

# **(rigid/affine) Image registration**

N. Duchateau – adapted from D. Sarrut + S. Rit

Master MISS / FSR-MED - Lyon, FR - 06/01/2022

## Course overview

### Theory:

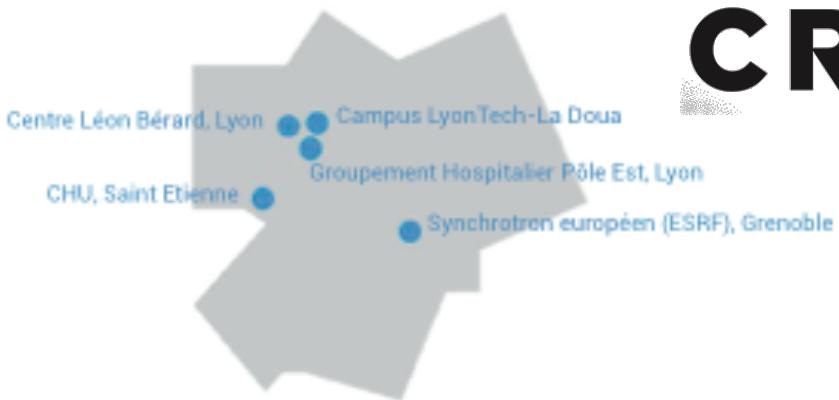
- 1) Rigid/affine + basics = 3h N. Duchateau
- 2) Non-rigid + advanced concepts = 3h + 1.5h D. Sarrut

### Labs:

- 1) Elastix / rigid-affine-nonrigid = 3h D. Sarrut
- 2) Motion estimation = 3h P. Clarysse
- 3) Quality check / temporal tracking = 3h N. Duchateau

Evaluation:      **Labs (x3) + Exam (1h30)**

## About (ND)



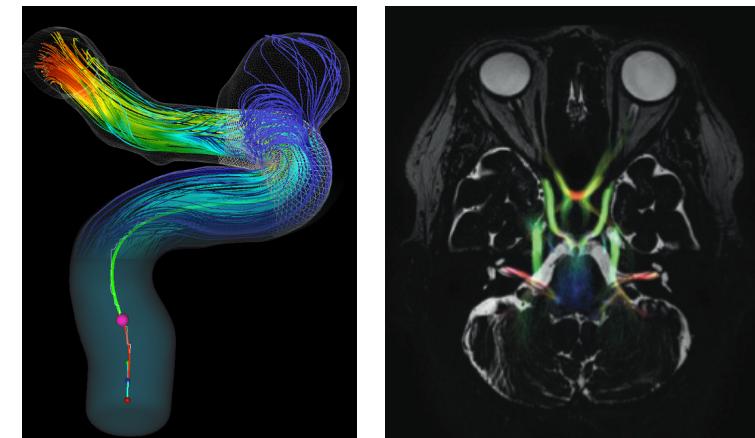
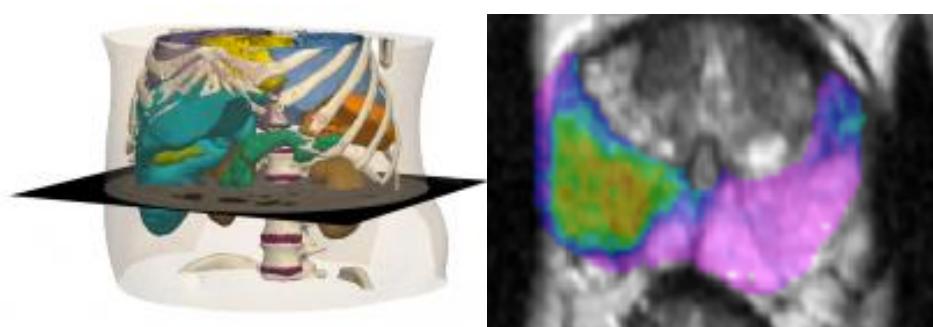
# CREATIS

## 4 équipes:

- Analyse d'images et modélisation
- Imagerie ultrasonore
- Imagerie tomographique + radiothérapie
- Imagerie RMN + optique

+ plate-forme d'imagerie multi-modale (PILOT)

+ plate-forme de calcul scientifique + VIP (Virtual Imaging Platform)



# Introduction

## Introduction

Definition =

“the process of aligning [images/meshes]  
so that corresponding *features* can easily be related”

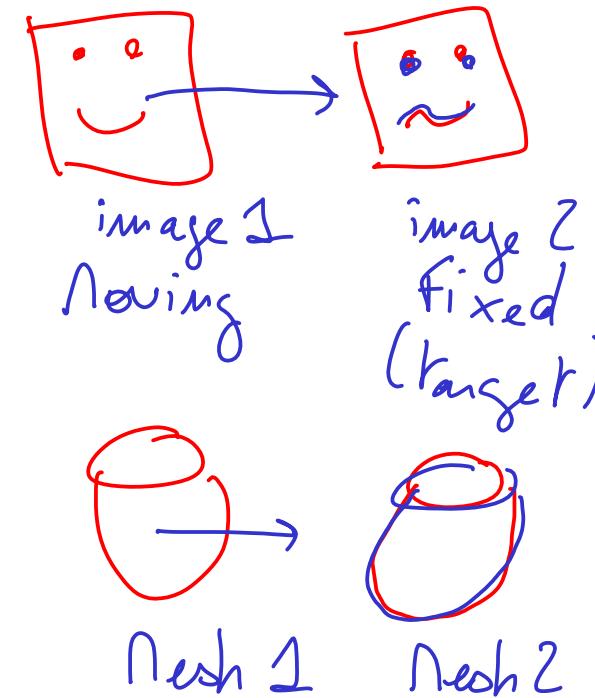
= registration      = *recalage*

= alignment

= geometrical correspondence

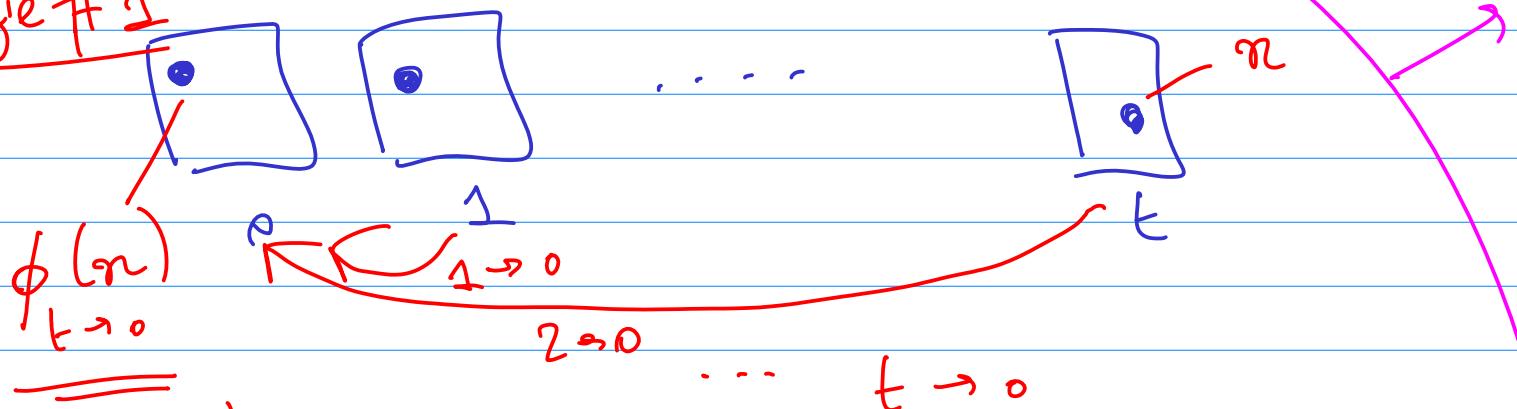
= matching

(= motion estimation ?)



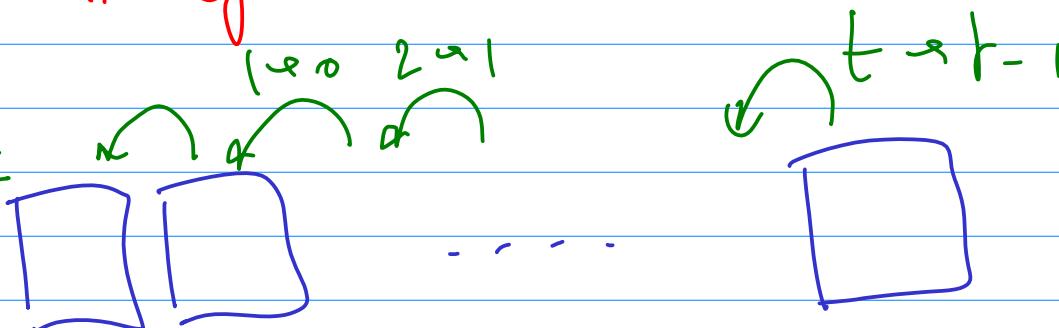
Sequence = estim. mer movement  
TP#3

Strategie #1



estimation  
de la transformation  $t \rightarrow 0$

Strategie #2



Compose

$$\phi_{1 \rightarrow 0} \circ \phi_{2 \rightarrow 1} \circ \dots$$

## Introduction

**Definition** = “the process of aligning [images/meshes]  
so that corresponding ***features*** can easily be related”

= registration

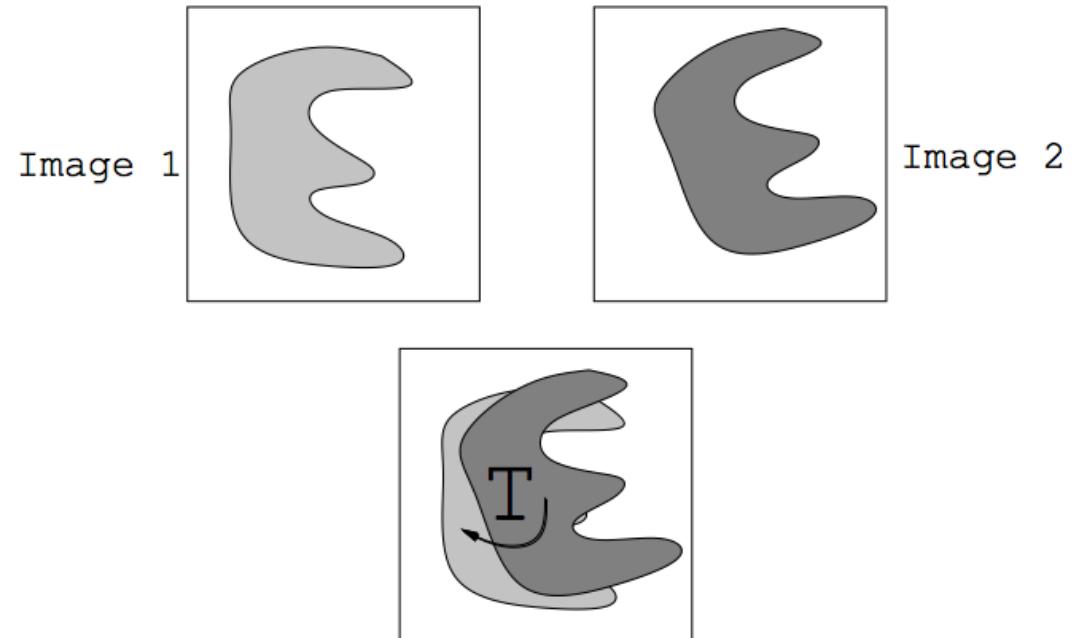
= alignment

= geometrical correspondence

= matching

(= motion estimation ?)

= find **spatial transformation  $T$**   
that matches two [images/meshes]



## Introduction

Definition = “the process of aligning [images/meshes]  
so that corresponding *features* can easily be related”

= registration

= alignment

= geometrical correspondence

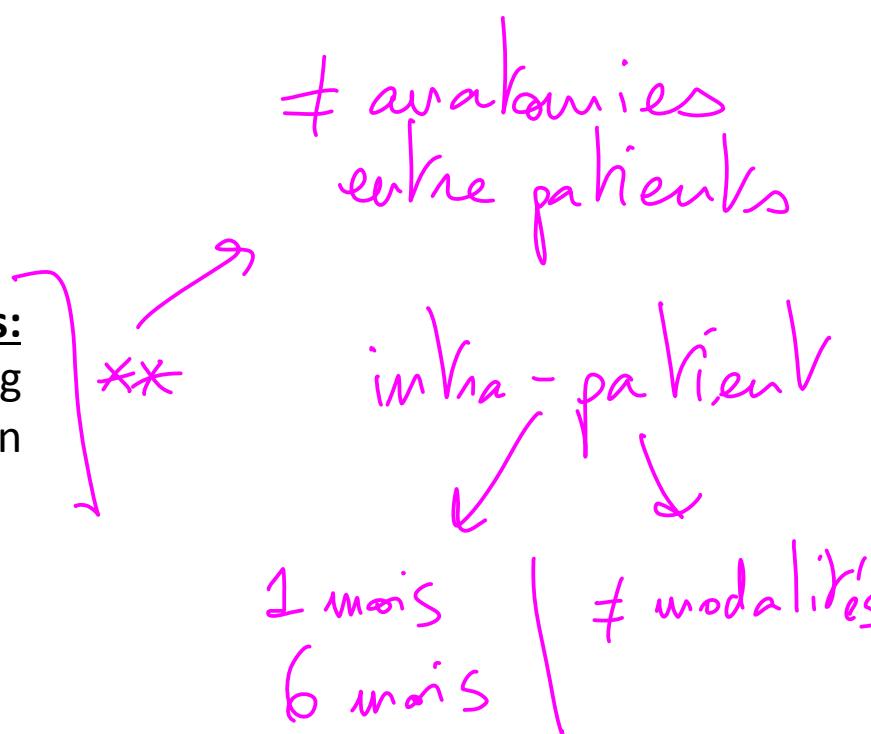
= matching

(= motion estimation ?)

## Important?

### Many applications:

- Medical imaging
- Computer vision
- ...



# Introduction

Definition = “the process of aligning [images/meshes]  
so that corresponding *features* can easily be related”

= registration

= alignment

= geometrical correspondence

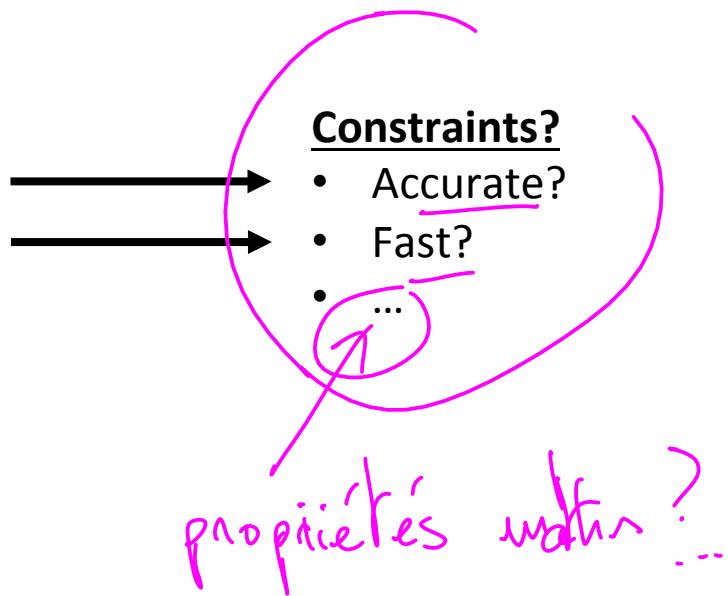
= matching

(= motion estimation ?)

## Important?

### Many applications:

- Medical imaging
- Computer vision
- ...



## Introduction

Definition = “the process of aligning [images/meshes]  
so that corresponding *features* can easily be related”

= registration

= alignment

= geometrical correspondence

= matching

(= motion estimation ?)

## Important?

### Many applications:

- Medical imaging
- Computer vision
- ...

### Constraints?

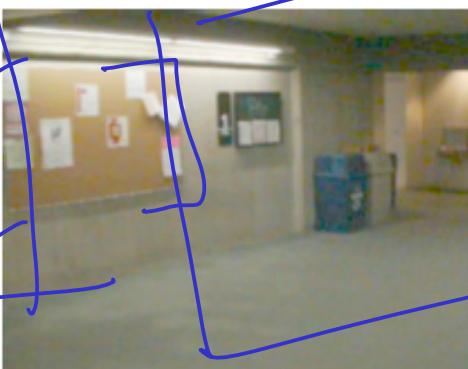
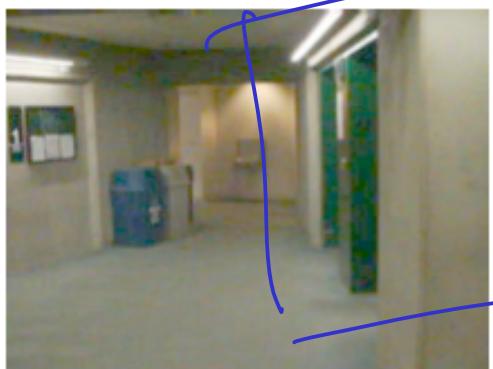
- Accurate?
- Fast?
- ...

### Basic elements used in many applications:

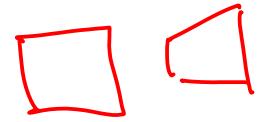
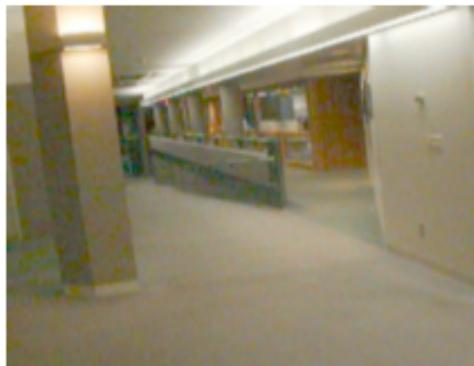
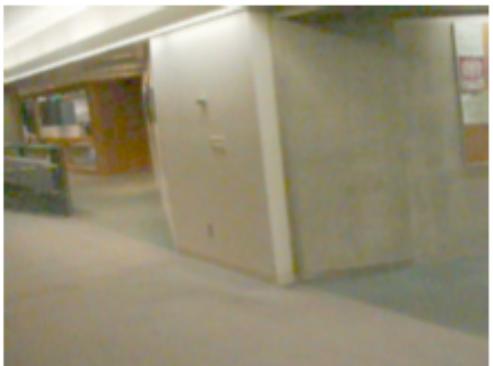
- Interpolation
- Similarity measure
- Optimization
- ...

## Examples

## Panoramic photography

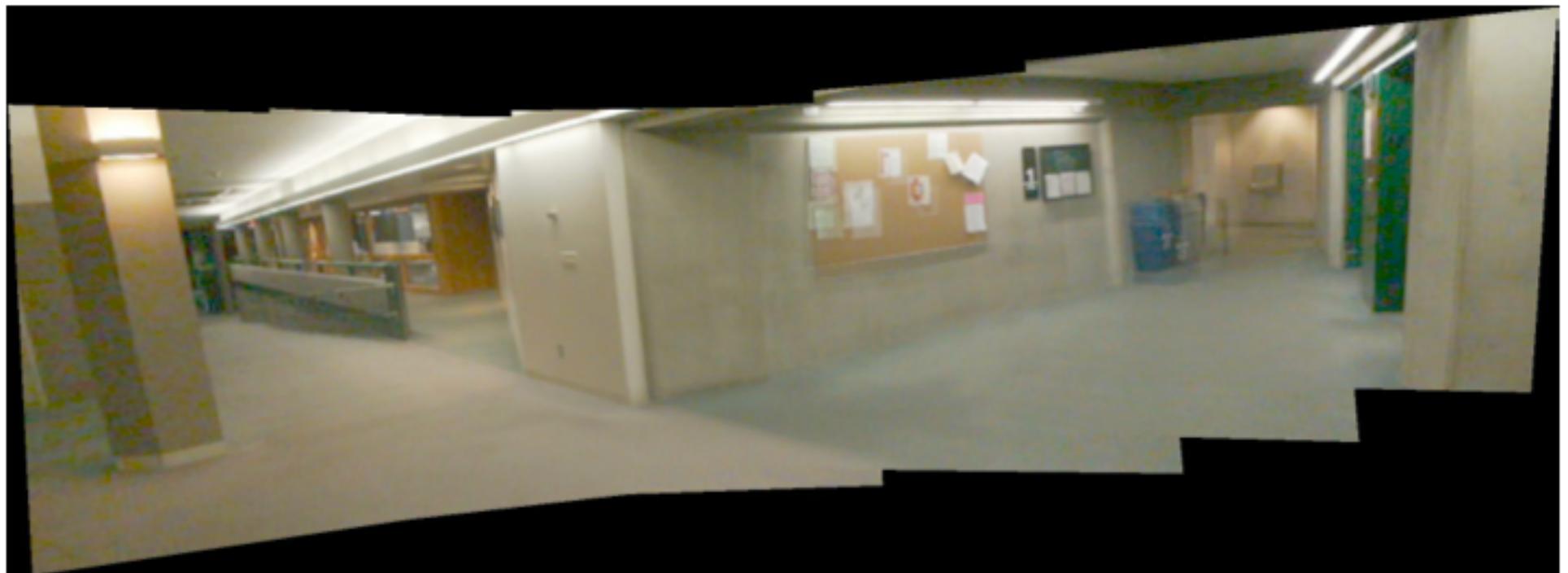


T simple:  
rotation  
translation  
affine



## Examples

### Panoramic photography



## Examples

### Satellite picture (e.g. GoogleMaps)

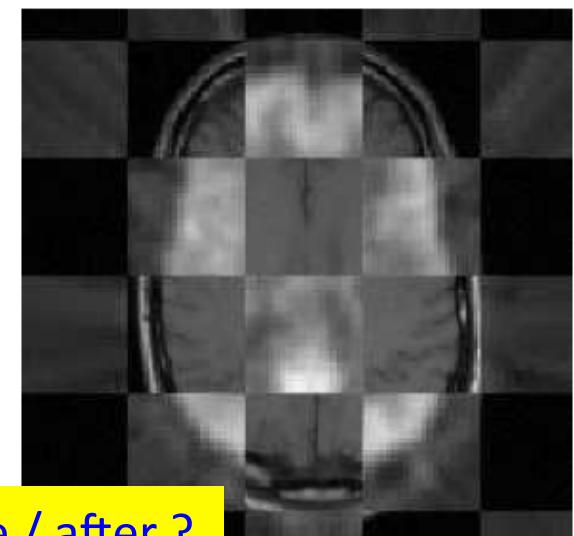
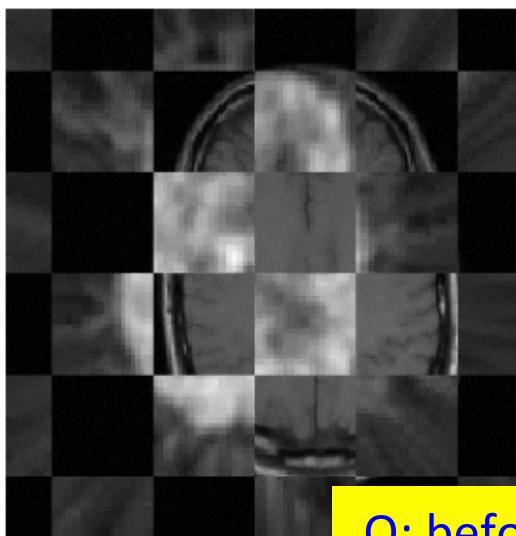
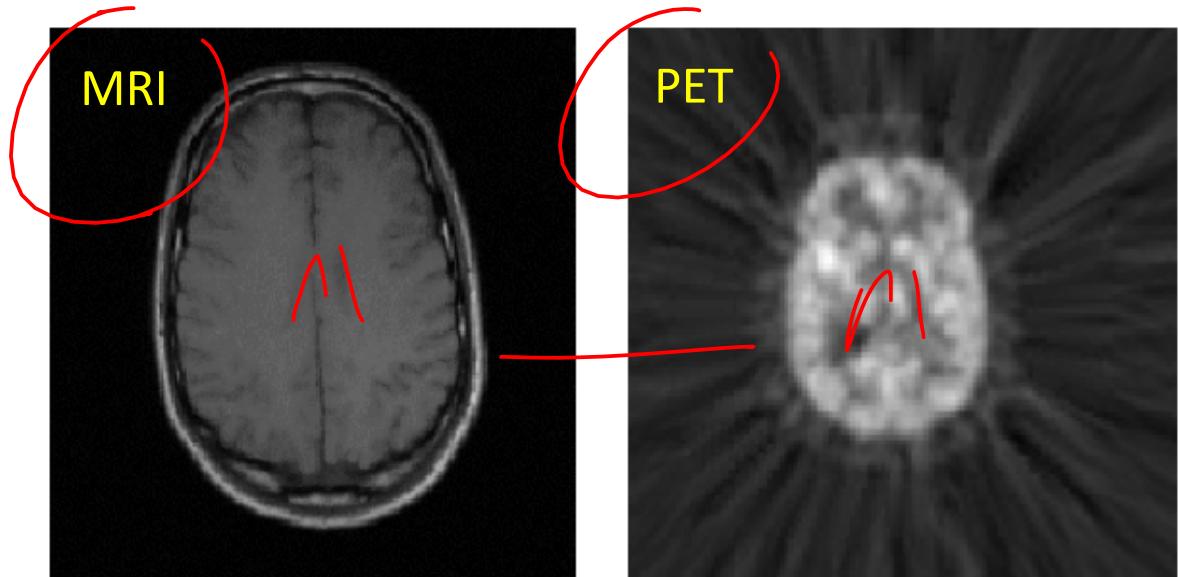


## Examples

### Multi-modal image fusion

T linéaire?  
(si c'est le

même patient)

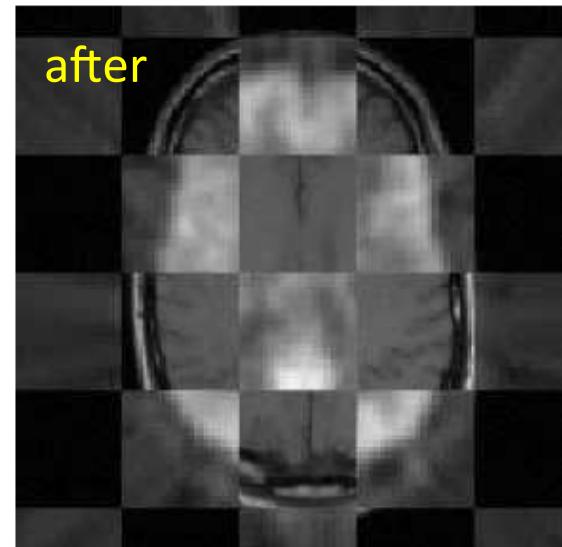
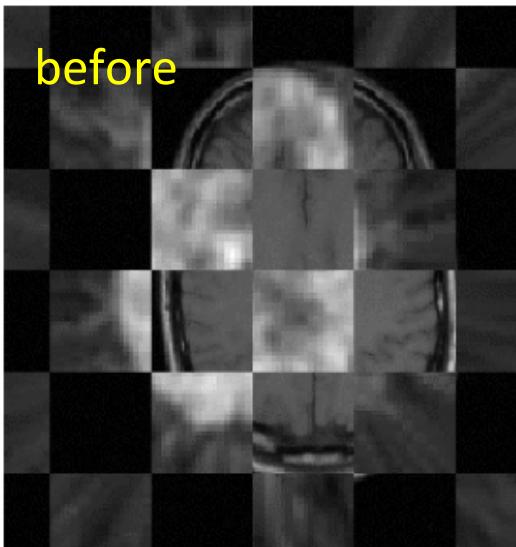
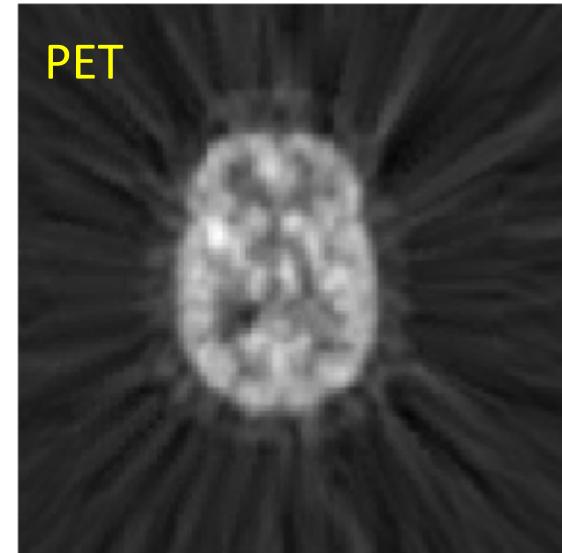
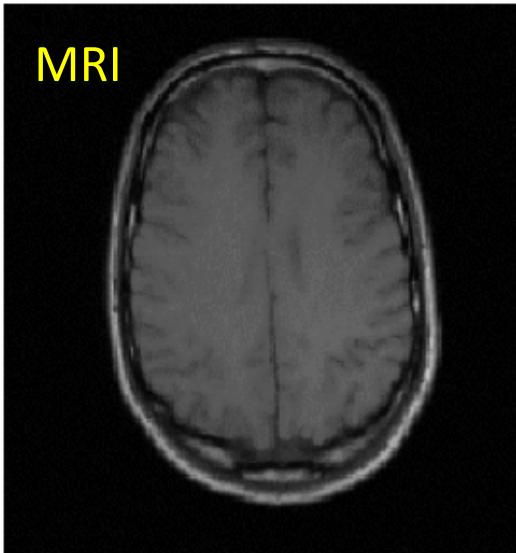


Q: before / after ?

## Examples

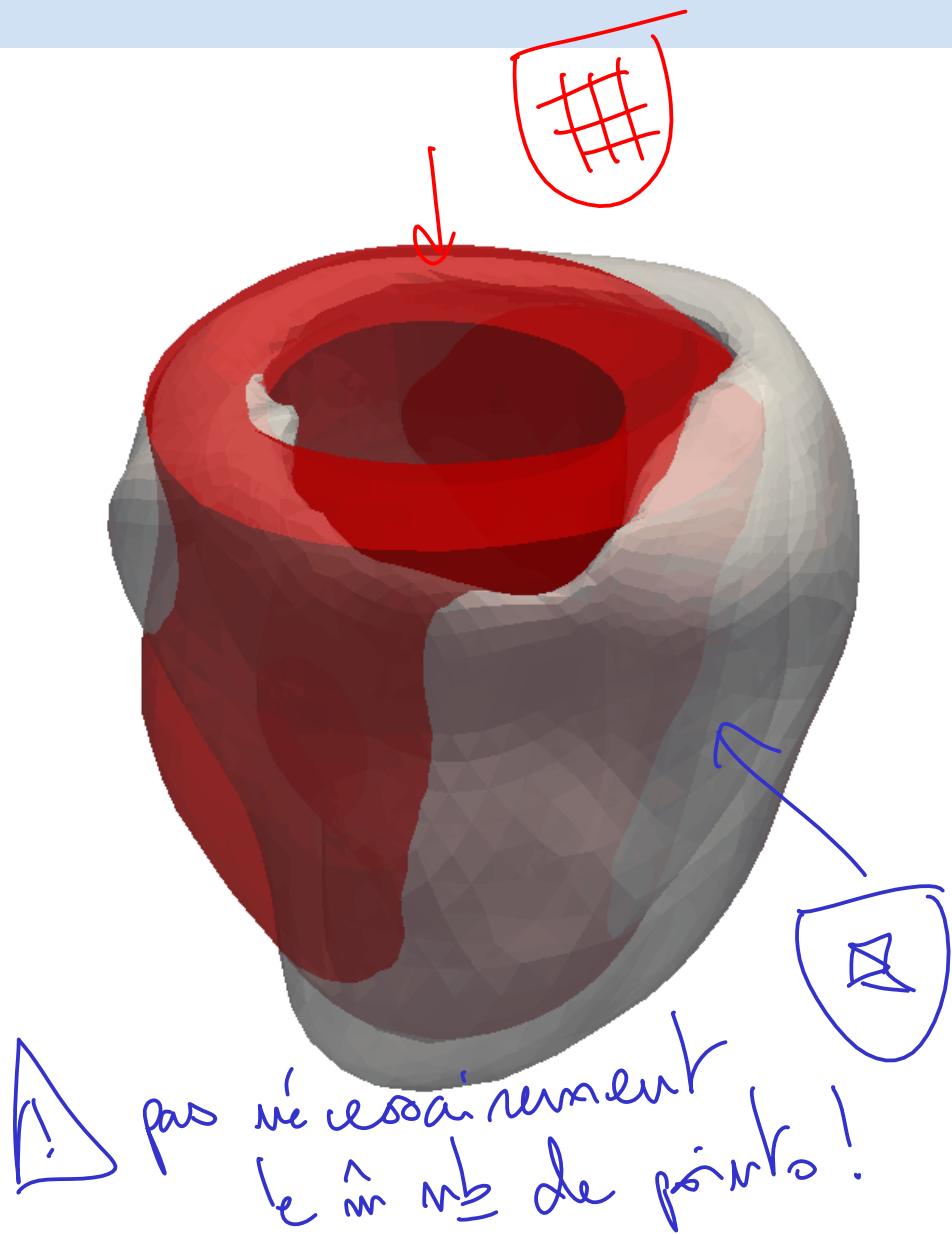
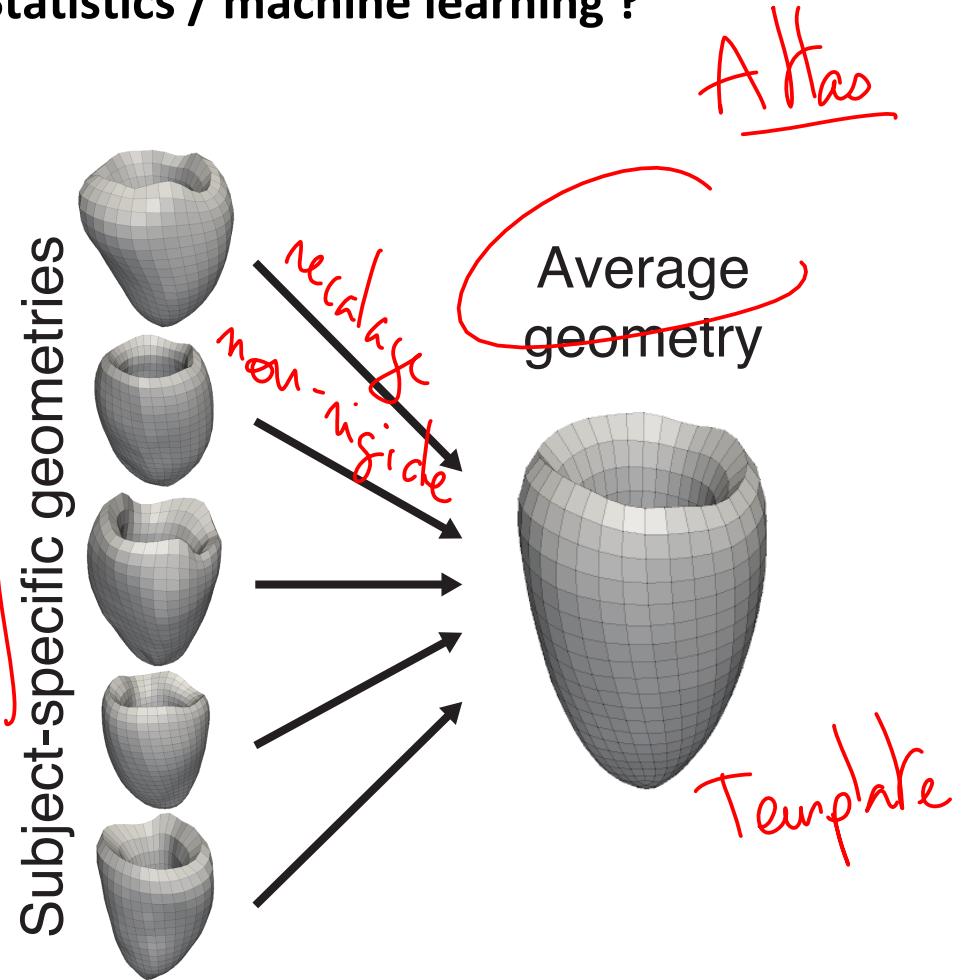
### Multi-modal image fusion

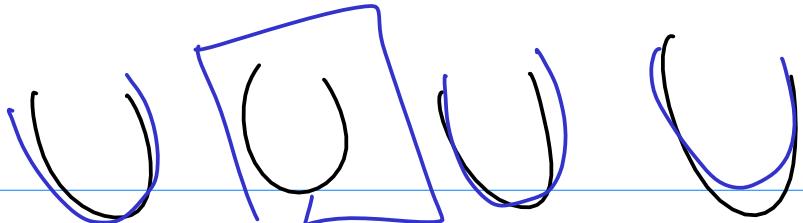
Crucial for accurate comparisons !!!



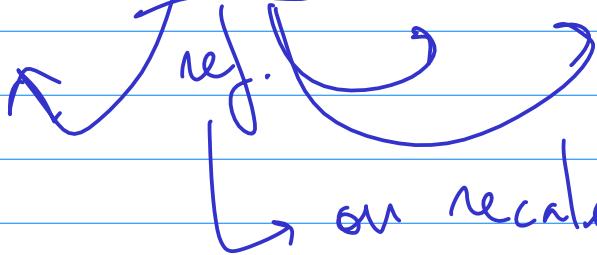
## Examples

Statistics / machine learning ?





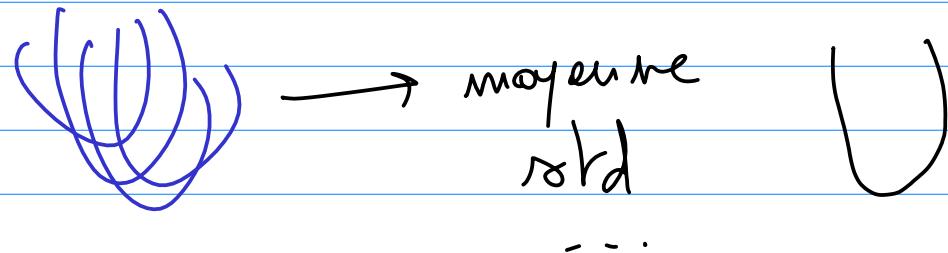
2 stratégies



on recalcule ~~af~~ sur les autres,

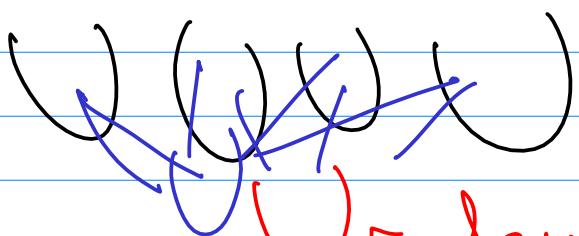
dans mb de points + conséq.

option #1



---

option #2



frame "centrale"

stats - own  
transf<sup>o</sup>  
+ process[er] i[eratif].

## Examples

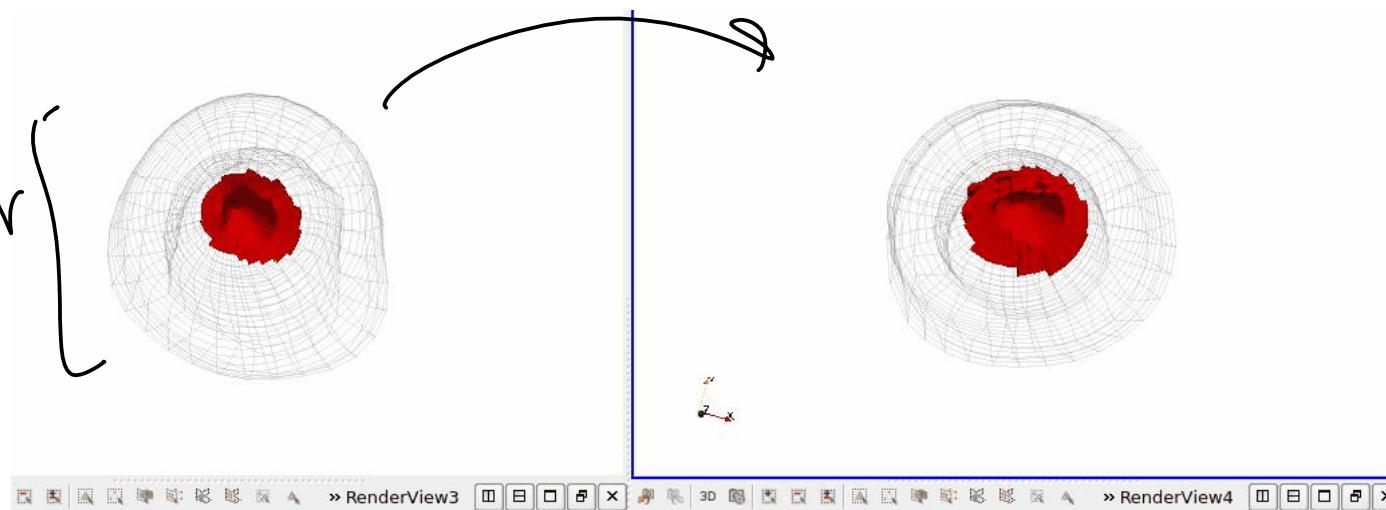
Statistics / machine learning ?

recalage +  
transport de l'information

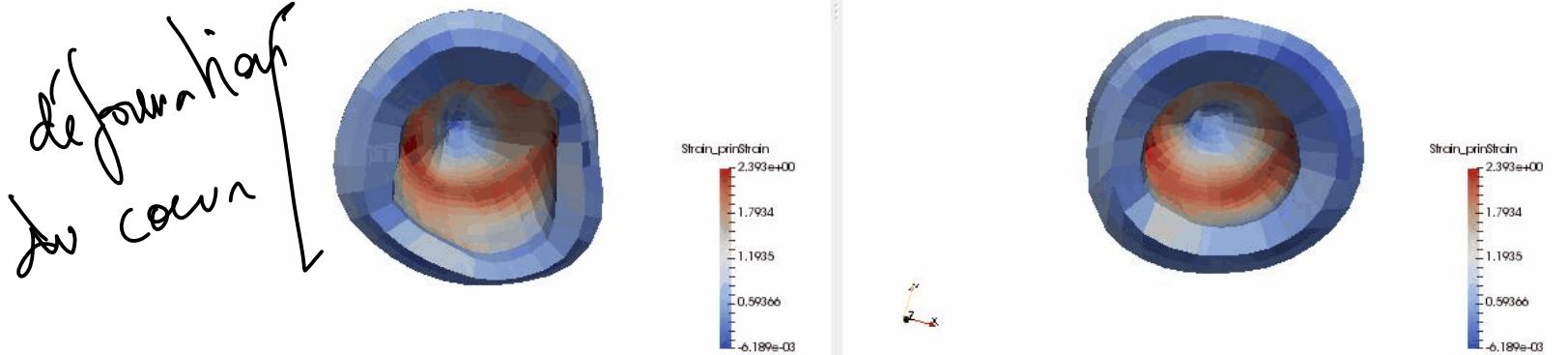
before

after

injuck



deformation  
de courb

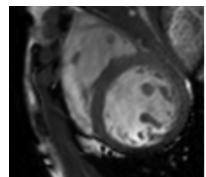


## Examples

## Motion analysis / tracking

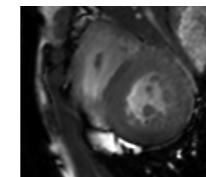
cf. TP3...

Registration [t,0]

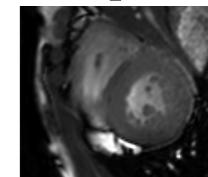


t=0

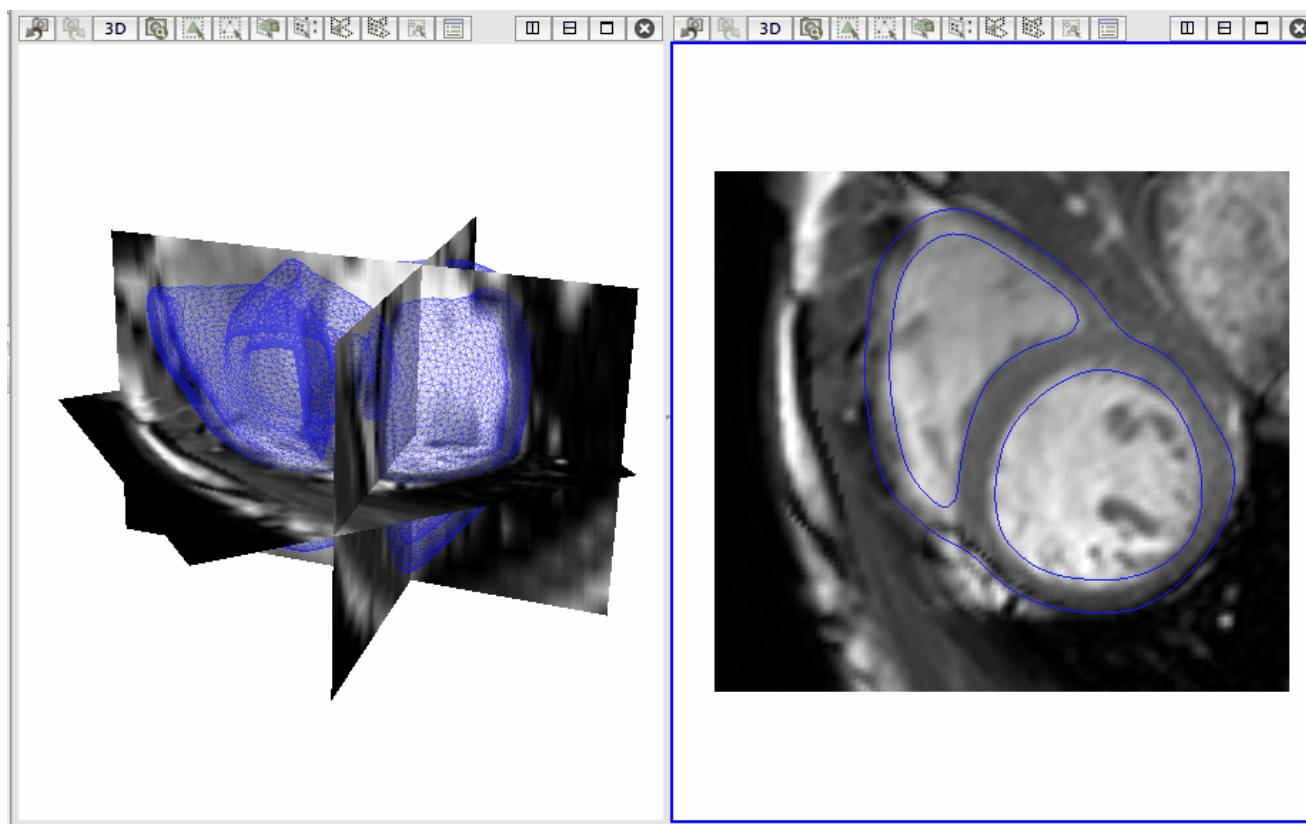
...



t-1



t



## Summary

- Find an “**optimal**” transformation
  - **Inputs = images/meshes**... may differ in:
    - Resolution
    - Time (longitudinal changes)
    - Space (2D / 2D+t / 3D / 3D+t)
    - Modality (MRI, US, CT, PET, ...)
    - Subject
    - ...
- + many possible output transformations



= many algorithms !!!

## Tools

- Code: ITK / VTK (C++), ...

images      meshes



Python

- Toolboxes: Elastix, Deformetrica, ...

TP#1      Elastix  
altavos



- Visualization: Paraview, ...



TP#3

## Formalism

images / maillages

Q:       $I = ?$   
           $J = ?$   
           $T = ?$   
           $S = ?$

$$\hat{T} = \underset{T \in \mathcal{T}}{\arg \max} S(I, J, T)$$

meilleure  
transf. candidates

Similarité ?  
(+ régularisation)

## Formalism

Q:  
 $I = ?$   
 $J = ?$   
 $T = ?$   
 $S = ?$

$$\hat{T} = \arg \max_{T \in \mathcal{T}} S(I, J, T)$$

## Notation

- $I$  = reference / fixed image
- $J$  = moving / floating image
- $T$  = transformation
- $\mathcal{T}$  = search space
- $S$  = similarity measure
- $\arg \max$  = optimization
- $\hat{T}$  = solution

faucille des possibles

## Digital images

- Size: nb pixels + spacing
- Pixel values: scalar (CT) / vector (RGB) / matrix (diffusion MRI)
- Colorscale / dynamic range
- Coordinates system: origin + orientation

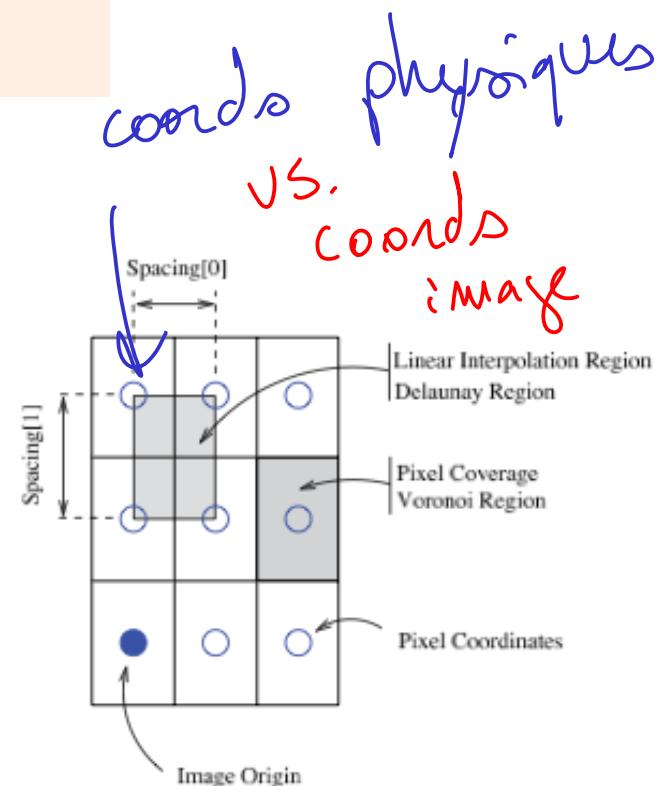
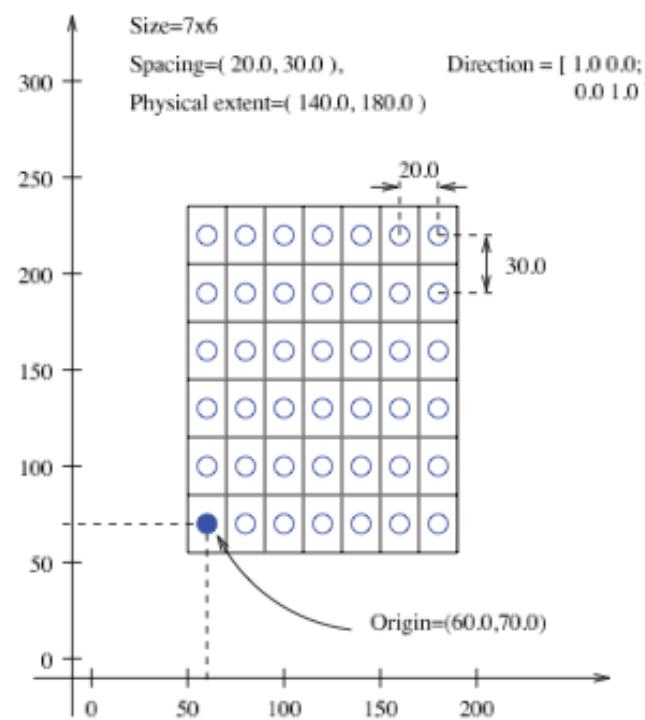
*m'solution**3x3 Tensor*[ItkSoftwareGuide.pdf](#)

Figure 4.1: Geometrical concepts associated with the ITK image.

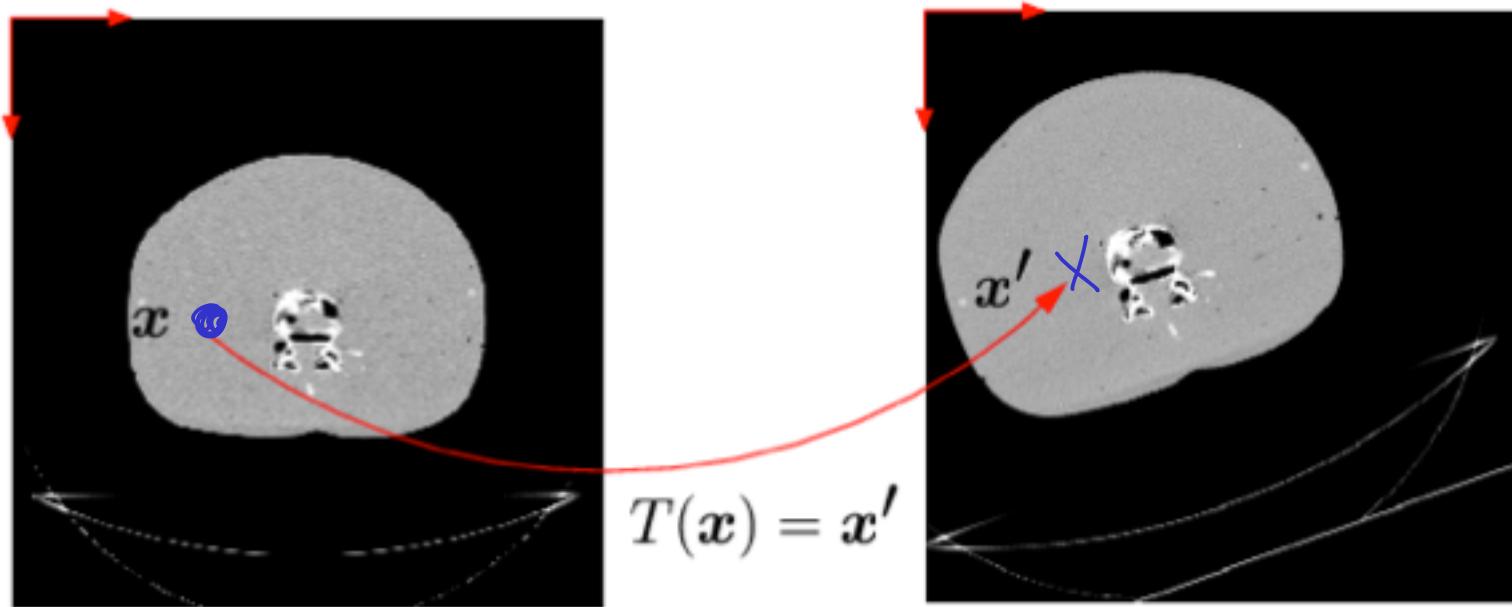
## Transformations

$$\hat{T} = \arg \max_{T \in \mathcal{T}} S(I, J, T)$$

## Notation

- $I$  = reference / fixed image
- $J$  = moving / floating image
- $T$  = transformation
- $\mathcal{T}$  = search space
- $S$  = similarity measure
- $\arg \max$  = optimization
- $\hat{T}$  = solution

## Transformations



$$T(\mathbf{x}) = \mathbf{x}'$$

- Cartesian coordinates:  $\mathbf{x} = (\underline{x}, \underline{y}, \underline{z}, t)^T$
- Transformation:  $T : \mathbb{R}^4 \rightarrow \mathbb{R}^4$   
 $\mathbf{x} \rightarrow T(\mathbf{x}) = \mathbf{x}'$

## Transformations: search space

One element of  $T$  is a transformation  $\tilde{T}$ .

Depends on:

- **Degrees of freedom** = number  $n$  of parameters needed to describe  $T$   
= dimension of the optimization problem
- **Boundaries** of each parameter (if any).

Success conditioned by:

- The solution belongs to the search space  $T$
- The search space  $T$  is small enough vs. the input data (1 solution can be reached)

pb dans l'optm<sup>e</sup> = bien définir l'espace des possibles

## Transformations types

Linear:

translation, rotation

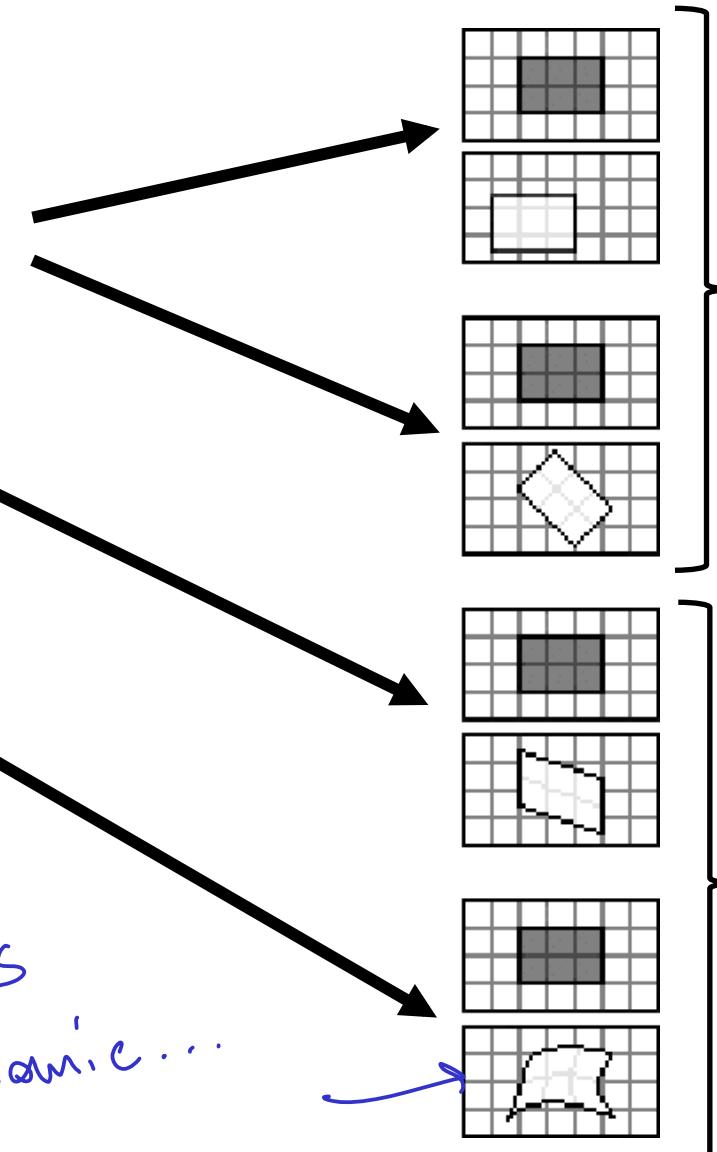
+ scaling

+ shearing (affine)

Non-linear:

- Deformable
- Elastic
- Fluid

proprites  
vs. anatomie ...



non - linear  
≠  
non - rigid

Rigid

Non-rigid

## Transformations types

Linear:

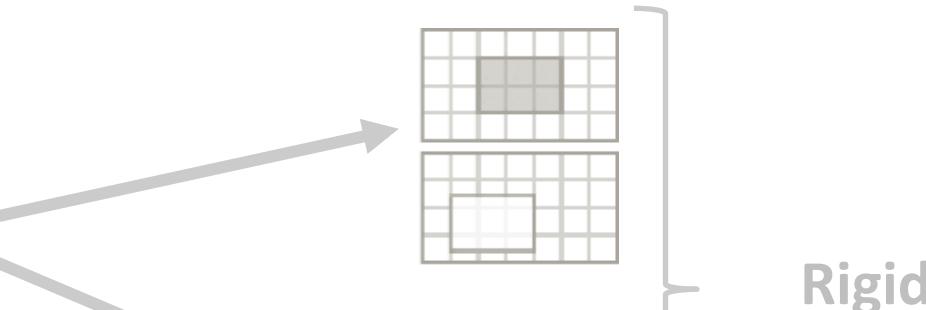
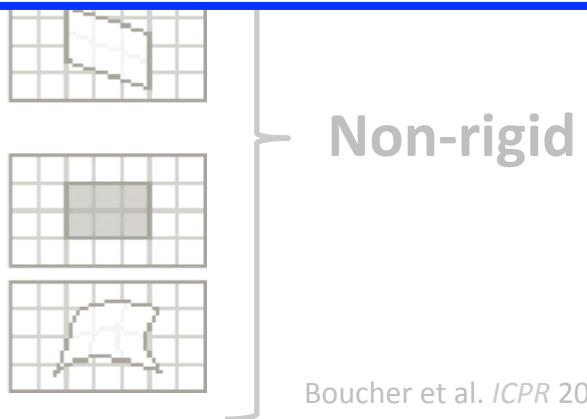
translation, rotation

+ scaling

+ shearing

Non-

- Deformation
- Elastography
- Fluid

**ALL are imposed MODELS...****Q: what if I want more freedom vs. the data?**

## Transformations types

Linear:

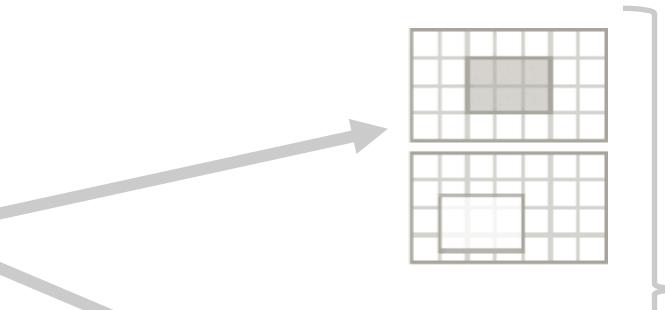
translation, rotation

+ scaling

+ shearing

Non-

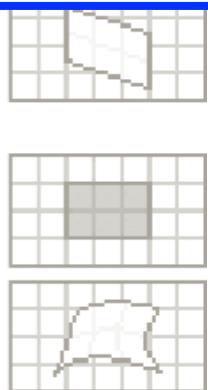
- Deformable
- Elastic
- Fluid



Rigid

**ALL are imposed MODELS...**

Q: what if I want more freedom vs. the data?

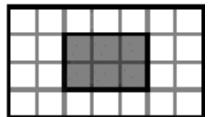
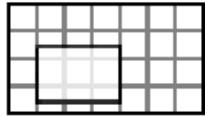
**... machine learning- based registration?***cov ~  
D. Smoother?*

Non-rigid

## Rigid transform



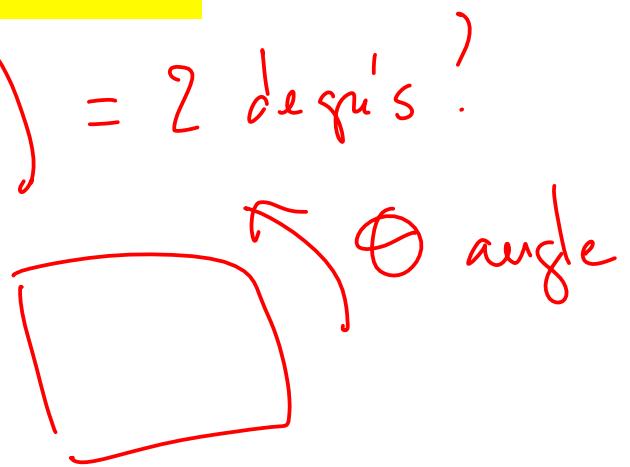
translation + rotation



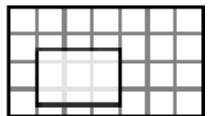
Q: degrees of freedom = parameters = ?

2D : transl :  $\begin{pmatrix} V \\ H \end{pmatrix}$  ) = 2 degr's ?

notation:



## Rigid transform



translation + rotation

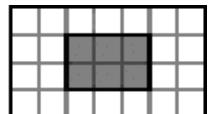
Q: degrees of freedom = parameters = ?

2D/2D: 3 parameters (2 translations + 1 rotation)

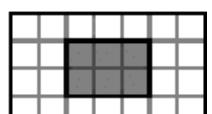
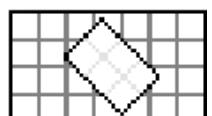
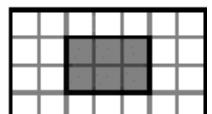
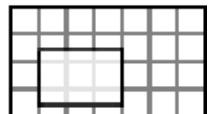
3D/3D: 6 parameters (3 rotations + 3 translations)

2D/3D: ~~6~~ parameters (3 rotations + 3 translations  
+ projection operator)

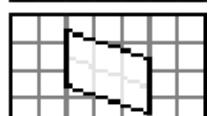
## Matrix representations



translation + rotation



affine



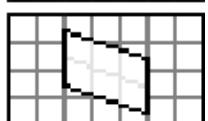
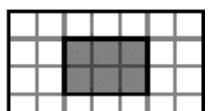
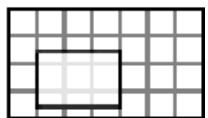
$$\begin{aligned} \text{Y} & \xrightarrow{\quad \quad \quad 2 \times 2 \quad \quad \quad 2 \times 1 \quad \quad \quad} \\ T_{\text{rigid}}(\mathbf{x}) &= \mathbf{R} \mathbf{x} + \mathbf{t} \\ T_{\text{affine}}(\mathbf{x}) &= \mathbf{A} \mathbf{x} + \mathbf{t} \end{aligned}$$

Q: dimension of  $\mathbf{x}$ ,  $\mathbf{R}$ ,  $\mathbf{M}$  and  $\mathbf{t}$ ?

$$\mathbf{n} = \begin{bmatrix} n_1 \\ n_2 \end{bmatrix} \quad \mathbf{t} = \begin{bmatrix} t_1 \\ t_2 \end{bmatrix}$$

$$\mathbf{R} = \begin{bmatrix} \cos \theta & -\sin \theta \\ \sin \theta & \cos \theta \end{bmatrix}$$

## Matrix representations



avec "biais"  
l'erreur de "bias"  
dans les réseaux  
neuro-synthétiques

\*\*\*  
"Homogeneous" coordinates:

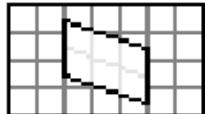
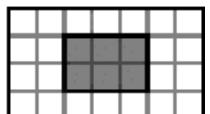
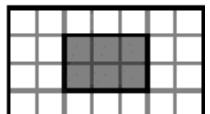
Q: dimension of  $x$ ,  $R$ ,  $M$  and  $t$  ?

(convenient for projection transforms)

$$\mathbf{x} = (x, y, z, 1) \rightarrow T_{\text{affine}}(\mathbf{x}) = \mathbf{M} \mathbf{x}$$

$$\mathbf{M} = \begin{bmatrix} R_{00} & R_{01} & R_{02} & t_0 \\ R_{10} & R_{11} & R_{12} & t_1 \\ R_{20} & R_{21} & R_{22} & t_2 \\ 0 & 0 & 0 & 1 \end{bmatrix}$$

## Matrix representations



Q: linear transform = ?

$$T_{\text{affine}}(\mathbf{x}) = \mathbf{M} \mathbf{x}$$

$$T(a \mathbf{x}_1 + \mathbf{x}_2) = a T(\mathbf{x}_1) + T(\mathbf{x}_2)$$

ex: Scaling:  $T(n) = \lambda n$

$$\begin{aligned} T(a n_1 + n_2) &= \lambda(a n_1 + n_2) \\ &= \lambda a n_1 + \lambda n_2 \\ &= a T(n_1) + T(n_2) \end{aligned}$$

## Matrix representations



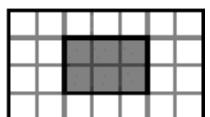
Q: linear transform = ?



$$T_{\text{affine}}(\mathbf{x}) = \mathbf{M} \mathbf{x}$$



$$T(a \mathbf{x}_1 + \mathbf{x}_2) = a T(\mathbf{x}_1) + T(\mathbf{x}_2)$$



Q: operations = ?



- Inverse:
- Composition:
- ...

*inverser de rotation  
en coordonnées homogènes*

$$\left. \begin{array}{l} T^{-1}(\mathbf{x}) = \mathbf{M}^{-1} \mathbf{x} \\ T_2( T_1(\mathbf{x}) ) = (T_2 \circ T_1)(\mathbf{x}) = \mathbf{M}_2 \mathbf{M}_1 \mathbf{x} \end{array} \right\} \quad (\text{if } \mathbf{M} \text{ is invertible})$$

Matrix representations: parameters  $\rightarrow \text{ex: } 3D$

$$\mathbf{M} = \begin{bmatrix} R_{00} & R_{01} & R_{02} & t_0 \\ R_{10} & R_{11} & R_{12} & t_1 \\ R_{20} & R_{21} & R_{22} & t_2 \\ 0 & 0 & 0 & 1 \end{bmatrix} \quad = 12 \text{ parameters to optimize !}$$

For registration:

- Option #1 = optimize the 12 parameters
- Option #2 = decompose in simpler transforms (rotation, translation, ...)

bien plus intéressant et plus complexe

Pause: important = dimensionnalité des objets:

Homogènes:

$$\vec{r} = \begin{pmatrix} x \\ y \\ z \\ 1 \end{pmatrix} \in \mathbb{R}^{4 \times 1}$$

$$T(\vec{r}) = M \vec{r}$$

$\mathbb{R}^{4 \times 1} \quad \mathbb{R}^{4 \times 4} \quad \mathbb{R}^{4 \times 1}$

$$M \in \mathbb{R}^{4 \times 4} \quad t \in \mathbb{R}^{3 \times 1}$$
$$= \begin{bmatrix} R \in \mathbb{R}^{3 \times 3} & | \\ 0 & 0 & 0 & 1 \end{bmatrix}$$

Concept de déterminant:  $\det$  = pour déformation

$$\det(R) \in \mathbb{R}$$

$R \in \mathbb{R}^{3 \times 3}$

Matrix representations: **translations**

$$\mathbf{M} = \begin{bmatrix} R_{00} & R_{01} & R_{02} & t_0 \\ R_{10} & R_{11} & R_{12} & t_1 \\ R_{20} & R_{21} & R_{22} & t_2 \\ 0 & 0 & 0 & 1 \end{bmatrix}$$

~~= 12 parameters to optimize !~~**2D: 2 parameters**

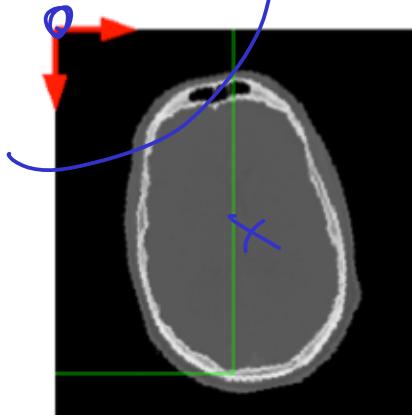
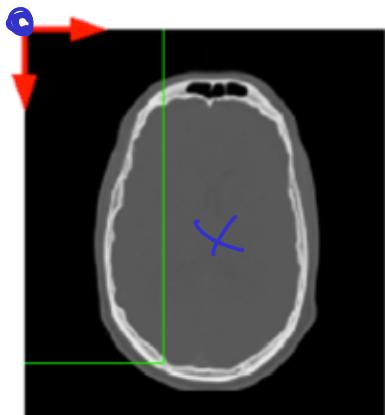
$$\mathbf{M} = \begin{bmatrix} 1 & 0 & t_0 \\ 0 & 1 & t_1 \\ 0 & 0 & 1 \end{bmatrix}$$

**3D: 3 parameters**

$$\mathbf{M} = \begin{bmatrix} 1 & 0 & 0 & t_0 \\ 0 & 1 & 0 & t_1 \\ 0 & 0 & 1 & t_2 \\ 0 & 0 & 0 & 1 \end{bmatrix}$$

Matrix representations: rotation

$$\mathbf{M} = \begin{bmatrix} R_{00} & R_{01} & R_{02} & t_0 \\ R_{10} & R_{11} & R_{12} & t_1 \\ R_{20} & R_{21} & R_{22} & t_2 \\ 0 & 0 & 0 & 1 \end{bmatrix}$$



= 12 parameters to optimize!

2D, around the origin: 1 parameter

$$\mathbf{M} = \begin{bmatrix} \cos \theta & -\sin \theta & 0 \\ \sin \theta & \cos \theta & 0 \\ 0 & 0 & 1 \end{bmatrix}$$

origin

## Matrix representations: rotation

$$\mathbf{M} = \begin{bmatrix} R_{00} & R_{01} & R_{02} & t_0 \\ R_{10} & R_{11} & R_{12} & t_1 \\ R_{20} & R_{21} & R_{22} & t_2 \\ 0 & 0 & 0 & 1 \end{bmatrix}$$

~~= 12 parameters to optimize!~~

**2D, around the origin: 1 parameter**

$$\mathbf{M} = \begin{bmatrix} \cos \theta & -\sin \theta & 0 \\ \sin \theta & \cos \theta & 0 \\ 0 & 0 & 1 \end{bmatrix}$$

Q: around another point?

$$\mathbf{p} = (p_0, p_1)$$

translation + rotation

$$\begin{bmatrix} 1 & 0 & p_0 \\ 0 & 1 & p_1 \\ 0 & 0 & 1 \end{bmatrix}$$

$$\begin{bmatrix} \cos \theta & -\sin \theta & 0 \\ \sin \theta & \cos \theta & 0 \\ 0 & 0 & 1 \end{bmatrix}$$

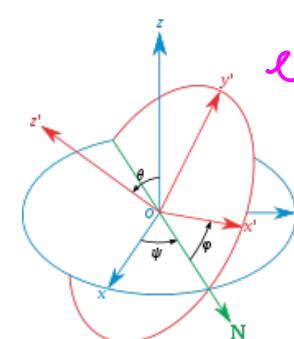
$$\begin{bmatrix} 1 & 0 & -p_0 \\ 0 & 1 & -p_1 \\ 0 & 0 & 1 \end{bmatrix}$$

## Matrix representations: 3D rotation

- Rotation = 3 x 3 matrix
- 3 degrees of freedom = 3 parameters → 3 angles
- Constraints  
Orthogonal matrix:  $\mathbf{R} \mathbf{R}^T = \mathbf{I}$  and  $\det(\mathbf{R}) = 1$   
(inverse:  $\mathbf{R}^{-1} = \mathbf{R}^T$ )

## Matrix representations: 3D rotation

- Rotation = 3 x 3 matrix
  - 3 degrees of freedom = 3 parameters
  - Constraints
  - Several decompositions (from 9 values to 3 parameters):
    - Euler angles
    - Axis angle
    - Quaternions
    - ...
- The parameters have a different meaning in each case
- Check illustrations  
➤ Q: pros vs. cons?
- comment bien définir ça ? ?
- ex: Euler angles



Intervalle :  $\det$

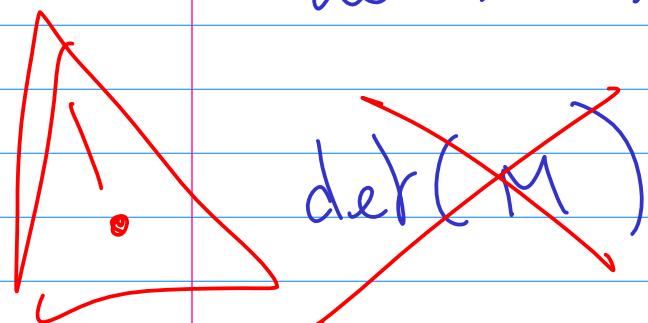
$$R = \begin{pmatrix} \cos \theta & -\sin \theta \\ \sin \theta & \cos \theta \end{pmatrix}$$

$$\det(R) = \cos^2 \theta - (-\sin^2 \theta)$$

\*\*

$$= 1$$

$\det$  : mesure de "déformation"



$\det(M)$

$\det(J_M)$

Sacobian

$$J = \begin{bmatrix} \frac{\partial f}{\partial x_1} & \dots & \frac{\partial f}{\partial x_n} \end{bmatrix} = \begin{bmatrix} \nabla^T f_1 \\ \vdots \\ \nabla^T f_m \end{bmatrix} = \begin{bmatrix} \frac{\partial f_1}{\partial x_1} & \dots & \frac{\partial f_1}{\partial x_n} \\ \vdots & \ddots & \vdots \\ \frac{\partial f_m}{\partial x_1} & \dots & \frac{\partial f_m}{\partial x_n} \end{bmatrix}$$

(2D)

$$\begin{bmatrix} \frac{\partial f_n}{\partial x} & \frac{\partial f_n}{\partial y} \\ \frac{\partial f_y}{\partial x} & \frac{\partial f_y}{\partial y} \end{bmatrix} \begin{bmatrix} \frac{\partial f_1}{\partial n_1} & \frac{\partial f_1}{\partial n_2} \\ \frac{\partial f_2}{\partial n_1} & \frac{\partial f_2}{\partial n_2} \end{bmatrix} \begin{matrix} f_1 \\ f_2 \end{matrix}$$

$$R n = \begin{bmatrix} \cos \theta & -\sin \theta \\ \sin \theta & \cos \theta \end{bmatrix} \begin{bmatrix} n_1 \\ n_2 \end{bmatrix} = \begin{bmatrix} \cos \theta n_1 - \sin \theta n_2 \\ \sin \theta n_1 + \cos \theta n_2 \end{bmatrix}$$

$$J_R = \begin{bmatrix} \frac{\partial f_1}{\partial n_1} = \cos \theta & -\sin \theta \\ \sin \theta & \cos \theta \end{bmatrix} = R !$$

Dans le cadre des rotations:  $J_R = R$

$$\det(J_R) = \det(R) = 1$$

on garde sa en tête ...

## Matrix representations: scaling

$$M = \begin{bmatrix} R_{00} & R_{01} & R_{02} & t_0 \\ R_{10} & R_{11} & R_{12} & t_1 \\ R_{20} & R_{21} & R_{22} & t_2 \\ 0 & 0 & 0 & 1 \end{bmatrix}$$

$$Mx \longleftrightarrow Rx + t$$

= 12 parameters to optimize!

3D: 3 parameters

$$R = \begin{bmatrix} s_x & 0 & 0 \\ 0 & s_y & 0 \\ 0 & 0 & s_z \end{bmatrix}$$

Rf notation

$$Rf = \begin{bmatrix} s_1 n_1 \\ s_2 n_2 \\ s_3 n_3 \end{bmatrix}$$

$\det(J_{\text{scaling?}})$

(20)

$$f(\alpha) = \begin{cases} s_1 n_1 \\ s_2 n_2 \end{cases}$$

$$J = \begin{bmatrix} \frac{\partial f_1}{\partial \alpha_1} & \frac{\partial f_1}{\partial \alpha_2} \\ \frac{\partial f_2}{\partial \alpha_1} & \frac{\partial f_2}{\partial \alpha_2} \end{bmatrix} = \begin{bmatrix} s_1 & 0 \\ 0 & s_2 \end{bmatrix}$$

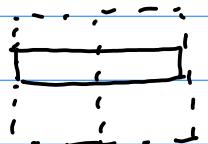
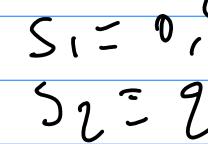
$s_1 = 1$   
 $s_2 > 1$

$\det(J) > 1$

ex:  

$s_1 > 1$   
 $s_2 = 1$

$\det(J) > 1$

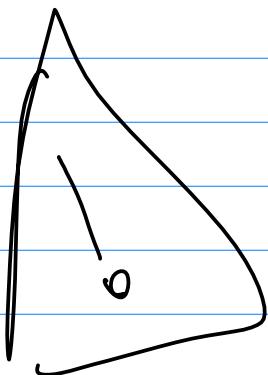
 

$s_1 = 0, s_2 = 2$

$\det(J) = 1$

$$\det(J) = s_1 s_2$$

intuition que  $\det(J) = \text{mesure de}$   
"déformation"



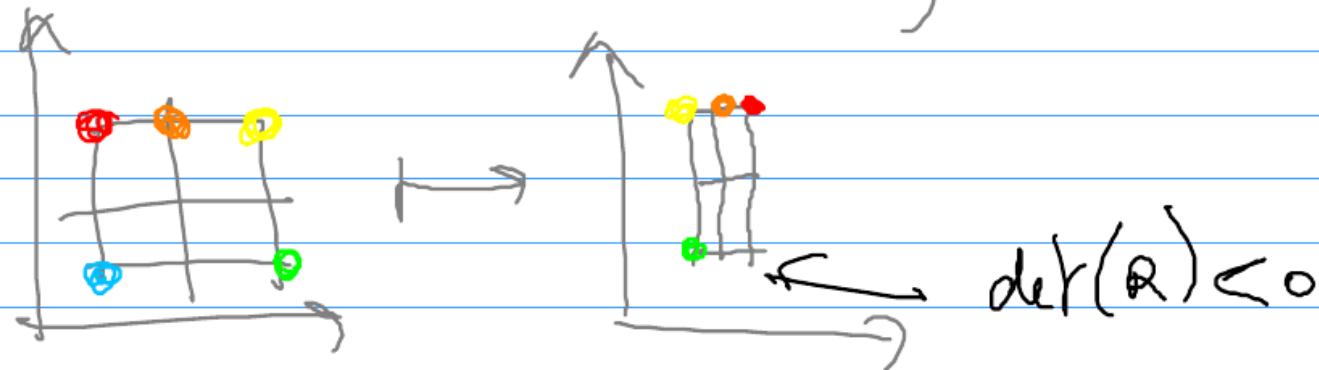
calculs, faits en un point précis !

Conclusion:  $\det(R)$

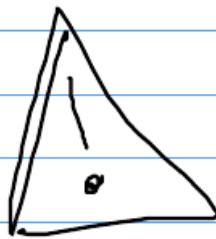
- $> 1$  grossit
- $= 1$  pareil (on peut translater)
- $< 1$  réduit

rotation

ex:  $\det(R) < 0$



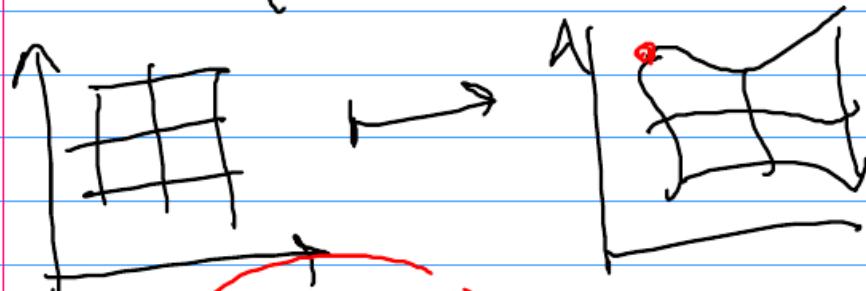
$\det(R)$  caractérise déformation



Transformations linéaires

$$T(\vec{v}) = M \vec{v}$$

Rq : Transformations non-linéaires



$$T(\vec{r}) = ?$$

pas de formulation matricielle

$$\det(R)$$

$$\det(?)$$

$$\det(J(\vec{r})) \leq 1$$

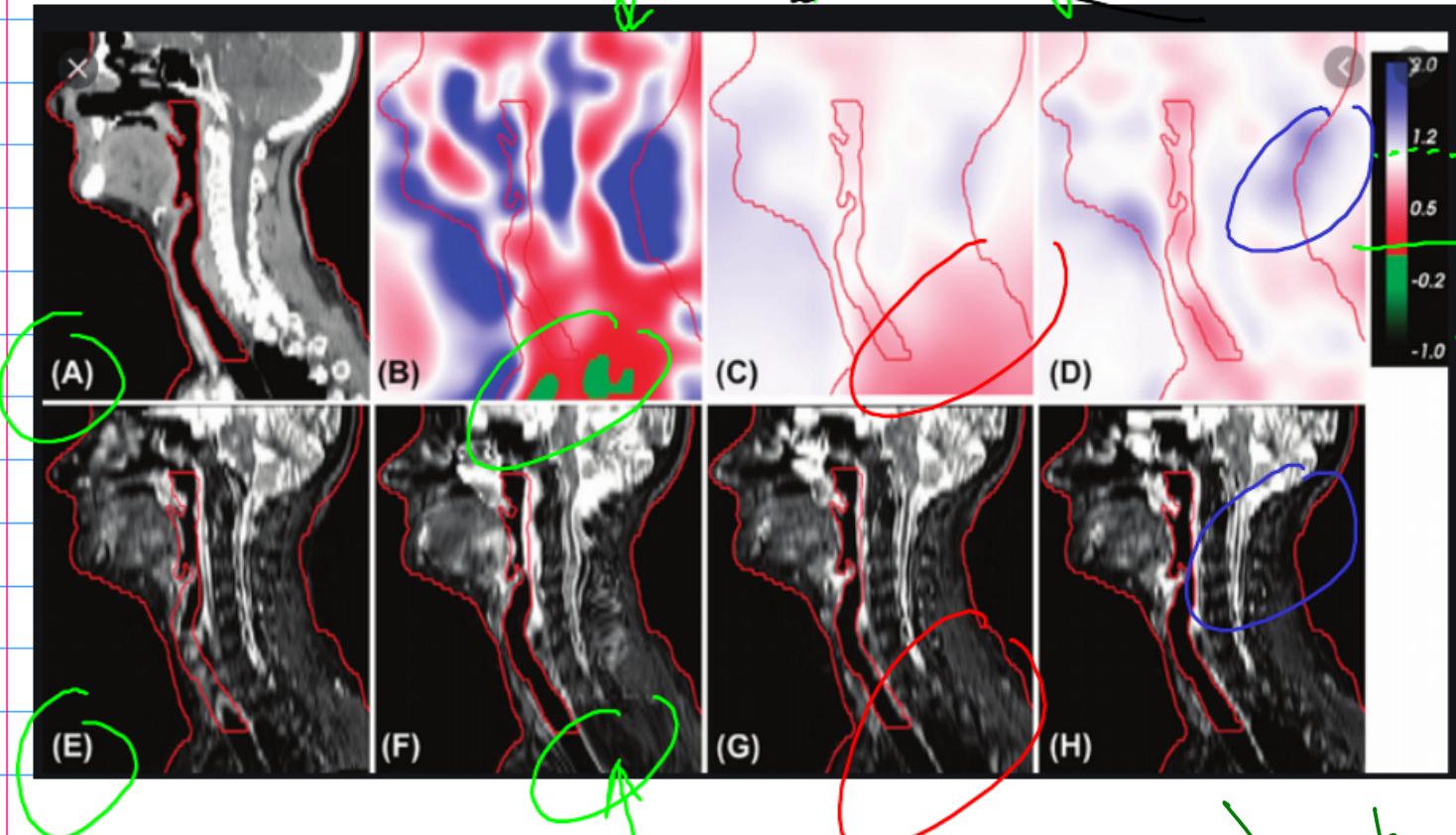
$$\text{Jacobiens} \begin{cases} > 1 \\ = 1 \\ < 1 \end{cases}$$

ex: 2D

$$J = \begin{bmatrix} \frac{\partial T_x}{\partial x} & \frac{\partial T_x}{\partial y} \\ \frac{\partial T_y}{\partial x} & \frac{\partial T_y}{\partial y} \end{bmatrix}$$

en chaque point

example



inverse  
localisation  
orientatie =  $P_b$

= knoopt van  
inverkeerde localisering  
pas sou hanteer  
imaging medicale

→ on cherchera des transf' difféomorphiques

(inversible + lisse + ...  
 $(\det(J) > 0)$

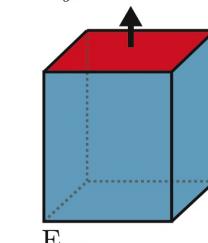
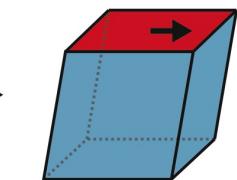
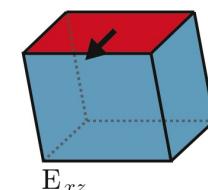
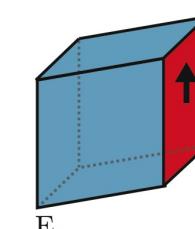
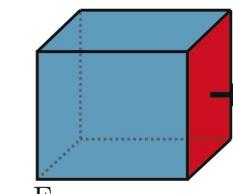
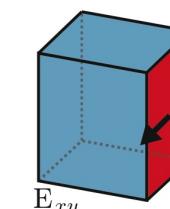
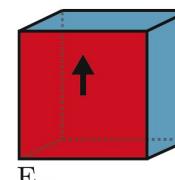
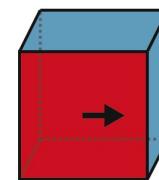
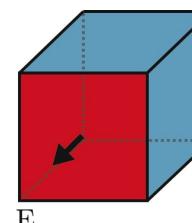
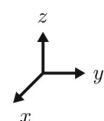
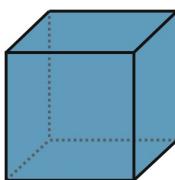
## Matrix representations: shear

$$M = R \begin{bmatrix} R_{00} & R_{01} & R_{02} & t_0 \\ R_{10} & R_{11} & R_{12} & t_1 \\ R_{20} & R_{21} & R_{22} & t_2 \\ 0 & 0 & 0 & 1 \end{bmatrix}$$

3D: 6 parameters

$$R = \begin{bmatrix} 1 & s_{yx} & s_{zx} \\ s_{xy} & 1 & s_{zy} \\ s_{xz} & s_{yz} & 1 \end{bmatrix}$$

ou 3 ?  
 (symétrie)



$$M \mathbf{x} \longleftrightarrow \mathbf{R} \mathbf{x} + \mathbf{t}$$

= 12 parameters to optimize !

Matrix representations. **skew**

$$M = \begin{bmatrix} R_{00} & R_{01} & R_{02} & t_0 \\ R_{10} & R_{11} & R_{12} & t_1 \\ R_{20} & R_{21} & R_{22} & t_2 \\ 0 & 0 & 0 & 1 \end{bmatrix}$$

$$M x \longleftrightarrow R x + t$$

= 12 parameters to optimize!

3D: 3 parameters

$$R = \begin{bmatrix} 1 & s_{yx} & s_{zx} \\ -s_{yx} & 1 & s_{zy} \\ -s_{zx} & -s_{zy} & 1 \end{bmatrix}$$

Close to shear, but antisymmetric:  $R^T = -R$

Matrix representations: **projection**  $\rightarrow$  2D  $\rightarrow$  3D

- $\underbrace{\text{2D}}_{(n-1) \times n \text{ matrix}}$   $\underbrace{\text{3D}}$

- Parallel or perspective
- ex: 2D/1D perspective proj. in homogeneous coords:

SDD = Source to Detector Distance

SID = Source to Isocenter Distance

$$\mathbf{M} = \begin{bmatrix} SDD & 0 & 0 \\ 0 & 1 & SID \end{bmatrix}$$

## Summary: transformations

**Linear matrices** can be decomposed in:

<http://mathworld.wolfram.com/MatrixDecomposition.html>  
<http://www.wikipedia.com/>

Translation parameters

straightforward

Rotation parameters

need caution, several solutions

Scaling parameters

straightforward

Skew parameters

less used

+ **Projection** for 2D/3D registration

## Summary: transformations

Linear matrices can be decomposed in:

<http://mathworks.molham.org/>  
<http://www.wikipedia.com/>

Translation parameters

Rotation parameters

Scaling parameters

Skew parameters

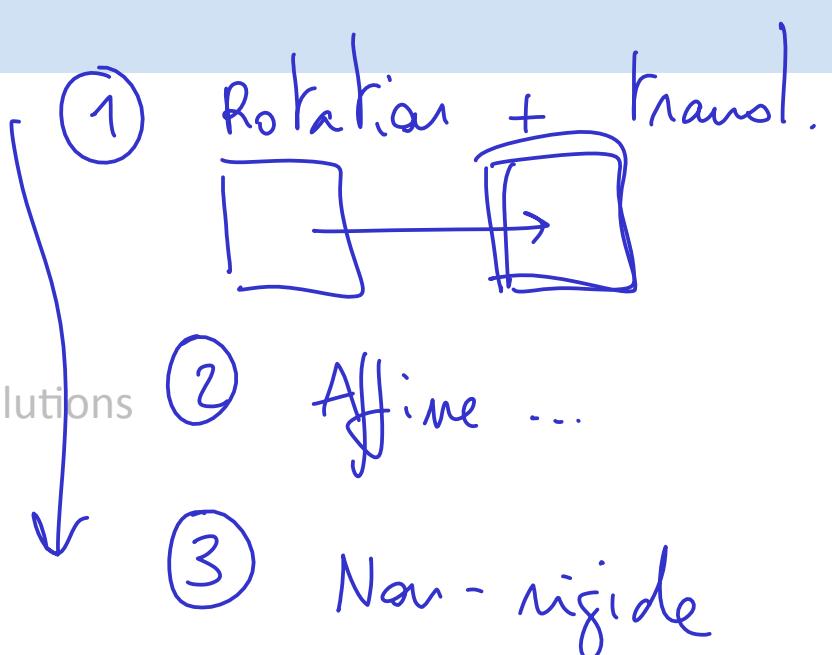
+ Projection for 2D/3D registration

straightforward

need caution, several solutions

straightforward

less used



Only a subset of the parameters can (should?) be optimized:

Rigid 6 parameters (translations + rotations)

Rigid + global scaling 7 parameters

Rigid + independent scaling 9 parameters

Affine 12 parameters

## Summary: transformations

### Tips and tricks:

#### Carefully choose the search space

- As little parameters as possible...
- ... but make sure the solution is in it !
- Bound the search space, e.g. translations < 2cm and rotations < 10°
- Scale the heterogeneous parameters, e.g. translations in mm and rotations in radians and degrees

bien définir espace des possibles.

Only a subset of the parameters can (should?) be optimized:

Rigid	6 parameters (translations + rotations)
Rigid + global scaling	7 parameters
Rigid + independent scaling	9 parameters
Affine	12 parameters

## Similarity measure

$$\hat{T} = \arg \max_{T \in \mathcal{T}} S(I, J, T)$$

## Notation

- $I$  = reference / fixed image
- $J$  = moving / floating image
- $T$  = transformation
- $\mathcal{T}$  = search space
- $S$  = similarity measure
- $\arg \max$  = optimization
- $\hat{T}$  = solution

## Similarity measure

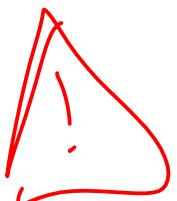
Measures the **alignment quality** of I and J:

$$\hat{T} = \arg \max_{T \in \mathcal{T}} S(I, J, T)$$

Q: Any idea?

$$S(I, J)$$

$$S(I, T(J))$$



$$S(I, J) = \sqrt{\frac{1}{N} \sum_{i=1}^N (I_i - J_i)^2}$$

RMSE

$$MAD = \frac{1}{N} \sum_{i=1}^N |I_i - J_i|$$

$$\text{ia } S(I, J) = 0$$

si similaires

$$\hat{T} = \operatorname{argmin} S(I, T(J))$$

corrélation :  $s(I, J) \approx 1$  si similaires  
0 sinon

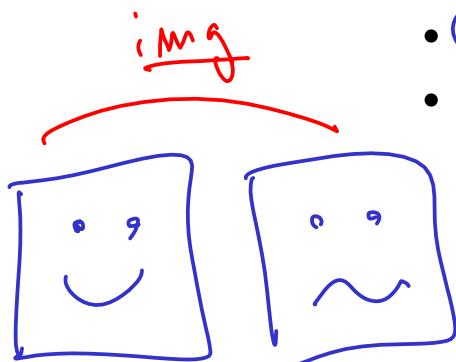
+ plein d'arts → à choisir vs. notre pb

## Similarity measure

Measures the **alignment quality** of I and J:  
on  $\arg\min$

$$\hat{T} = \arg \max_{T \in \mathcal{T}} S(I, J, T)$$

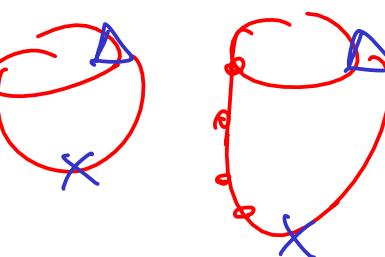
Q: Any idea?



- Two families:
- Feature-based registration
  - Image-based registration

en tout point

extraction des points



distance

$$D(I, T(J))$$

## Similarity measure: **feature-based**

Features are extracted **prior to the similarity measure**:

- Points/landmarks
- Lines
- Vectors
- Surfaces
- Volumes
- ...

$$\hat{T} = \arg \max_{T \in \mathcal{T}} S(\mathcal{F}_I, T \circ \mathcal{F}_J)$$

### Notation

- $\mathcal{F}_I$  = feature set of reference / fixed image
- $\mathcal{F}_J$  = feature set of moving / floating image
- $T$  = transformation
- $\mathcal{T}$  = search space
- $S$  = similarity measure
- $\arg \max$  = optimization
- $\hat{T}$  = solution

## Similarity measure: feature-based

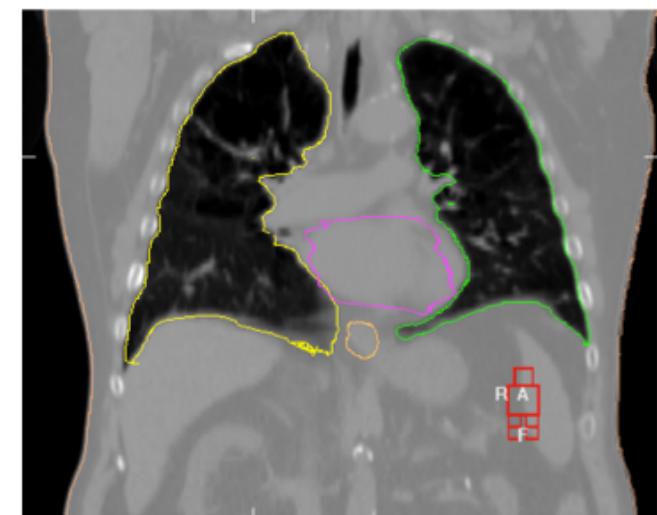
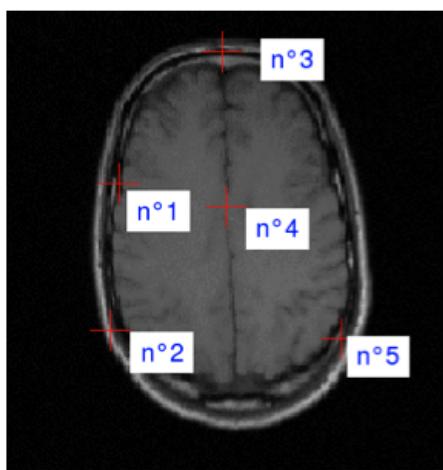
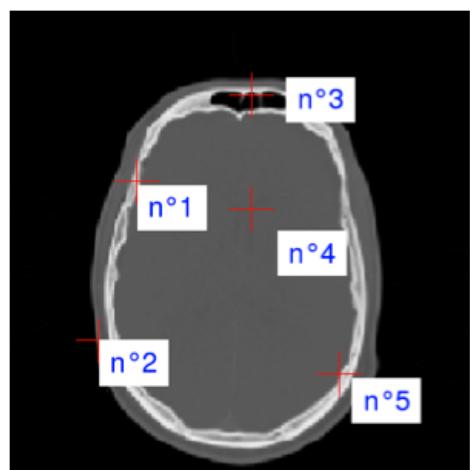
Features are extracted prior to the similarity measure:

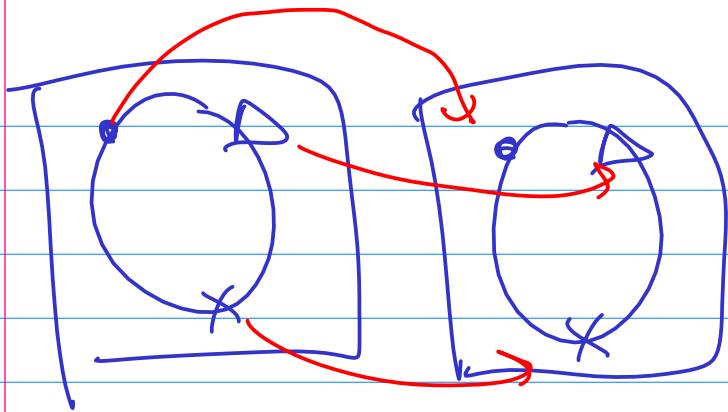
- Points/landmarks
- Lines
- Vectors
- Surfaces
- Volumes
- ...

- Manually or automatically extracted
- Can be unpaired = not same sampling, amount, ...

(machine learning?)

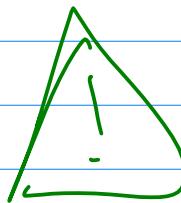
landmarks  
non-identified





Q: que se passe-t-il entre  
les points clés ?

interpoler entre



Régularisation \*\*\*



TIP#3

Similarity measure: **feature-based**(Dis-)similarity measure for features:

- Sum of distances:

- L1 norm

$$|\mathbf{x}|_1 = \sum_{r=1}^n |x_r| \rightarrow \text{MAD}$$

- L2 norm

$$|\mathbf{x}|_2 = \sqrt{\sum_{r=1}^n x_r^2} \leftarrow \begin{matrix} \text{SSD} \\ \text{RMSE} \end{matrix}$$

- Quadratic sum (faster):

$$\sum_{r=1}^n x_r^2 \quad \dots$$

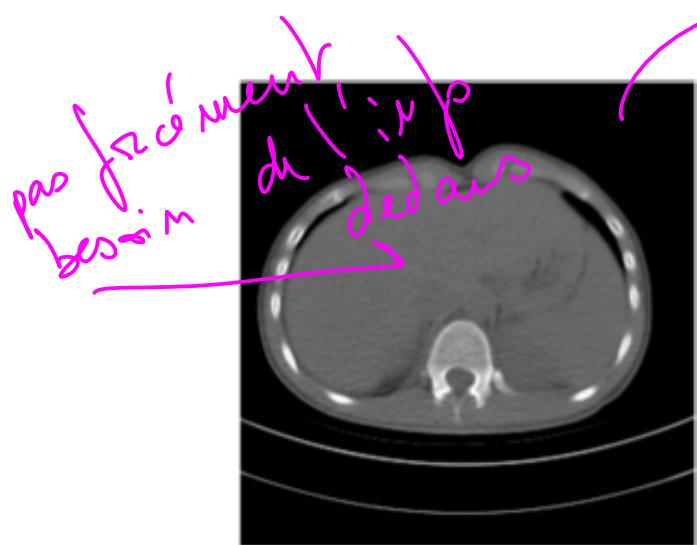
- ...

Similarity measure: **feature-based**

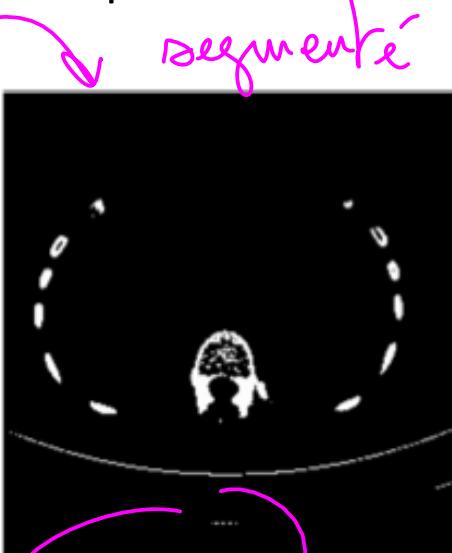
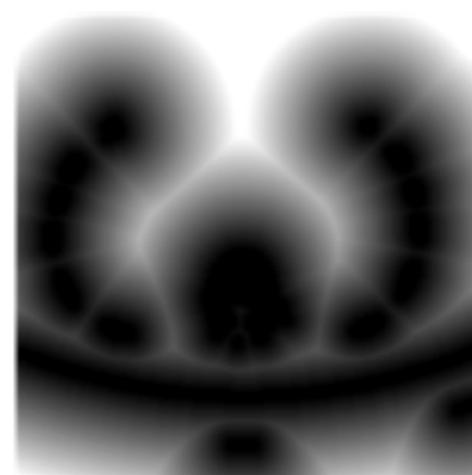
(Dis-)similarity measure for features:

**+ distance map**

= map of the distance between each point and the closest feature  
= fast + useful for unpaired features

*+ flexible*

Image

*segmenté***Features = bones**

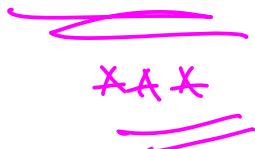
Distance map

- **Computed only once** for the reference image
- **Fast computations:** Chamfer distance (not Euclidean)  
separable algorithms [Coeurjolly et al. PAMI 2007]

Similarity measure: **feature-based**



Symmetry?



$$S(\mathcal{F}_I, T \circ \mathcal{F}_J) \text{ or } S(\mathcal{F}_J, T \circ \mathcal{F}_I)?$$

Different !

or combiner les deux ?

→ algo explicitement symétriques

$$S(I, J, T) = \sum_{F \in \mathcal{F}_J} d_I \circ T(F)$$

$$S(I, J, T) = \sum_{F \in \mathcal{F}_I} d_J \circ T(F)$$



ex :

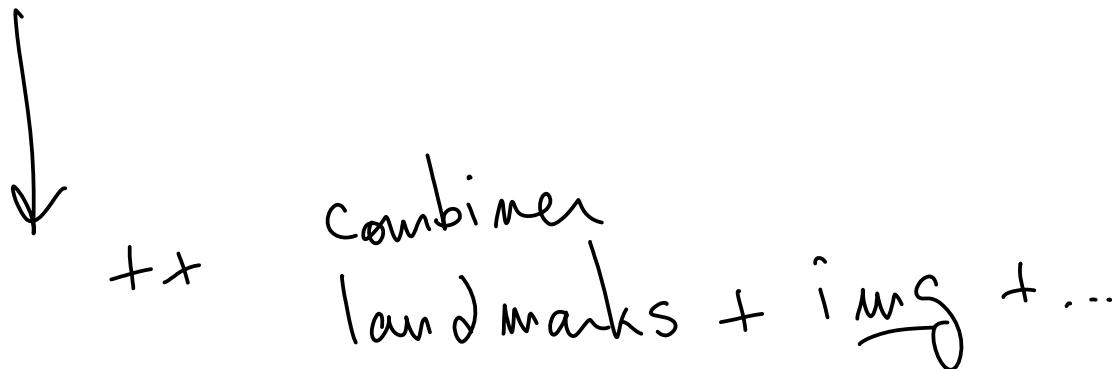
- Use the distance map of the image with the easiest-to-extract features

Similarity measure: **feature-based**

### Summary:

- Preliminary steps to identify / extract features
- Pairing algorithm may be required
- + Generally fast (depends on the feature type and size)
- + Registration of some features only...

➤ Still interesting, especially for non-rigid registration...



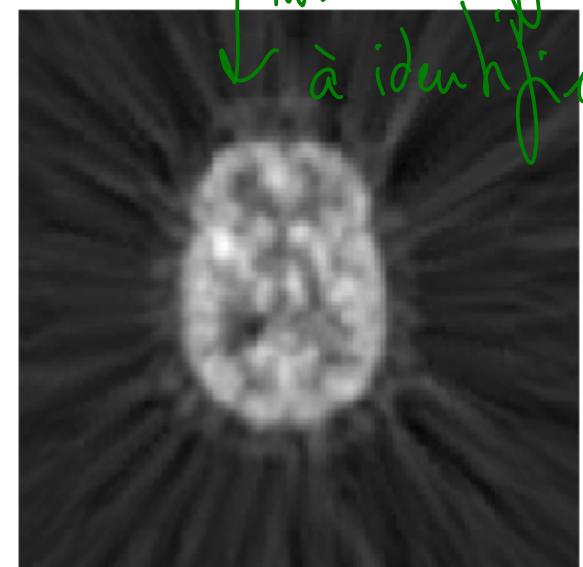
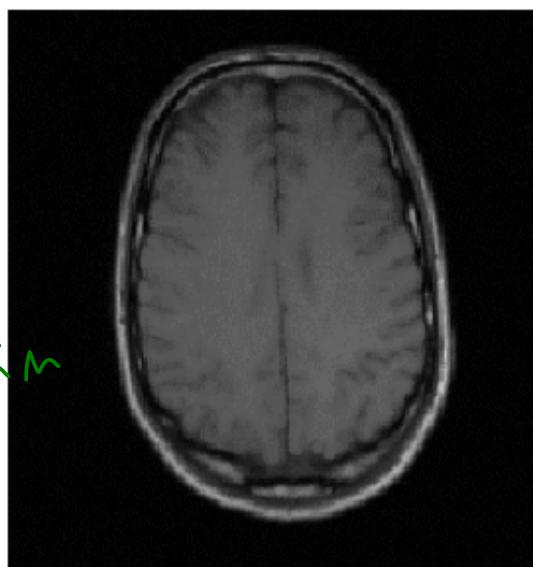
Similarity measure: **feature-based**

### Summary:

- Preliminary steps to identify / extract features
- Pairing algorithm may be required
- + Generally fast (depends on the feature type and size)
- + Registration of some features only...
  - Still interesting, especially for non-rigid registration...

### Another limitation:

- **Multi-modal images**
  - ex: NMI d. + bin
- **Image-based registration**



landmarks  
points /  
mais difficiles  
à identifier

Similarity measure: **intensity-based**

(assuming a functional dependence between the pixel intensities)

$$\hat{T} = \arg \max_{T \in \mathcal{T}} S(I, J \circ T)$$

### Notation

- $I$  = reference / fixed image
- $J$  = moving / floating image
- $T$  = transformation
- $\mathcal{T}$  = search space
- $S$  = similarity measure
- $\arg \max$  = optimization
- $\hat{T}$  = solution

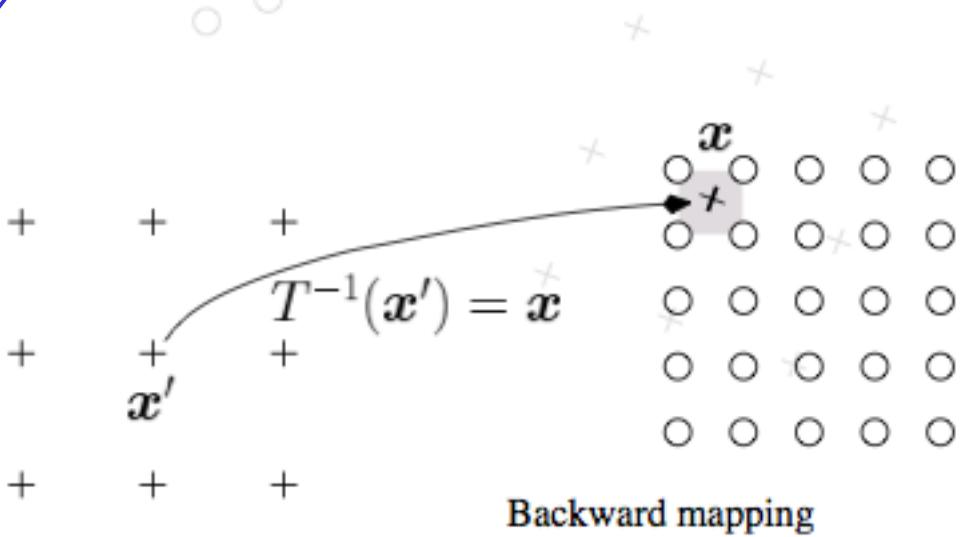
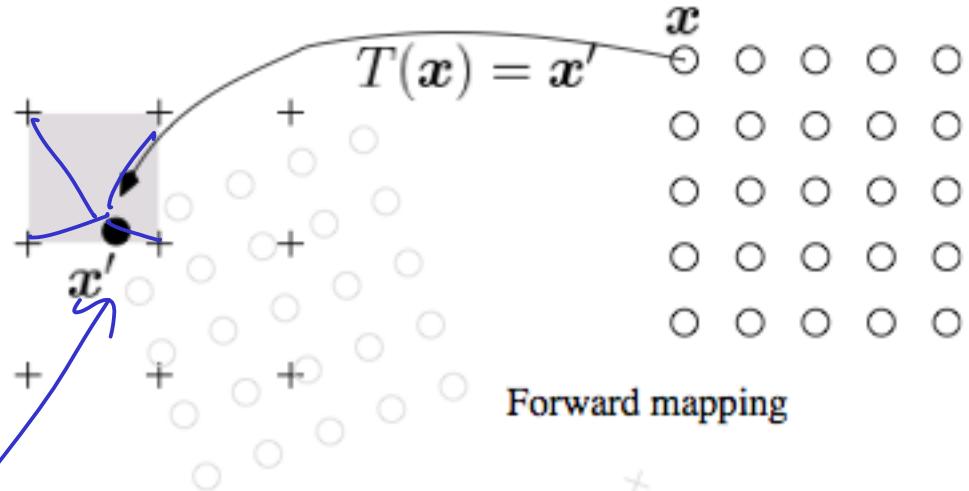
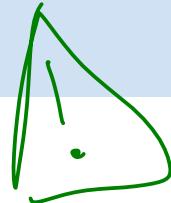
Similarity measure: **intensity-based**

Forward and backward mappings:

Required to compute  ~~$J \circ T$~~   $+ (J)$

interpolation! \*\*\*

ne tombe pas  
fréquemment sur  
un pixel



Similarity measure: **intensity-based**

Forward and backward mappings:

Required to compute  $J \circ T$

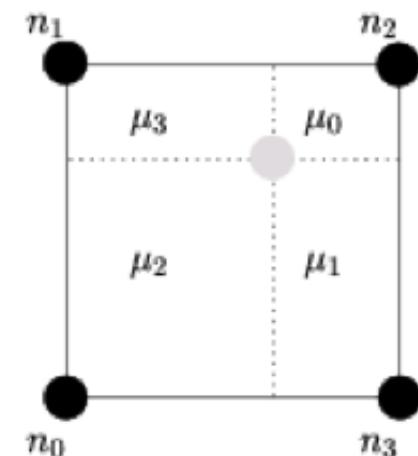
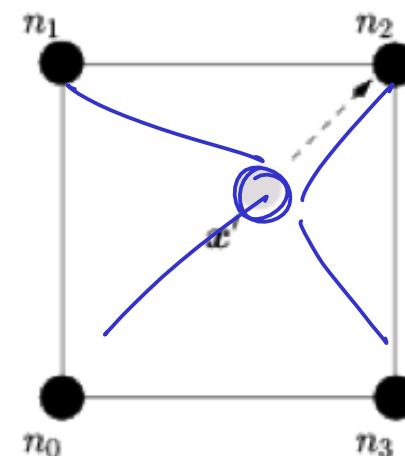
**Forward** requires an additional weight map + potential holes

**Backward** usually preferred: not a problem

- If  $T$  is invertible (ok with affine, otherwise should be enforced)
- Optimize  $T^{-1}$  and invert it at the end

Beware: **interpolation required !**

- Nearest neighbor
  - Linear
  - More accurate: quadratic, spline, ...
- = compromise between speed and accuracy



Similarity measure: **intensity-based**

(Dis-)similarity measure for intensities:

**Sum-of-squared differences (SSD):**

- Fast
- Needs to be **normalized** to the domain size (in voxels)
- Functional dependence between intensities:
  - **Ok if same modality + noise**
  - **Impossible if multimodal !**
- Similar to *Sum-of-absolute differences (SAD)*

$$SSD(I, J) = \sum_{x \in \Omega} (I(x) - J(x))^2$$

*sensible au b/w*

Similarity measure: **intensity-based**(Dis-)similarity measure for intensities:

Sum-of-squared differences (SSD)

**(Pearson) correlation coefficient:**

- Range = [ -1 , 1 ]
  - 1 = perfect increasing linear relationship
  - 1 = ...
  - 0 = independent ↙

&gt; Similarity measure = | CC( I , J ) |

- Fast (with computational tricks) ↘

$$\begin{aligned} CC(I, J) &= \frac{\text{cov}(I, J)}{\sigma_I \sigma_J} \\ &= N \frac{\sum_i (I_i(\mathbf{x}) - m_I)(J_i(\mathbf{x}) - m_J)}{\sqrt{\sum_i (I_i(\mathbf{x}) - m_I)^2} \sqrt{\sum_i (J_i(\mathbf{x}) - m_J)^2}} \end{aligned}$$

## Similarity measure: **intensity-based**

### (Dis-)similarity measure for intensities:

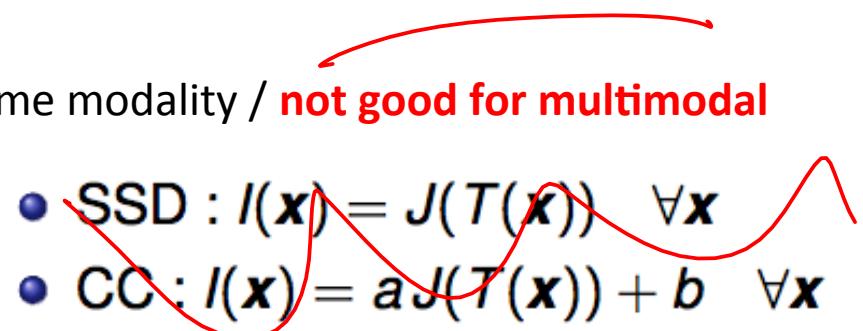
Sum-of-squared differences (SSD)

(Pearson) correlation coefficient:

- Range = [ -1 , 1 ]
    - 1 = perfect increasing linear relationship
    - 1 = ...
    - 0 = independent
- > Similarity measure = | CC( I , J ) |

- Fast (with computational tricks)
- More robust than SSD** for 2 images from the same modality / **not good for multimodal**
- Functional dependence between intensities:
  - SSD :  $I(\mathbf{x}) = J(T(\mathbf{x})) \quad \forall \mathbf{x}$
  - CC :  $I(\mathbf{x}) = aJ(T(\mathbf{x})) + b \quad \forall \mathbf{x}$

$$\begin{aligned} CC(I, J) &= \frac{\text{cov}(I, J)}{\sigma_I \sigma_J} \\ &= N \frac{\sum_i (I_i(\mathbf{x}) - m_I)(J_i(\mathbf{x}) - m_J)}{\sqrt{\sum_i (I_i(\mathbf{x}) - m_I)^2} \sqrt{\sum_i (J_i(\mathbf{x}) - m_J)^2}} \end{aligned}$$



## Similarity measure: **intensity-based**

### (Dis-)similarity measure for intensities:

Sum-of-squared differences (SSD)

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- Range = [ -1 , 1 ]
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- Other functional relationship? = **information theory**:

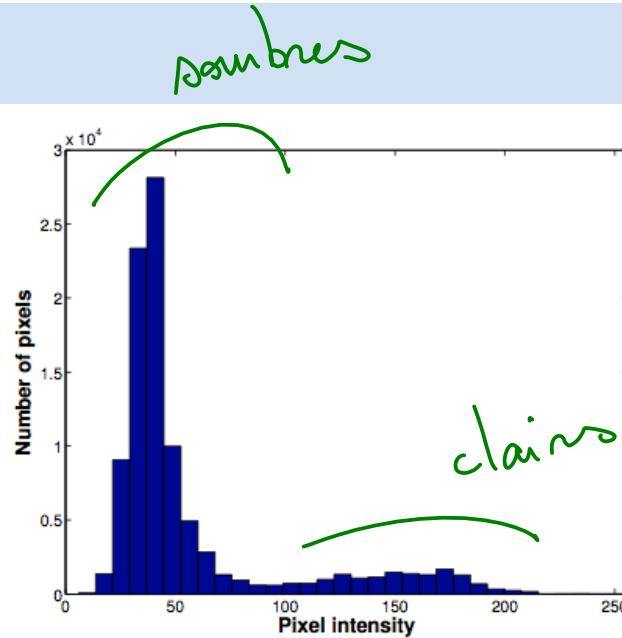
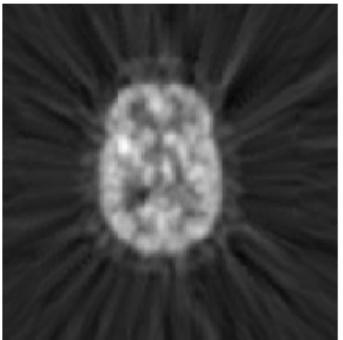
- Mutual information, coefficient ratio...

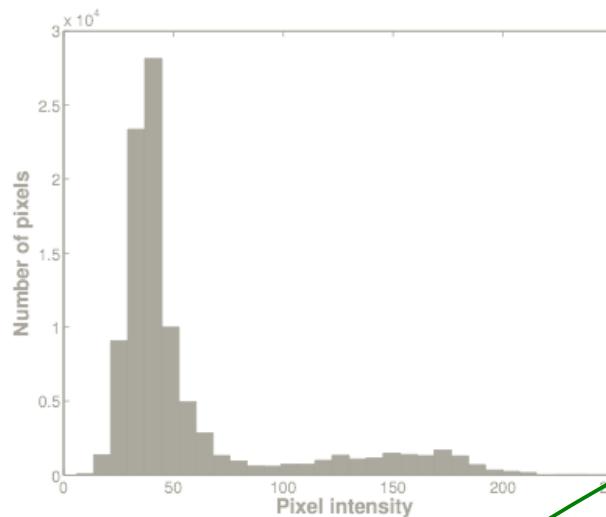
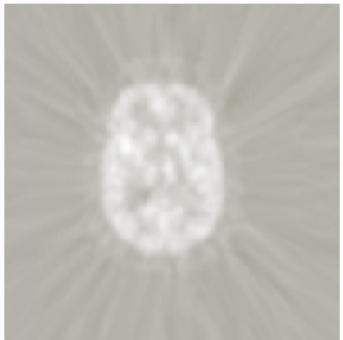
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- CC :  $I(\mathbf{x}) = aJ(T(\mathbf{x})) + b \quad \forall \mathbf{x}$

both require **joint histograms**

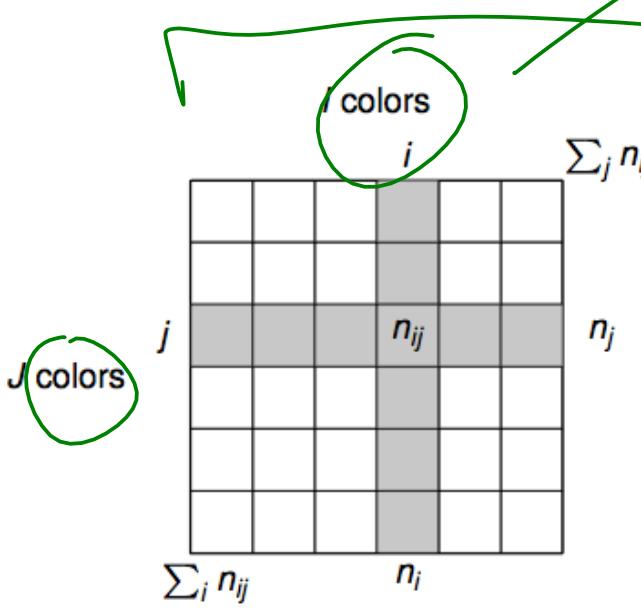
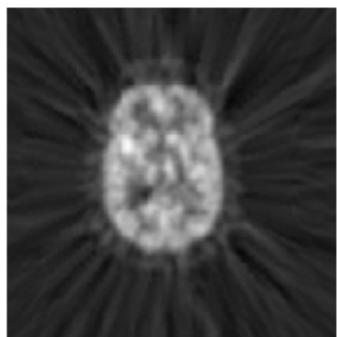
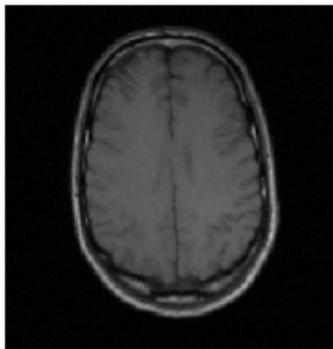
Similarity measure: **intensity-based**

Histogram:



Similarity measure: **intensity-based**Histogram:

pas pixels  
mais mb  
bins  
dans l'histogramme

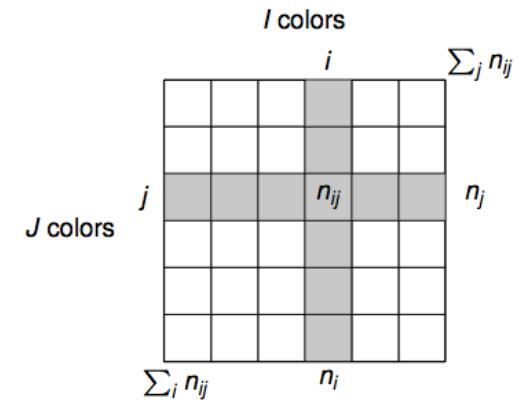
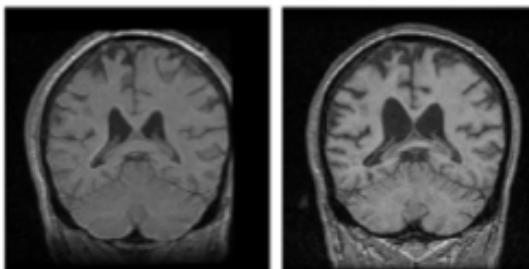
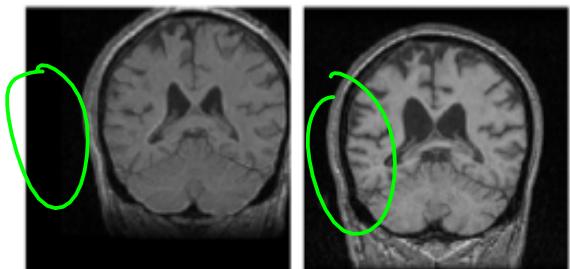
Joint histogram:

Not pixel indices !

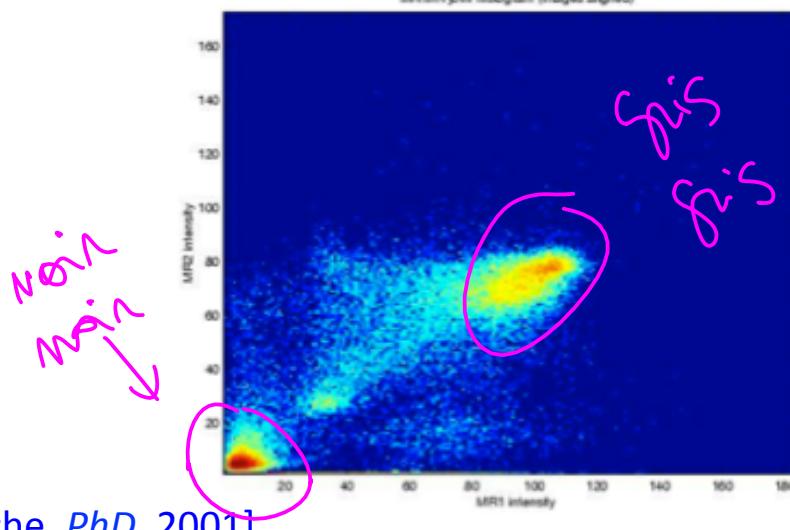
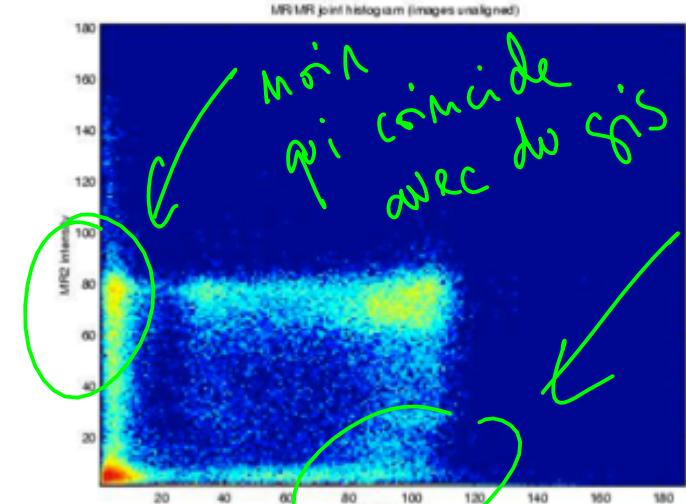
- $i, j$  : (binned) colors
- $n_{ij}$ : number of pixels with color  $i$  in  $I$  and  $j$  in  $J$
- $n_i, n_j$ : marginal values, i.e., histograms of  $I$  and  $J$

Similarity measure: **intensity-based**Joint histogram:

- **2D distribution** of the intensity pairs at each voxel
- Recomputed at each step of the optimisation



**Mono-modal**

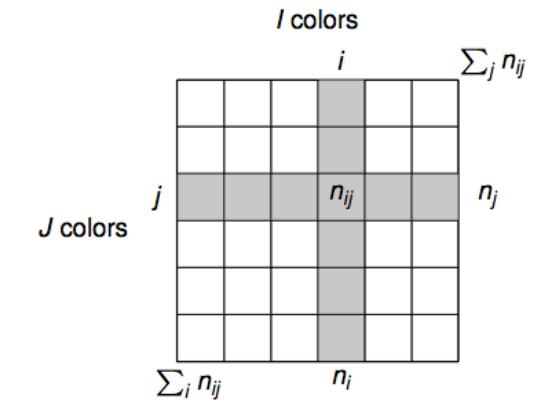
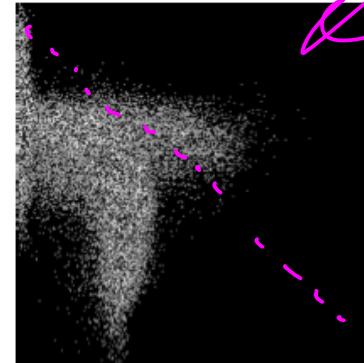
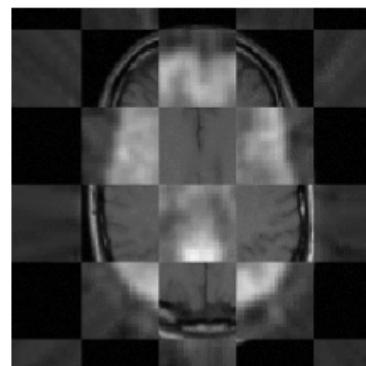
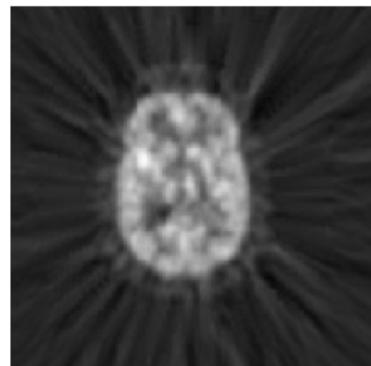
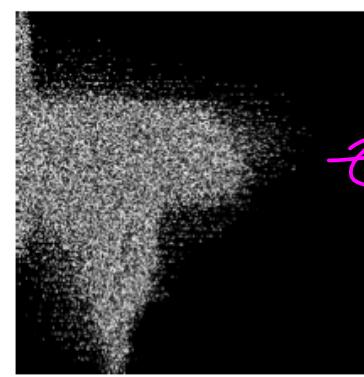
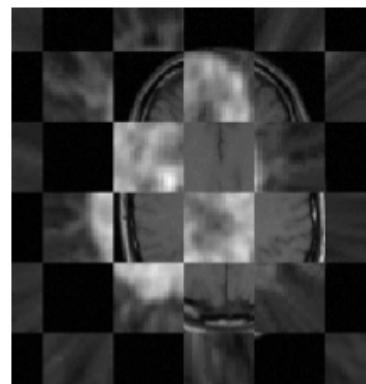
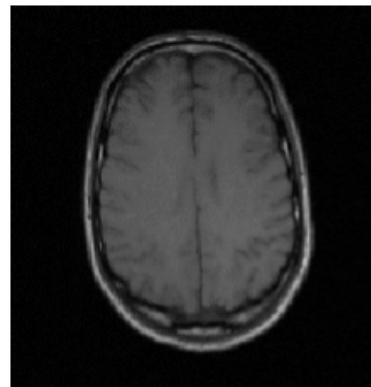


[Roche, PhD, 2001]

Similarity measure: **intensity-based**Joint histogram:

- **2D distribution** of the intensity pairs at each voxel
- Recomputed at each step of the optimisation

plus subtîl!



Multi-modal

diagonale inversée?  
(sans cette p'titi)

Similarity measure: **intensity-based**Joint histogram: what for?

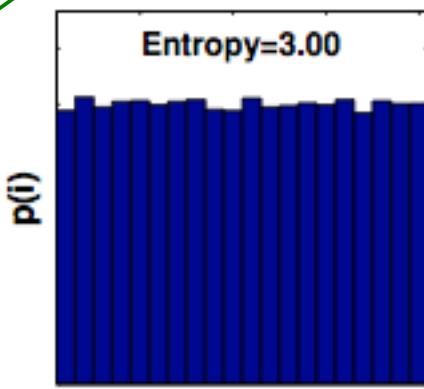
- Mutual information – entropy = measure of information as registration metric

Entropy (Shannon-Wiener):

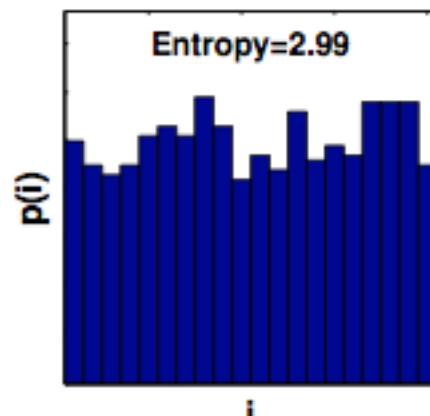
 $\approx$  désordre

histogramme  
à 1 octave image

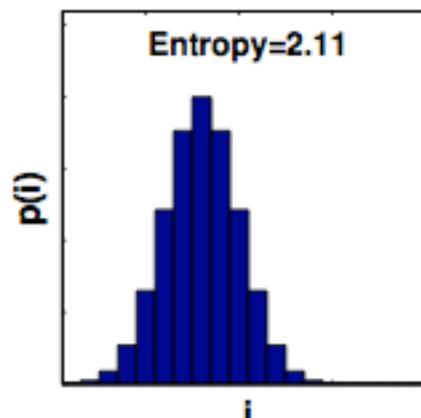
$$H = \sum_i p_i \log \frac{1}{p_i} = - \sum_i p_i \log p_i$$



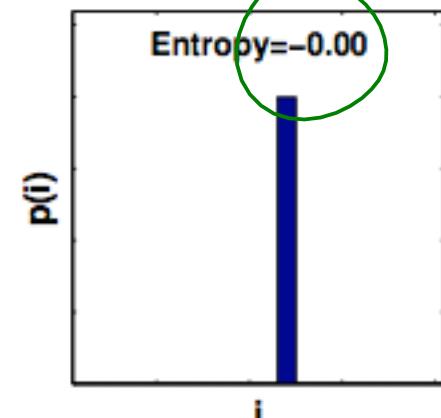
buit pur



img bruitée



img

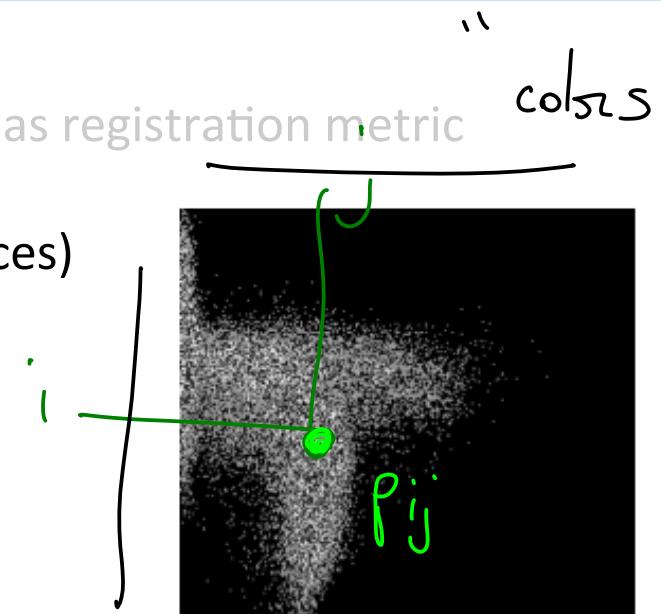


img constante

Similarity measure: **intensity-based**Joint histogram: what for?

- Mutual information – entropy = measure of information as registration metric
- Mutual information – joint entropy (2 images = 2 subindices)

$$H = - \sum_{i,j} p_{ij} \log p_{ij}$$



&gt; The more similar the distributions...

cf. &  
lim

= the lower the joint entropy  
(compared to the sum of individual entropies)

$$H(I, J) \leq H(I) + H(J)$$

Similarity measure: **intensity-based**Joint histogram: what for?

- Mutual information – entropy = measure of information as registration metric
- Mutual information – joint entropy (2 images = 2 subindices)

$$H(I, J) \leq H(I) + H(J)$$

- **Mutual information:**

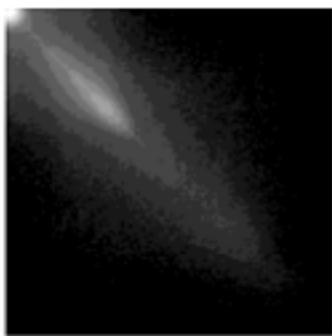
$$MI(I, J) = \sum_{i,j} p_{ij} \log \frac{p_{ij}}{p_i p_j}$$

$$MI(I, J) = H(I) - H(J|I,J) = H(J) - H(I|J) = H(I) + H(J) - H(I, J) \geq 0$$

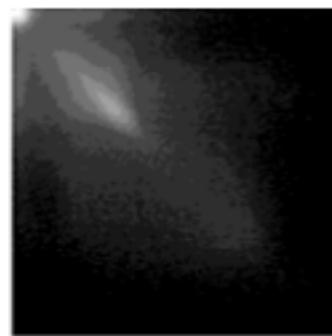


3.82

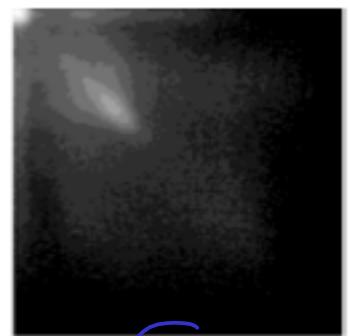
[Pluim et al. IEEE TMI, 2003]



6.79



6.98



7.15

## Similarity measure: **intensity-based**

### Joint histogram: what for?

- Mutual information – entropy = measure of information as registration metric
- Mutual information – joint entropy (2 images = 2 subindices)

- Mutual information:

$$MI(I, J) = \sum_{i,j} p_{ij} \log \frac{p_{ij}}{p_i p_j}$$

- Positive

- Symmetric:  $MI(I, J) = \underbrace{MI(J, I)}_{**}$

- $H(J | I) = H(I, J) - H(I) = \text{conditional entropy}$   
~~~~~

## Similarity measure: intensity-based

### Joint histogram: what for?

- Mutual information – entropy = measure of information as registration metric
- Mutual information – joint entropy (2 images = 2 subindices)
- Mutual information
- Normalized mutual information
  - Goal = overcome the **sensitivity of MI** to variations of the overlap size between  $I$  and  $J$  or  $T$
  - Several versions
  - Example = [Studholme et al. Patt Recogn, 1999]
  - It's a reference !!!
    - First articles: 1995 (Maes et al. [Belgium], Viola et al. [US])
    - First journal articles: 1997
    - Among the most cited articles in the field...*

$$NMI(I, J) = \frac{H(I) + H(J)}{H(I, J)}$$

## Similarity measure: **intensity-based**

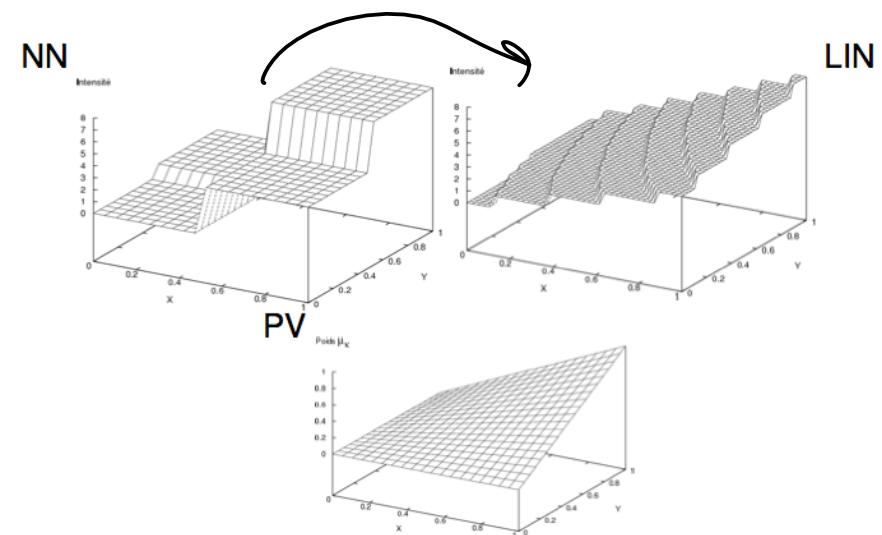
### Joint histogram: what for?

- Mutual information – entropy = measure of information as registration metric
- Mutual information – joint entropy (2 images = 2 subindices)
- Mutual information
- Normalized mutual information
  - **Currently = #1 similarity measure !    Multi-modal + robust +++**
  - Implementation matters (as usual...)
    - Survey of methods = [Pluim et al. *IEEE TMI* 2003]

## Similarity measure: **intensity-based**

### Joint histogram: what for?

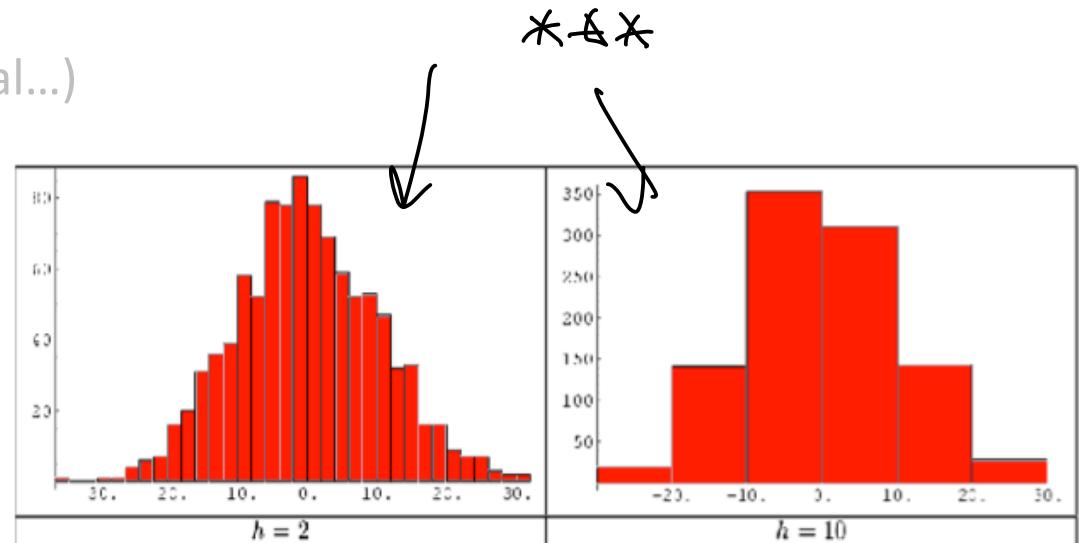
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    - Specifically:
      - Interpolation



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      - Binning



## Similarity measure: **intensity-based**

### Joint histogram: what for?

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- Mutual information
- Normalized mutual information
  - Currently = #1 similarity measure ! Multi-modal + robust +++
  - Implementation matters (as usual...)
    - Specifically:
      - Interpolation
      - Binning
      - Probability computation
      - Normalization (NMI) /

## Similarity measure: **intensity-based**

### Summary:

- The measures can be classified vs. hidden variables [Malandain, *HDR*, 2006]

• SSD / SAD

0 variable

• CC

2 variables: slope + intercept

• MI

size of the joint histogram matrix

- **MI is more general**, hence its success...
- ... but more susceptible to **fail in simple situations**

/ where SSD is enough  
(e.g. time series)

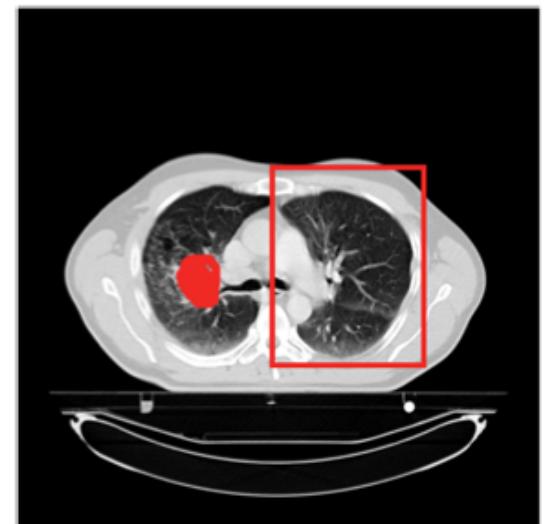
## Similarity measure: **intensity-based**

### Summary:

- The measures can be classified vs. hidden variables [Malandain, *HDR*, 2006]
  - SSD / SAD      0 variable
  - CC                2 variables: slope + intercept
  - MI                size of the joint histogram matrix
- 
- MI is more general, hence its success...
  - ... but more susceptible to fail in simple situations

Remark: region of interest (ROI) ↗

> applied to the reference and/or the target  
(depending on the similarity measure)



## Optimization

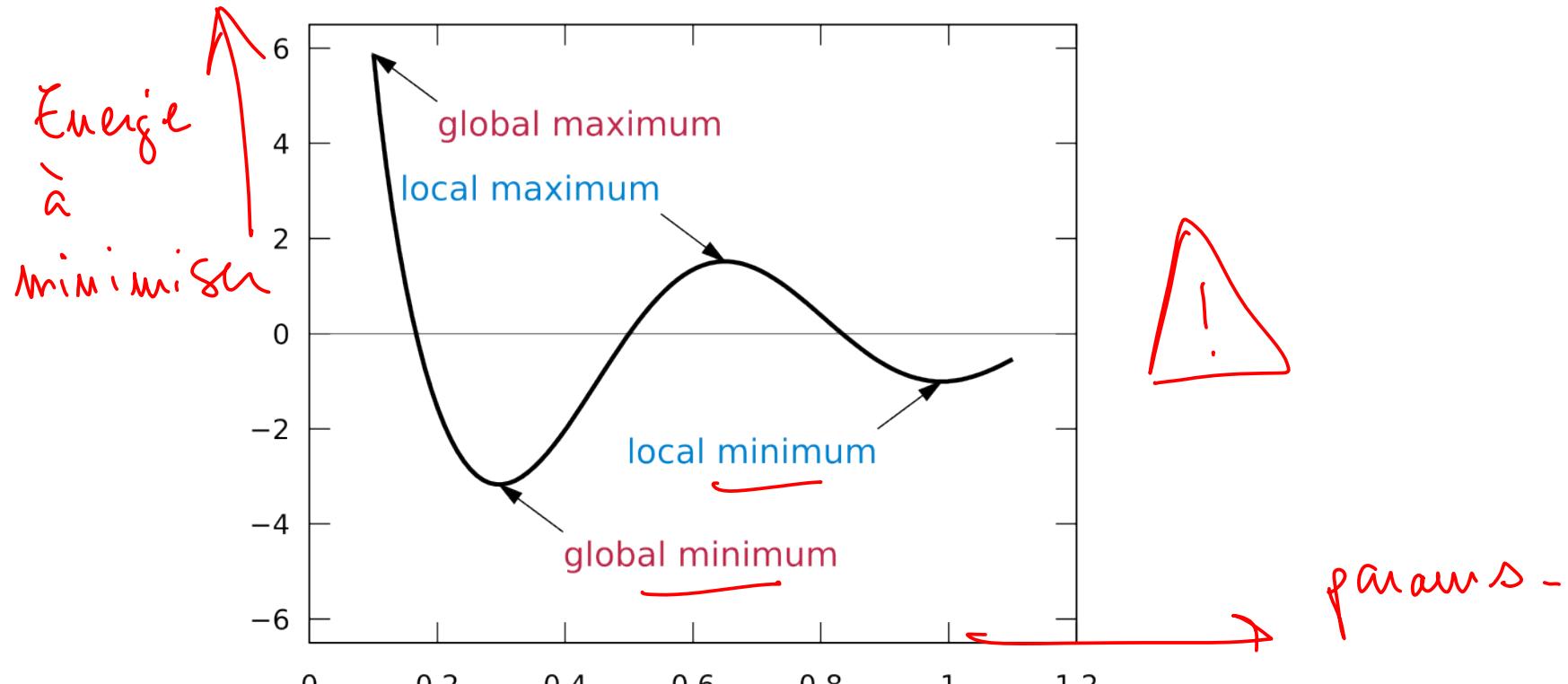
$$\hat{T} = \arg \max_{T \in \mathcal{T}} S(I, J, T)$$

## Notation

- $I$  = reference / fixed image
- $J$  = moving / floating image
- $T$  = transformation
- $\mathcal{T}$  = search space
- $S$  = similarity measure
- $\arg \max$  = optimization
- $\hat{T}$  = solution

## Optimization

Challenge = find the **global** minimum... as fast as possible



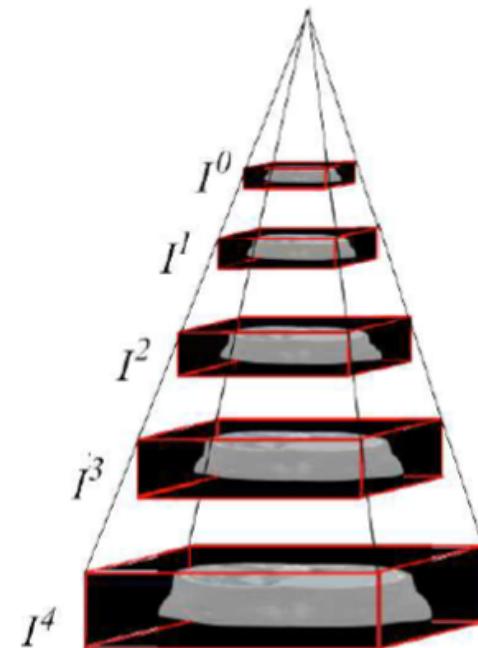
ex : deep : params = points de réseaux  
ex : recette : params :  $\begin{bmatrix} a_1 & a_2 & a_3 \end{bmatrix} = R$   
 $a_{\dots}$

## Optimization

Challenge = find the **global** minimum... as fast as possible

- **Maximize or minimize** (depending on the energy formulation)  

- With or without gradient (if  $f$  is differentiable)  

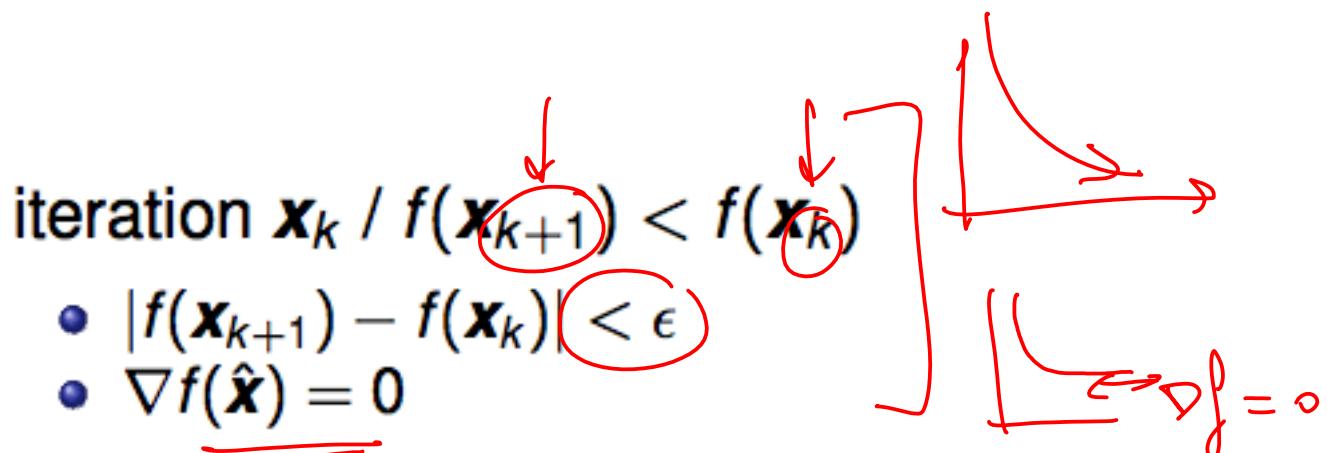
- **Multi-resolution**
  - Convergence
  - Robustness to noise + local minima
  - Large deformations

$$T_{\text{final}} = T_4 \circ T_3 \circ \dots$$

## Optimization

Challenge = find the **global** minimum... as fast as possible

- Maximize or minimize (depending on the energy formulation)
- With or without gradient (if  $f$  is differentiable)
- Multi-resolution
- Iterative process
- Stopping criterion =
  - $|f(\mathbf{x}_{k+1}) - f(\mathbf{x}_k)| < \epsilon$
  - $\nabla f(\hat{\mathbf{x}}) = 0$
- Update = **search  $\delta \mathbf{x}_k$**  /  $\mathbf{x}_{k+1} = \mathbf{x}_k + \delta \mathbf{x}_k$



## Optimization

Challenge = find the **global** minimum... as fast as possible

- Maximize or minimize (depending on the energy formulation)
- With or without gradient (if  $f$  is differentiable)
- Multi-resolution
- Iterative process      **iteration**  $\mathbf{x}_k / f(\mathbf{x}_{k+1}) < f(\mathbf{x}_k)$
- Stopping criterion =
  - $|f(\mathbf{x}_{k+1}) - f(\mathbf{x}_k)| < \epsilon$
  - $\nabla f(\hat{\mathbf{x}}) = 0$
- Update =    **search**  $\delta\mathbf{x}_k / \mathbf{x}_{k+1} = \mathbf{x}_k + \delta\mathbf{x}_k$
- Importance of initialization !!!     **$\mathbf{x}_0$** 
  - Pre-align image origins / centers / mass centers
  - Rigid alignment before non-rigid



## Optimization

Process / methods = cf. supplementary slides [RIT-SARRUT\_optimization.pdf]

# Optimization

## Summary:

- **Without gradient**

- Simple
- Powell (conjugate directions)

- **With gradient:**

- Gradient descent
- Conjugate gradient
- Quasi-Newton
- Levenberg-Marquart

- **Also:**

- Genetic algorithms
- Simulated annealing
- ...

## Validation

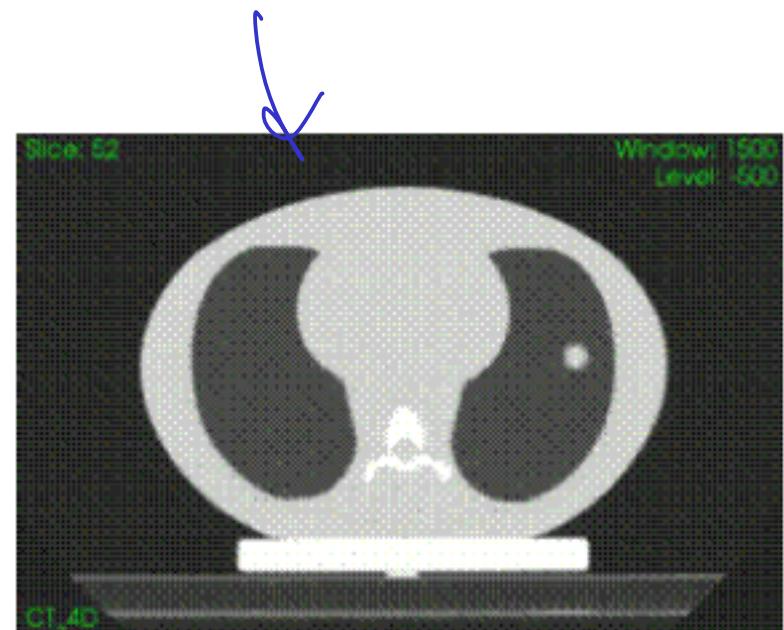
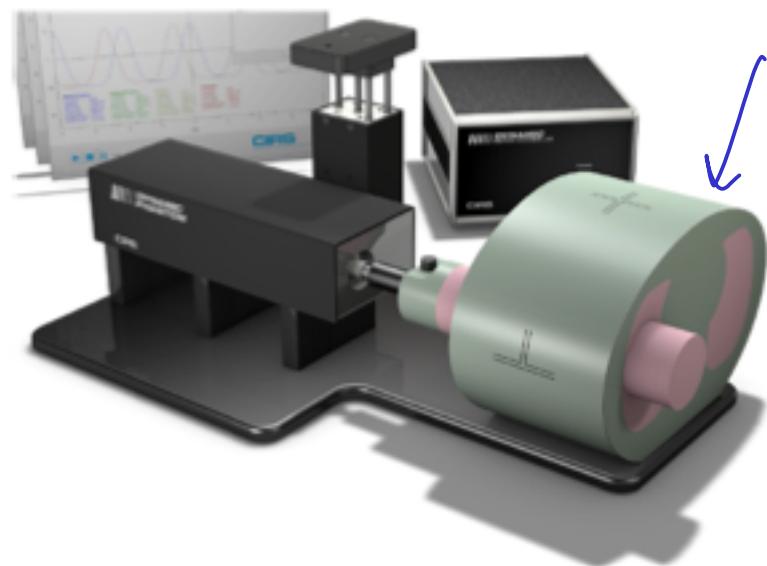
(further developed in D. Sarrut's part)

|          | Clinical data | Cadavers | Physical phantoms | Realistic simulations | Numerical simulations |
|----------|---------------|----------|-------------------|-----------------------|-----------------------|
| Easy     |               | -        |                   |                       | +                     |
| Control  |               |          |                   |                       |                       |
| Clinical |               |          |                   |                       |                       |
| Realism  |               | +        | ←                 |                       | -                     |

## Validation

(further developed in D. Sarrut's part)

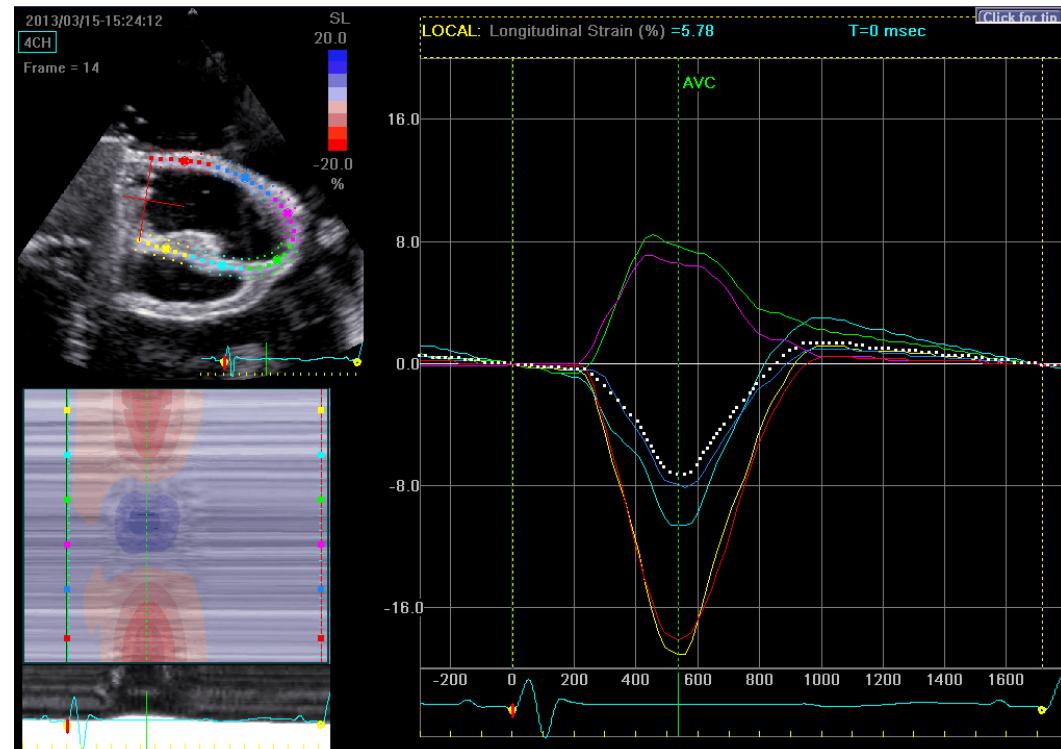
|                  | Clinical data | Cadavers | Physical phantoms | Realistic simulations | Numerical simulations |
|------------------|---------------|----------|-------------------|-----------------------|-----------------------|
| Easy Control     | —             | —        | —                 | → +                   |                       |
| Clinical Realism | + ←           |          |                   | —                     |                       |



## Validation

(further developed in D. Sarrut's part)

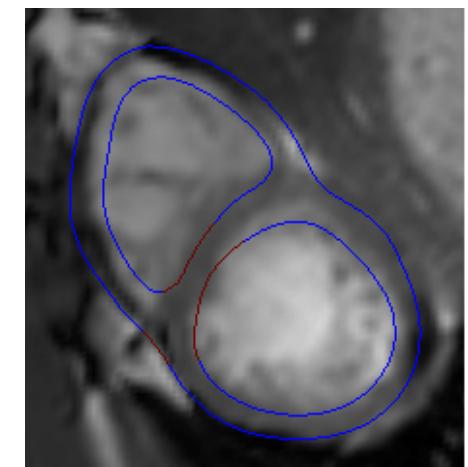
|                  | Clinical data | Cadavers | Physical phantoms | Realistic simulations | Numerical simulations |
|------------------|---------------|----------|-------------------|-----------------------|-----------------------|
| Easy Control     | —             | —        |                   | +                     |                       |
| Clinical Realism | +             | ←        |                   | —                     |                       |



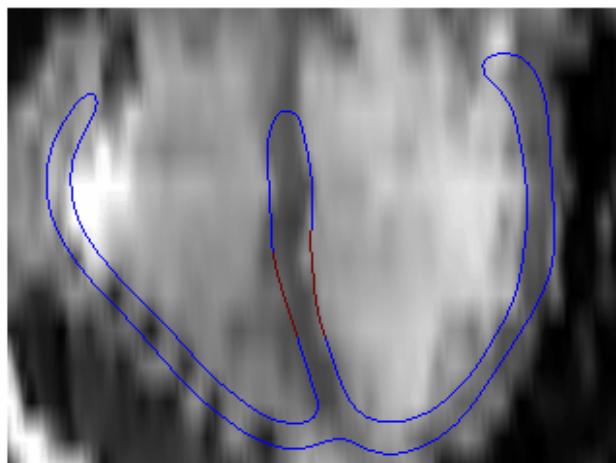
## Validation

(further developed in D. Sarrut's part)

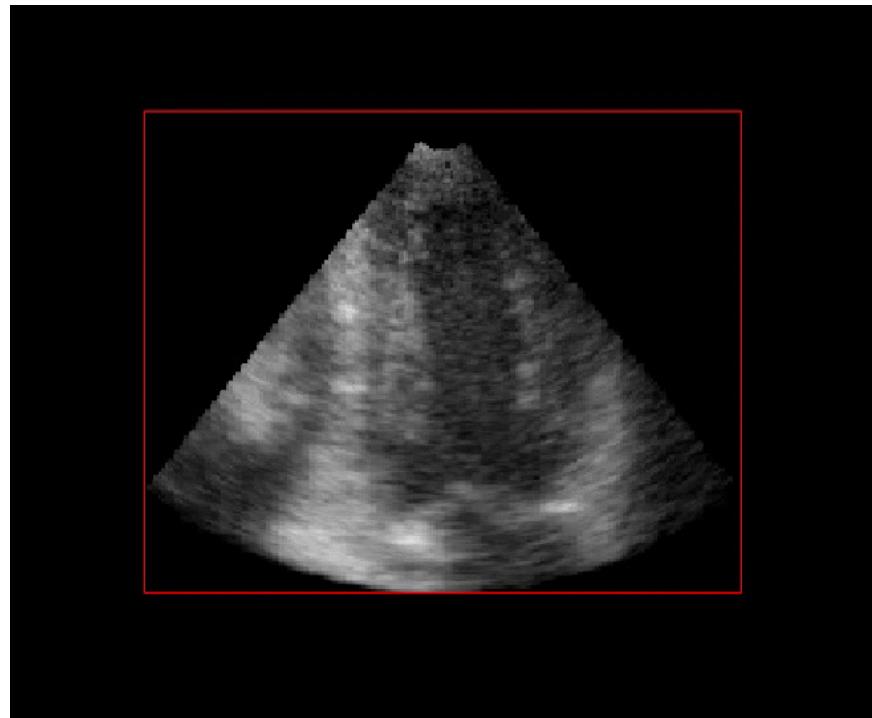
|                  | Clinical data | Cadavers | Physical phantoms | Realistic simulations | Numerical simulations |
|------------------|---------------|----------|-------------------|-----------------------|-----------------------|
| Easy Control     |               | —        | —                 | → +                   |                       |
| Clinical Realism |               | + ←      |                   |                       | —                     |



[Duchateau et al. IEEE TMI, 2018]



[Zhou et al. IEEE TMI, 2018]



## Validation

(further developed in D. Sarrut's part)

Kaggle ...

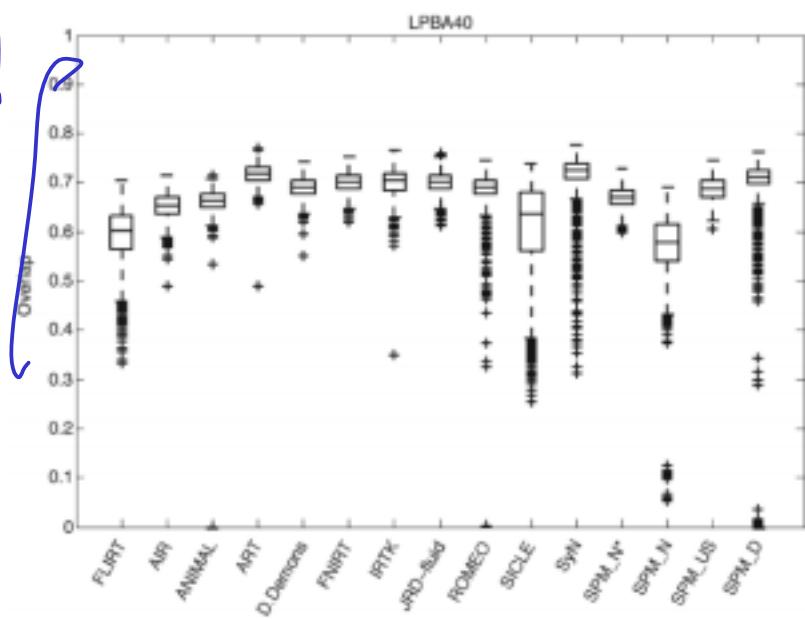
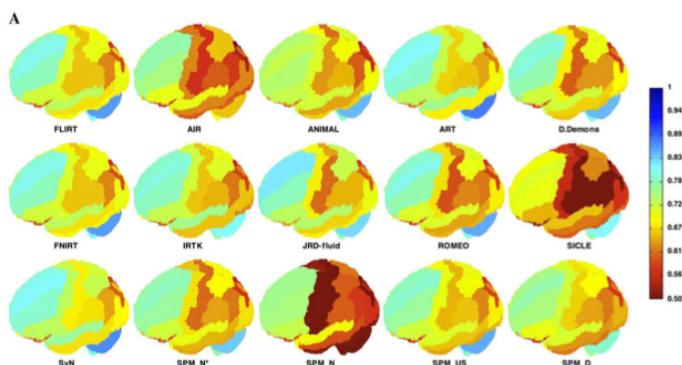
conf: niccai...

|          | Clinical data | Cadavers | Physical phantoms | Realistic simulations | Numerical simulations |
|----------|---------------|----------|-------------------|-----------------------|-----------------------|
| Easy     |               | —        |                   |                       | +                     |
| Control  |               | —        |                   |                       |                       |
| Clinical |               | +        | ←                 |                       | —                     |
| Realism  |               |          |                   |                       |                       |

Neuroimage. 2009 July 1; 46(3): 786–802. doi:10.1016/j.neuroimage.2008.12.037.

### Evaluation of 14 nonlinear deformation algorithms applied to human brain MRI registration

Arno Klein<sup>a,\*</sup>, Jesper Andersson<sup>b</sup>, Babak A. Ardekani<sup>c,d</sup>, John Ashburner<sup>e</sup>, Brian Avants<sup>f</sup>, Ming-Chang Chiang<sup>g</sup>, Gary E. Christensen<sup>h</sup>, D. Louis Collins<sup>i</sup>, James Gee<sup>f</sup>, Pierre Hellier<sup>j,k</sup>, Joo Hyun Song<sup>h</sup>, Mark Jenkinson<sup>b</sup>, Claude Lepage<sup>i</sup>, Daniel Rueckert<sup>m</sup>, Paul Thompson<sup>g</sup>, Tom Vercauteren<sup>n,l</sup>, Roger P. Woods<sup>o</sup>, J. John Mann<sup>a</sup>, and Ramin V. Parsey<sup>a</sup>

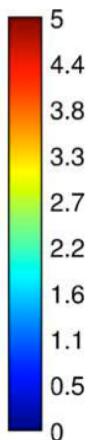
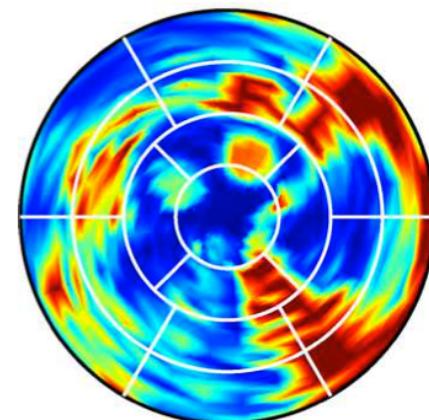
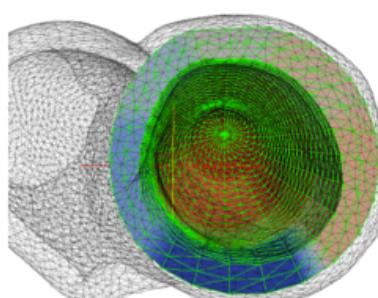
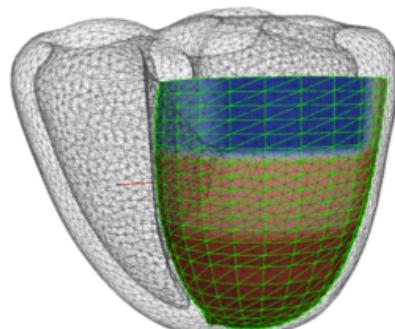
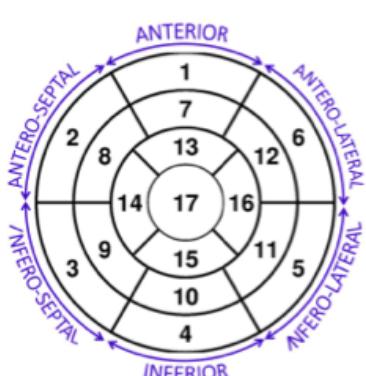


## Validation

(further developed in D. Sarrut's part)

+ challenges

|                  | Clinical data | Cadavers | Physical phantoms | Realistic simulations | Numerical simulations |
|------------------|---------------|----------|-------------------|-----------------------|-----------------------|
| Easy Control     |               | —        | —                 | → +                   |                       |
| Clinical Realism | + ←           |          |                   | —                     |                       |

IEEE Trans Med Imaging. 2016 Aug;35(8):1915-26. doi: 10.1109/TMI.2016.2537848. Epub 2016 Mar 3.**Detailed Evaluation of Five 3D Speckle Tracking Algorithms Using Synthetic Echocardiographic Recordings.***mouvement cardiaque*Alessandrini M, Heyde B, Queiros S, Cygan S, Zontak M, Somphone O, Bernard O, Serresant M, Delingette H, Barbosa D, De Craene M, O'Donnell M, Dhooge J.

Conclusion

$$\hat{T} = \arg \max_{T \in \mathcal{T}} S(I, J, T)$$

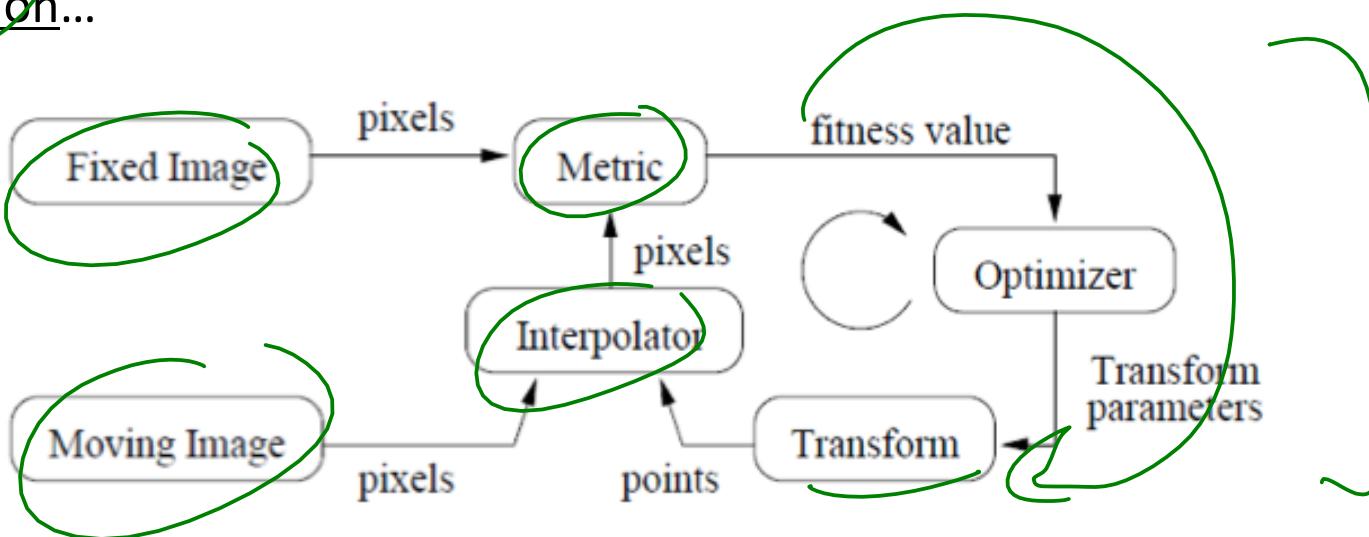
par  $\Delta$   
recalage  
non-rigide

+ Regularisation  
 $Reg(T)$

## Conclusion

$$\hat{T} = \arg \max_{T \in \mathcal{T}} S(I, J, T)$$

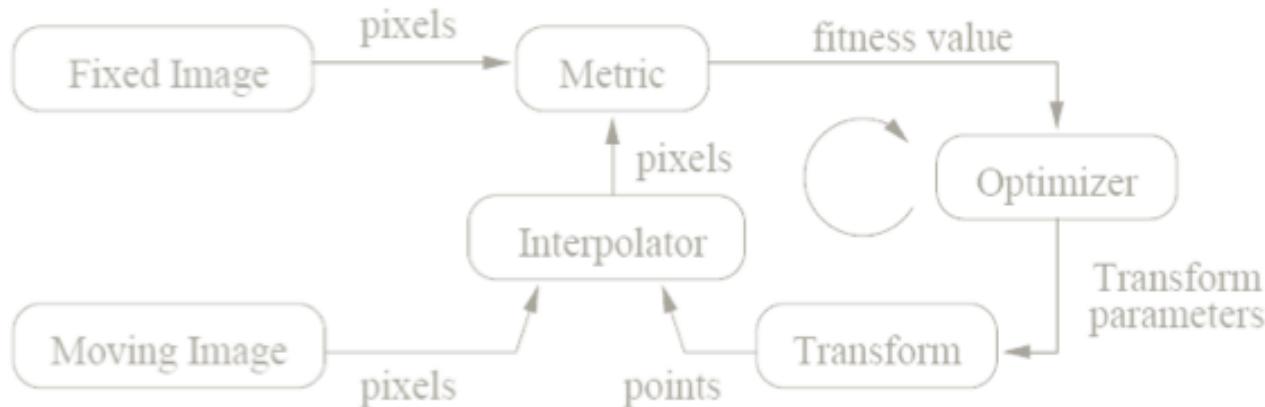
Your own solution...



## Conclusion

$$\hat{T} = \arg \max_{T \in \mathcal{T}} S(I, J, T)$$

Your own solution...



... or existing solutions

- ITK / VTK
- Elastix, Deformetrica, ...





