registration</th>
</tr>
</thead>
<tbody>
<tr>
<td>Lek1</td>
<td>X Y Z</td>
<td>X Y Z</td>
</tr>
<tr>
<td>88.12</td>
<td>92.31</td>
<td>115.37</td>
</tr>
<tr>
<td>88.34</td>
<td>92.60</td>
<td>115.37</td>
</tr>
<tr>
<td>88.54</td>
<td>92.98</td>
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<td>88.74</td>
<td>93.36</td>
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</tr>
<tr>
<td>88.94</td>
<td>93.74</td>
<td>115.37</td>
</tr>
<tr>
<td>Lek2</td>
<td>X Y Z</td>
<td>X Y Z</td>
</tr>
<tr>
<td>101.31</td>
<td>104.5</td>
<td>115.37</td>
</tr>
<tr>
<td>101.51</td>
<td>105.8</td>
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<td>106.5</td>
<td>115.37</td>
</tr>
<tr>
<td>101.91</td>
<td>107.1</td>
<td>115.37</td>
</tr>
<tr>
<td>Lek3</td>
<td>X Y Z</td>
<td>X Y Z</td>
</tr>
<tr>
<td>102.11</td>
<td>105.1</td>
<td>115.37</td>
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<tr>
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<tr>
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<td>105.1</td>
<td>115.37</td>
</tr>
<tr>
<td>102.71</td>
<td>105.1</td>
<td>115.37</td>
</tr>
<tr>
<td>Average Error</td>
<td>1.84</td>
<td>Maximum error</td>
</tr>
</tbody>
</table>

The error presented is the distance between the 3D coordinate given by the Leksell registration and the 3D coordinates given by the hybrid registration.
The error is the absolute difference between a component given by the Leksell registration and a component given by the hybrid registration (e.g. X or Y or Z).

### DISCUSSION

The intensity-based method has been found to be very precise but relatively time consuming and suitable for the registration of images produced by the same physical phenomenon (e.g. X-rays for CT and DSA) because of their statistical dependence. For example, Murphy (10) had proposed an intensity-based approach to match CT and DSA for a similar application in radiosurgery. His approach has been proved to be precise. Unfortunately, it cannot be applied in our case for two reasons: Murphy’s method required paired projections and we work on MRA images.

Feature-based methods are very fast but have a significant imprecision for radiosurgery. Furthermore, the many intermediate steps (e.g. segmentation) limit their use.

Finally, in most of cases proposed in the literature for intensity or feature-based method, registration relies on paired DSA images, which is not the case of the hybrid approach.

During the evaluation stage and after registration, the measurements of the distances on the phantom gave perfectly acceptable results, since the maximum distance error observed was always less than one millimetre.

The accuracy evaluated on clinical data confirmed that the epipolar constraints were verified after registration. In practice, the epipolar straight lines were only a few tenths of a millimetre apart (max. error 0.8mm, SD 0.3). This weak error confirms the good accuracy of the registration when using injected MRA.

For the validation stage during the radiosurgery protocol, the measurements showed an acceptable error on the X and Z axes, but revealed a significant error on the Y axis. This observation led us to reconsider the quality of the TOF images used for the validation because a such a phenomenon does not appear when using gadolinium.

It appears that the TOF images required very high spatial coding gradients that are potentially non-linear, and could be the source of major distortions in the XY plane (acquisition plane for axial images).

We were able to confirm this hypothesis by taking measurements of the distances on the Leksell frame on axial images (Fig. 20) taken with a TOF sequence (a) and with a standard sequence (b) resulting in much smaller gradients. It was clearly evident that the TOF images were subjected to a deformation in the Y axis. This deformation led to an error during the reconstruction of the registration volume, which made determination of localization matrices imprecise. Extra-fast sequences do not therefore seem to be very suitable for registration, unless a method can be developed for correcting the distortions caused, which is not currently the case.

### CONCLUSION

In this study we have described the validation steps required for applying registration of multimodality images in radiosurgery. These registration techniques will eventually allow us to replace the DSA localization exam by the standard diagnostic exam.

The protocol presented here is based on an evaluation of accuracy and on a validation under clinical conditions, where validation means comparing intrinsic registration with a gold standard technique.

The validation procedure that we have applied has revealed a lack of accuracy (maximum error 2.83mm, SD 0.54), making the method unsuitable for radiosurgery treatment. However, a study of the errors observed has highlighted problems associated with TOF images. Their intrinsic distortions in the XY (axial) plane make it difficult to perform an accurate registration. But the routine use of MRA/DSA registration could still be envisaged. MRA with the injection of a contrast agent seems to be more suitable for this application, as shown by the accuracy study.

The studies conducted on the phantom and on patients are however fairly conclusive and lead us to think that routine use is very probable in the future. The degree of accuracy obtained is equivalent to the accuracy when using external markers. It therefore seems possible to quantify an AVM by using...
frameless and non-paired DSA images, since MRA acquisition is increasingly being requested by physicians. For the reasons exposed in the introduction, the substitution of the localization DSA by the diagnostic exam would greatly improve the treatment protocol. If the method is to be used routinely in clinical practice, further studies are needed on the 3D mapping of epipolar errors, which would constitute a model of registration error and which would allow us to observe the spatial limits of registration. A complete clinical study will also be needed. The clinical test protocols, defined by the various specialists involved in Gamma Knife treatment (Neurosurgeons, Neuroradiologists and Radiotherapists) will require the combined use of localization and diagnostic images. The localization images are used to plan the operation under standard conditions, whereas the diagnostic images are used in parallel. The dosimetry produced for each type of image could therefore be compared. The measurements shown through the form of Bland and Altman (19) graphs should be used to study the consistency of the dosimetry with and without localization images. It would demonstrate the interchangeability between the Gold Standard and the use of our frameless DSA registration approach. In that manner, our method could be applied in daily practice to improve effectiveness and simplicity of the surgical procedure.

REFERENCES

2. Wan Rui, Li Minglu, “An Overview of Medical Image Registration”, Fifth International Conference on Computational Intelligence and Multimedia Applications (ICCIMA ’03), 2003.