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A Modified ICP Algorithm for Normal-Guided Surface Registration

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ABSTRACT

The iterative closest point (ICP) algorithm is probably the most popular algorithm for fine registration of surfaces. Among its key properties are: a simple minimization scheme, proofs of convergence as well as the easiness to modify and improve it in many ways (e.g. use of fuzzy point correspondences, incorporation of a priori knowledge, extensions to non-linear deformations, speed-up strategies, etc.) while keeping the desirable properties of the original method. However, most ICP-like registration methods suffer from the fact that they only consider the distance between the surfaces to register in the criterion to minimize, and thus are highly dependent on how the surfaces are aligned in the first place. This explains why these methods are likely to be trapped in local minima and to lead to erroneous solutions. A solution to partly alleviate this problem would consist in adding higher-order information in the criterion to minimize (e.g. normals, curvatures, etc.), but previous works along these research tracks have led to computationally intractable minimization schemes. In this paper, we propose a new way to include the point unit normals in addition to the point coordinates to derive an ICP-like scheme for non-linear registration of surfaces, and we show how to keep the properties of the original ICP algorithm. Our algorithm rests on a simple formula showing how the unit normal changes when a surface undergoes a small deformation. The use of this formula in an ICP-like algorithm is made possible by adequate implementation choices, most notably the use of a local, differentiable, parametrization of the surfaces and a locally affine deformation model using this local parametrization. Then we experimentally show the strong added value of using the unit normals in a series of controlled experiments.

1. INTRODUCTION

Non-linear registration is a key tool for many applications in medical imaging: morphometric analyses, follow-up studies, computer-assisted surgery, etc. Surfaces represented by point clouds or meshes are especially convenient to represent a subject’s anatomy, as they allow to focus the subsequent analyses on the structure of interest. Many surface registration algorithms have been proposed in the literature. One of the most popular is the ICP algorithm proposed by Besl & McKay. Over the years it has led to numerous successful variants and extensions providing many ICP-like algorithms using fuzzy correspondences, a priori knowledge, selection of points for faster convergence, robust techniques to deal with outliers, etc. Extensions to non-linear deformations have also been successfully developed. In essence, the ICP algorithm minimizes the distance between the two surfaces. However, some works have tried to add higher-order information in the ICP criterion, to alleviate the well-known local convergence problem of the algorithm, but they have led to intractable solutions.

For instance, Feldmar & Ayache have shown how to include the normals and curvatures. After showing how these quantities are modified by a locally affine deformation, they devised a criterion combining point coordinates, normals and curvatures and showed how to minimize it in an iterative manner similarly to the original ICP algorithm. However, solving for the optimal transformation given correspondences in this framework is a non-linear problem which makes the overall scheme computationally expensive and difficult to implement in case of large surfaces. Another significant work along these lines has been done by Kambhamettu and colleagues in a series of papers where they showed how to properly derive a simple solution to this last problem, using linear algebra, when the hypothesis of a special type of deformations (a subclass of affine transformations) is made. However, in their work this solution was not encompassed in an ICP-type algorithm, and an exhaustive search over all possible pairs of correspondences was made instead to register the two surfaces. Moreover, the
point-to-point distances was not taken into account in their algorithm, while these are of the utmost importance when the normals cannot be faithfully estimated.

In this paper, 1) we propose to use the locally affine transformations subtype proposed by Kambhamettu and colleagues to devise a proper ICP-type registration criterion using these transformations, the point coordinates and the point unit normals, 2) we show how to derive an iterative solution to optimise this criterion, possessing the desirable convergence and simplicity properties of the original ICP algorithm, and 3) we experimentally show that adding the unit normals in the ICP criterion significantly improves the registration.

2. GENERAL SCHEME

The problem of surface registration consists in finding the transformation which best superposes a surface \( P \) with a reference surface \( X \). In this paper we consider surfaces represented by point sets or triangle meshes. We call these sets \( P = \{p_i\}, i = 1 \ldots N_P \) and \( X = \{x_j\}, j = 1 \ldots N_X \). A very fruitful approach to tackle this problem is to introduce point correspondences as intermediate variables. Following this idea, Besl & McKay have proposed the ICP algorithm, whose criterion to minimize has been later modified by Chui and Rangarajan\(^2\) as well as Combès and Prima,\(^3\) and that can be formulated in a very general way as follows:

\[
\varepsilon(T, A) = \sum_{i=1}^{N_P} \sum_{j=1}^{N_X} A_{ij} d_T(x_j, p_i) + \beta L(T) + \gamma \sum_{i=1}^{N_P} \sum_{j=1}^{N_X} A_{ij} \log(A_{ij}) \quad \text{with } \forall i, \sum_j A_{ij} = 1 \text{ and } \forall (i,j), A_{ij} \geq 0 \quad (1)
\]

- \( T \) is the transformation best superposing the surfaces (and that can be rigid, affine, elastic, etc.).
- \( A \) is a \( N_P \times N_X \) matrix that describes the fuzzy correspondences between the surface points. The greater \( A_{ij} \), the more likely the point \( x_j \in X \) to be the correspondent of the point \( p_i \in P \). \( A \) can be viewed as a hidden variable of the registration problem and will be called the match matrix.
- \( d_T(x_j, p_i) \) is a general distance measuring the discrepancy between \( x_j \) and \( T(p_i) \).
- \( L \) is a regularizer allowing to penalize discontinuities of \( T \) over the space, with \( \beta \geq 0 \) weighting its influence.
- \( \sum_{i=1}^{N_P} \sum_{j=1}^{N_X} A_{ij} \log(A_{ij}) \) is a barrier function allowing to control the fuzziness of \( A \) (the higher \( \gamma \geq 0 \), the greater the fuzziness). In practice, this term convexifies the criterion.

The strategy first proposed by Besl & McKay\(^4\) to minimize \( \varepsilon \) consists in alternating the estimation of the point-to-point fuzzy correspondences \( A \) and of the transformation \( T \). This has been shown to converge to an at least local minimum of the criterion \( \varepsilon \). The iterative algorithm can then be simply written as follows:

| Step 0: Initialize \( T^0 \) |
| Step 1: \( A^{n+1} = \arg \min_A \varepsilon(T^n, A) \) |
| Step 2: \( T^{n+1} = \arg \min_T \varepsilon(T, A^{n+1}) \) |
| Step 3: if \( |T^{n+1} - T^n| \geq \mu \) go to Step 1 else exit |

Solving Steps 1 and 2 mainly depends on the choice of \( d_T, L \) and \( T \), and the main challenge in this framework is to make realistic choices while keeping Step 1 and Step 2 computationally tractable. In the original ICP algorithm, \( d_T \) is simply the Euclidean distance between \( x_j \) and \( T(p_i) \), \( T \) is a rigid-body transformation, \( L \) is the null function and \( \gamma \) is set to zero. Step 1 is then solved setting \( A^{n+1}_{ij} \) to 1 if and only \( x_j \) is the closest point of \( T^n(p_i) \) in \( X \) and zero else (typically using a kd-tree). Solving Step 2 then consists in finding the rigid-body transformation best superposing the correspondences established during Step 1 in the least-squares sense, which can be done in a closed-form using for instance unit quaternions\(^17,18\) or the SVD\(^19\).

This iterative algorithm is very satisfying, as it is very simple and can be easily extended to cope with some specific types of non-linear transformations. However, the lack of a good initial transformation \( T \) makes it likely to be trapped in a local minimum, mostly because \( d_T \) is only based on the Euclidean distance between point coordinates. In the following we propose to include the unit normals in addition to the point coordinates to devise a new distance term \( d_T \). We show how an adequate transformation model and regularization term can be used to keep Steps 1 and 2 of the above-mentioned iterative scheme tractable.
3. CHOICE OF THE DISTANCE TERM

Instead of using the Euclidean distance between surface points $d_T(x_j, p_i) = ||x_j - T(p_i)||^2$ (as in the original ICP), we propose to use instead: $d_T(x_j, p_i) = ||x_j - T(p_i)||^2 + \lambda ||n_{x_j} - n_{T(p_i)}||^2$ where $n_{x_j}$ and $n_{T(p_i)}$ are the unit normal vectors at points $x_j \in X$ and $T(p_i) \in T(P)$, and $\lambda$ is a positive value weighing the influence of the normal-to-normal distance term over the point-to-point distance term.

It turns out that $n_{T(p_i)}$ can be expressed simply in terms of $n_p_i$ (unit normal vector at point $p_i \in P$) if additional hypotheses are made. We rely on the work by Kambhamettu and colleagues\cite{Kambhamettu} to derive a local differentiable orthogonal parametrization for the surface $P$. These constraints would be possible, at the cost of more complex computations. Relaxing this formula is independent of the chosen parametrization, provided it is orthogonal. The orthogonality of the parametrization and the hypothesis of a small deformation mostly allow to simplify the computations. Relaxing these constraints would be possible, at the cost of more complex computations.\cite{Kambhamettu}

As a summary, we design the distance term $d_T$ as follows:

$$d_T(x_j, p_i) = ||x_j - p_i - s_i||^2 + \lambda ||n_{x_j} - n_{p_i} - n_{p_i} \times \left( \frac{p_{u_i} \times s_{u_i}}{p_{u_i} \cdot p_{u_i}} + \frac{p_{v_i} \times s_{v_i}}{p_{v_i} \cdot p_{v_i}} \right)||^2$$

(3)

4. MINIMIZATION OF THE CRITERION

4.1 Derivatives of the Distance and Regularization Terms w.r.t. the Transformation

Before tackling the minimization of the whole ICP criterion, we need to further specify the deformation model and the associated regularization term. As suggested by Kambhamettu and colleagues,\cite{Kambhamettu} we propose to use a locally affine deformation model:

$$s(u_i, v_i) = u_i a_i + v_i b_i$$

We also propose to design a first-order regularization term that we define as:

$$L(T) = \sum_{(k, l) \in C_2(P)} ||u_k a_k + v_k b_k - (u_l a_l + v_l b_l)||^2$$

where $C_2(P)$ is the set of second-order cliques of $P$. In practice, the cliques are determined using a mesh: $(k, l) \in C_2 \Leftrightarrow$ there exists an edge between points $p_k$ and $p_l$. These choices are especially convenient, as they allow Step 2 of our ICP-type algorithm to stay tractable, as we will see in Section 4.2.2.

Below, we provide the derivatives of the distance and regularization terms with respect to the transformation parameters $a_i$ and $b_i$. Using $I$ as the identity matrix in $\mathbb{R}^{3 \times 3}$ and writing everything in block matrix form, we get:
4.2 Minimization of the Criterion

Using the choices made in Sections 3 and 4.1 for the distance and the regularization terms, the general criterion (1) becomes:

\[ \varepsilon(T, A) = \sum_{i=1}^{N_P} \sum_{j=1}^{N_S} A_{ij} \left( \frac{\partial}{\partial a_i} \frac{\partial}{\partial b_i} \right) \left( ||x_j - p_i - (u_i a_i + v_i b_i)||^2 + \lambda \right) \left( n_{x_j} - n_{p_i} \times \left( \frac{p_{u_i, x_s}}{p_{u_i, p_s}} + \frac{p_{v_i, x_s}}{p_{v_i, p_s}} \right) \right) \]

\[ + \beta \sum_{(k,l) \in C_2(P)} \left( ||u_k a_k + v_k b_k - (u_i a_i + v_i b_i)||^2 + \gamma \sum_{i=1}^{N_P} \sum_{j=1}^{N_S} A_{ij} \log(A_{ij}) \right) \]

We now proceed to the actual iterative minimization of this criterion, by first computing the optimal match matrix A when the transformation is given, and second computing the optimal transformation T for a given match matrix. These optima are computed as values where the derivatives vanish.

4.2.1 Step 1: Point Correspondences

Minimizing the criterion with respect to \( A_{ij} \) given the constraints \( \forall i, \sum_j A_{ij} = 1 \) and \( \forall (i,j), A_{ij} \geq 0 \) can be done in a straightforward way using Lagrange multipliers and give the following results:

\[ \forall (i,j), \frac{A_{ij}}{\alpha_j} = \exp \left( \frac{||x_j - p_i - (u_i a_i + v_i b_i)||^2 + \lambda \left( n_{x_j} - n_{p_i} \times \left( \frac{p_{u_i, x_s}}{p_{u_i, p_s}} + \frac{p_{v_i, x_s}}{p_{v_i, p_s}} \right) \right)^2}{-\gamma} \right) \]

where \( \alpha_j \) is such that \( \sum_j A_{ij} = 1 \). To reduce the computational cost of estimating the optimal transformation in Step 2, we choose to compute one-to-one correspondences from the fuzzy match matrix A, as proposed by Chui & Rangarajan. That is once we have estimated A we consider that \( p_i \) is matched with the virtual point \( x_i = \sum_j A_{ij} x_j \). Thus we consider that one-to-one correspondences between P and S are achieved in this first step.
4.2.2 Step 2: Transformation

After matching each point \( p_i \in P \) with a point \( \hat{x}_i \), we use the formulas (4), (5) and (6) obtained in Section 4.1 to minimize the criterion (7) with respect to \( a_i \) and \( b_i \), \( \forall i \in 1 \ldots N_P \). This yields a system of \( 6N_P \) equations:

\[
\forall i \in 1 \ldots N_P, \quad \left( (1 + \beta|V(i)|)u_i^2 + \lambda \frac{n_{p_i,n_{p_i}^T}}{v_i p_{v_i} v_{v_i}} \left( (1 + \beta|V(i)|)u_i v_i I + \lambda \frac{n_{p_i,n_{p_i}^T}}{v_i p_{v_i} v_{v_i}} \right) \right) \left( \begin{array}{c} a_i \\ b_i \end{array} \right) = \begin{array}{c} u_i[(\hat{\mathbf{x}}_i - \mathbf{p}_i)] + \beta \sum_{k \in V(i)} \{u_k a_k + v_k b_k\} \\ v_i[(\hat{\mathbf{x}}_i - \mathbf{p}_i)] + \beta \sum_{k \in V(i)} \{u_k a_k + v_k b_k\} - \lambda \frac{\{n_{s_i} - n_{p_i}\}_p}{p_{v_i} v_{v_i}} n_{p_i} \end{array} \right)
\]

The point \( \hat{x}_i \) does not necessarily belong to the surface \( X \). To compute \( n_{k_i} \), we simply project the point \( \hat{x}_i \) on the surface \( X \) and use the unit normal at this point in the formulas. The complete set of formulas obtained when keeping the entire fuzzy match matrix as computed in Step 1 using Equation (8) would be easy to derive, but we omit them for the sake of simplicity.

It could appear at first that the 6 equations obtained by differentiating with respect to the unknowns \( a_i \) and \( b_i \) are enough to compute them. It is not the case, because \( a_i \) and \( b_i \) also appear in the equations obtained when differentiating with respect to \( a_k \) and \( b_k \), when \( k \in V(i) \). Thus, in order to minimize the criterion with respect to the set \( (a_i, b_i), i \in 1 \ldots N_P \) we use an Iterated Coordinate Descent (ICD) algorithm, where we minimize with respect to each pair of parameters \( (a_i, b_i) \) in turn, while keeping all the other pairs fixed:

- **Step 2.0:** Initialize \((a_i, b_i) \forall i \in 1 \ldots N_P\)
- **Step 2.1:** \( \forall i \in 1 \ldots N_P \) \( \hat{a}_j, \hat{b}_j = \arg \min z((\hat{a}_0, \hat{b}_0) \ldots (a_i, b_i) \ldots (\hat{a}_{N_P}, \hat{b}_{N_P}), A) \)
- **Step 2.2:** If \( |\bar{e} - e| \geq \mu \) go to Step 2.0 else exit

Thus, each Step 2.1 consists of \( N_P \) linear systems with a simple \( 6 \times 6 \) matrix inversion, given by Equation (9). It can be shown that using the ICD algorithm does not change the convergence properties of the algorithm.

5. IMPLEMENTATION DETAILS

We look for a Monge patch representation at each point \( p \) of the surface \( P \). Such a representation is of the form \((x, y, z) = (u, v, f(u, v))\) and is an especially simple type of orthogonal parametrization. To obtain this representation, we first compute the unit normal at each point \( p \in P \) using the widely used VTK library
Rotation + TPS deformation

<table>
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<tr>
<th>Rotation in degrees</th>
<th>4</th>
<th>5</th>
<th>6</th>
<th>7</th>
<th>8</th>
<th>9</th>
</tr>
</thead>
<tbody>
<tr>
<td>Our algorithm (10 iterations)</td>
<td>0.52</td>
<td>0.56</td>
<td>0.63</td>
<td>0.72</td>
<td>0.83</td>
<td>0.98</td>
</tr>
<tr>
<td>Reference algorithm (10 iterations)</td>
<td>1.63</td>
<td>1.87</td>
<td>2.35</td>
<td>3.67</td>
<td>5.48</td>
<td>6.40</td>
</tr>
<tr>
<td>Reference algorithm (120 iterations)</td>
<td>1.04</td>
<td>1.10</td>
<td>1.29</td>
<td>1.64</td>
<td>4.32</td>
<td>5.48</td>
</tr>
</tbody>
</table>

Table 1. Mean error of registration for rotated, non-linearly deformed (using thin plate splines) ventricles using our algorithm (after 10 iterations) and using the reference algorithm (after 10 and 120 iterations).

We use the point coordinates and the unit normals of surface $X$ to build a kd-tree ($k=6$) for an efficient implementation of Equation (8). The use of a kd-tree is made possible by introducing a threshold function

$$
\rho_{\delta}(x) = \begin{cases} 
    x : |x| < \delta 
    \delta : \text{else} 
\end{cases}
$$

and thus replacing the original distance term by a term of the form $d_{T,\delta}(\cdot) = \rho_{\delta}(d_T(\cdot))$.

It can be shown that this change does not affect the convergence properties of the algorithm. It allows us to use range queries in the kd-tree to compute the values $A_{ij}$ and thus reduce the time complexity of the algorithm. Moreover, the matrix $A$ becomes sparse, reducing the space complexity. Another positive side effect of this threshold function is to make the algorithm less vulnerable to outliers. Note that when $\gamma$ becomes close to 0, Equation (8) boils down to a simple nearest neighbor query.

6. EXPERIMENTS AND RESULTS

As outlined in the introduction, the convergence of the classical ICP algorithm depends on the initial position of the surfaces: badly aligned surfaces in the first place may trap the algorithm in a local minimum. We show in this section that our algorithm may help to avoid this kind of problems. We generate the data in the following way:

- We segment several lateral ventricles from 3T T1-weighted brain isotropic MRI data using itk-SNAP (http://www.itksnap.org), which gives surfaces $X$ of about 10,000 points.
- We generate $P$ from $X$ by applying 1) a random thin plate spline transformation (to simulate such a transformation, we select a set of 8 landmarks on $X$ and randomly move them independently in a sphere of radius 20 mm around their initial position) and then 2) rotations with increasing angles. This gives us ground truth pairs $(X, P)$.
- We register each such obtained pair using 1) our algorithm (with the following parameters: $\beta = 50, \gamma = 0, \lambda = 1000$) and 2) a reference algorithm (where $\lambda = 0$, i.e. only the point coordinates are used to register $X$ and $P$).
- We evaluate the registration error by computing the mean distance between homologous points after registration. The values shown in Table 1 are the mean of these mean errors over the different ventricle pairs and results on one of the pairs are displayed on Figure 1.

The conclusions of these preliminary experiments are twofold: 1) the accuracy of our algorithm is much higher and 2) its convergence is much faster compared to the reference algorithm.

7. DISCUSSION

In this paper we have presented a novel algorithm for non-linear registration of surfaces. We have shown how to include the unit normals in addition to the point coordinates in an ICP-type criterion and we have shown how to minimize it in a simple way when a special type of small deformations is assumed between the two surfaces. Numerous adaptations and variants proposed in the literature for the original ICP algorithm could be applied in a straightforward way to our algorithm, such as the use of a prior affinity distance term as proposed by Combès.
Future work could also consist in adding a term based on the curvature in the criterion, using another set of formulas derived by Kambhamettu and colleagues. We have presented a set of preliminary experiments, and extending the validation dataset would be necessary. Evaluating different sets of parameters and different multiresolution or multiscale schemes would also be needed. At last, our algorithm needs to be compared with others, especially with landmark-free approaches and with other methods that do not resort to point-to-point correspondences.

REFERENCES


