

# Medical Image Analysis - Tensor B-Spline based warp exercise

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## 1 Introduction

The purpose of this exercise is to evaluate the transformation between two mid-sagittal brain MR slices (cf. figure 1) as a tensor B-spline. We will use a Gauss-Newton optimization to perform the registration.

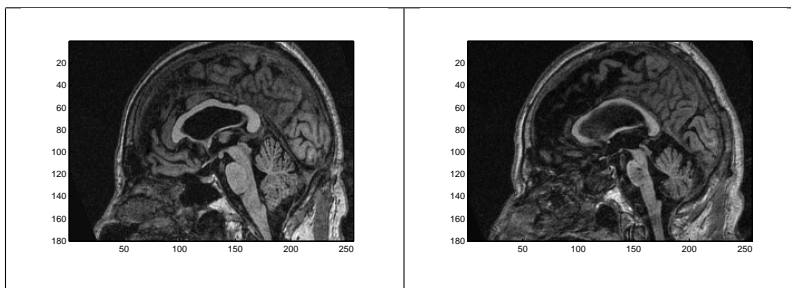


Figure 1: Two mid-sagittal brain MR slices (the left one is used as reference and the right one is the template)

The algorithm minimizes the dissimilarity (defined as the sum-of-squared differences) between the template and the reference as well as a regularizer (defined as the sum-of-squared B-Spline weights, to keep the transformation small between two iterations). The initial dissimilarity is shown in figure 2.

## 2 Algorithm

The algorithm minimizes the objective function  $J(w) = \|T(x + Qw) - R\|^2 + \alpha \|w\|^2$  where  $w$  is the spline weights vector,  $x$  represents the spatial identity,  $Q$  contains the values of the bidimensional spline at each knot,  $T$  is the template,  $R$  is the reference and  $\alpha$  is a parameter. The first term is the dissimilarity and the second the regularizer.

The number of iterations necessary for convergence is set experimentally. We found that, for the values of  $\alpha$  and the number of knots used, after 50 iterations the template hardly evolves any more.

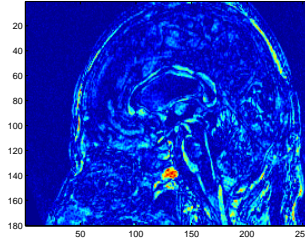


Figure 2: Initial dissimilarity

## 2.1 Initialization

```

1 % Load MR scans. "mr1" is the reference and "mr2" the
  template.
2 load mr.mat
3
4 % Size of images.
5 [R C]=size(mr1);
6
7 % Number of knots (total number is "p*p").
8 p=3;
9
10 % Parameter "alpha"
11 alpha=100;
12
13 % Number of iterations.
14 k_max=50;
15
16 % Definition of the unidimensional sets of knots. "s" is
    for the vertical and "t" for the horizontal. The knots
    are equidistantly distributed along their direction.
    3 additional knots are placed on top of the boundary
    knots so the spline will be equal to zero outside of
    the image.
17 s=augknt(linspace(0,R,p),3);
18 t=augknt(linspace(0,C,p),3);
19
20 % Construction of the splines, "B1" and "B2". We define "
    Q1" and "Q2" containing the values of the splines at
    each unidimensional knot. "Q" contains the values of
    the bidimensional spline at each knot.
21 B1=spmak(s,eye(p));
22 B2=spmak(t,eye(p));

```

```

23 Q1=fnval(B1,1:R)';
24 Q2=fnval(B2,1:C)';
25 Q=kron(speye(2),kron(Q2,Q1));
26
27 % "x1" and "x2" contain all the image coordinates with no
    transformation.
28 [x1 x2]=ndgrid(1:R,1:C);
29
30 % The initial value of the spline weights vector "w" is
    0.
31 w=zeros(size(Q,2),1);
32
33 % Main loop. The loop keeps going on until "stop" equals
    "true". The number of iterations is stored in "k".
34 stop=false;
35 k=0;
36 while ~(stop)

```

## 2.2 Transformation update

The coordinates transformation is defined as the identity plus the displacement:  $y(x) = x + Qw$  where  $Qw$  is the displacement given by the spline for the current weights.

```

1 % Computation of the transformation
2 y=[x1(:); x2(:)]+Q*w;
3
4 % "y1" contains the horizontal component of the
    transformation and "y2" the vertical component.
5 y1=reshape(y(1:end/2),R,C);
6 y2=reshape(y(end/2+1:end),R,C);

```

## 2.3 Transformed image and its spatial derivatives computation

```

1 % "Ty" is the transformed image.
2 Ty=interp2(mr1,y2,y1,'linear',0);
3
4 % "GradTy" is a matrix composed of two blocks. The left
    one is a diagonal matrix which diagonal is the
    vertical gradient. The right one is the equivalent for
    the horizontal gradient.
5 [GradTy2 GradTy1]=gradient(Ty);
6 GradTy=[spdiags(GradTy1(:),0,R*C,R*C) spdiags(GradTy2(:)
    ,0,R*C,R*C)];

```

## 2.4 Minimization of the objective function

We compute the new value of  $w$  as  $w + \Delta w$  with  $\Delta w$  minimizing  $J(w + \Delta w)$  by solving the equation  $(A^T A + \alpha I) \Delta w = A^T (R - T(y)) - \alpha w$  where  $A = \nabla T(y) Q$ .

```

1 % Computation of "A".
2 A=GradTy*Q;
3
4 % Minimization .
5 AtA=A'*A;
6 delta_w=(AtA+alpha*eye(size(AtA)))\'(A'*(mr2(:)-Ty(:))-
    alpha*w);
7
8 % Update of the value of "w"
9 w=w+delta_w;

```

## 2.5 End

```

1 % Update number of iteration
2 k=k+1;
3
4 % If the total number of iterations is reached , we stop
   the loop .
5 if k==k_max
6     stop=true ;
7 end
8
9 % End of the main loop .
10 end

```

## 3 Results

We tested the algorithm with  $3 \times 3$  knots,  $5 \times 5$  knots,  $7 \times 7$  knots for  $\alpha \in \{0.1; 1; 10; 100\}$ . Cf figure 3.

Although very similar, the best results are obtained for the higher number of knots, which seems logical as it implies a better adaptation to the complexity of the transformation. The results for higher values of  $\alpha$  seem slightly better as well: the zones of high dissimilarity (the bright zones) are smaller.

### 3.1 Conclusion

A tensor B-spline model for this non linear transform computed with a Gauss-Newton method is very efficient: the zones of strong dissimilarity are solved and the computing time is limited. However, the fact that the best results were obtained with a strong regularizer shows that in the present case, the initial dissimilarity was quite low and thus more important transform might not perform that well.

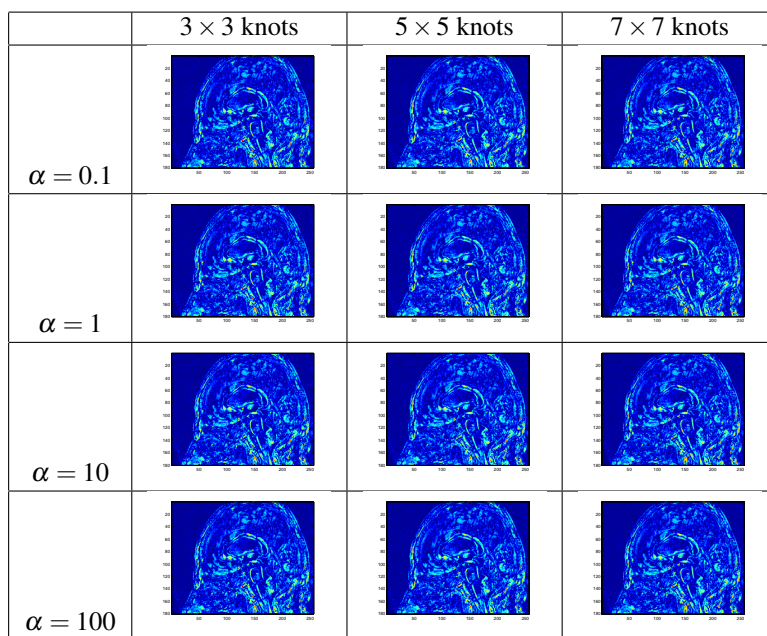


Figure 3: Resulting dissimilarities