



Algebraic Reconstruction and beyond in Tomography

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Overview



- Algebraic reconstruction
- Limited angle tomography
- Bayesian Restoration of X-ray images



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Description of X-ray interaction

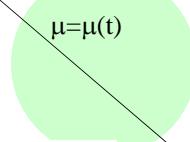


- An X-ray tube emitting monochromatic radiation (from 20 to 150 keV);
- An object absorbing X-rays (absorption + Compton effect);

- A sensor to measure the X-ray intensity.

X-ray tube

A
 $I=I_0$
t



B
Sensor

$$\frac{\Delta N}{N} = -(\tau + \sigma)\Delta x = -\mu\Delta x$$

At the limit for $\Delta x \rightarrow 0$

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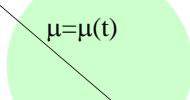


Tomography as a linear problem



$$\int_x^{x+\Delta x} \frac{\Delta N}{N} dx = - \int_x^{x+\Delta x} \mu dx$$

X-ray tube
A
 $I=I_0$
t



B
Sensor

With $x = A$ and $\Delta x = B$, $I_0 = I_A$:

$$I_B = I_0 \cdot e^{- \int_A^B \mu(t) dt}$$

$$\ln \left(\frac{I_B}{I_0} \right) = - \int_A^B \mu(t) dt$$

→ De Beer's law

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μ components



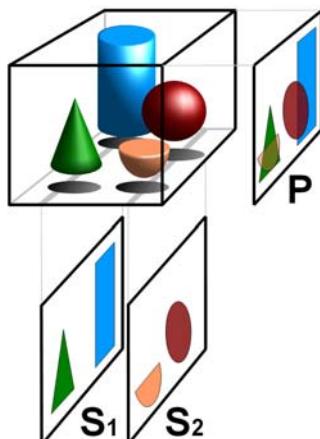
- The X-ray is not monochromatic, but has a (wide) spectrum.
- Absorption is not linear. Lower energy photons are absorbed by soft tissue leaving higher energy photons for bony tissue (Beam hardening).
- Overestimate in few regions (soft tissue) and underestimate in bony regions (underestimate).
- Compton effect does occur.
- Sensors with anti-scatter grid are usually employed.
- **For highly accuracy reconstruction system, in a refinement stage beam hardening could be taken into account.**

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Tomography is a typical inverse problem



We start from N images of MxP pixels each

Volume as a set of parallel planes or voxels.

Interpolation between adjacent planes (Marching cubes)

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 **CT machines** 



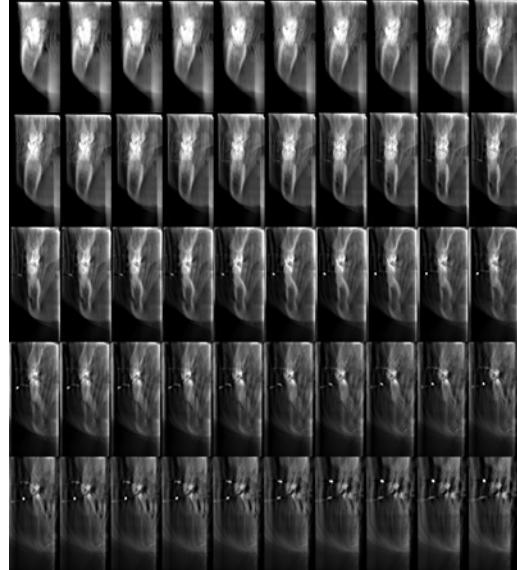
Skyview - Myray



Somaton Emotion - Siemens

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 **Ricostruzione “continua” del volume** 



Go to Matlab visualization
SW: Tomografia3D
M\VFW

**Slices from 11 images 40°,
60kV,10mA**

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Approccio analitico



- E' basato sulla trasformata di Radon:

$$P_\theta(t) = \int_{-\infty}^{\infty} \int_{-\infty}^{\infty} f(x, y) \delta(x \cos \theta + y \sin \theta - t) dx dy.$$

- E sulla sua antritrasformata filtrata (caso fan-beam).

$$f(x, y) = \int_0^\pi Q_\theta(x \cos \theta + y \sin \theta) d\theta$$

$$Q_\theta(t) = \int_{-\infty}^{\infty} S_\theta(w) |w| e^{j2\pi w t} dw.$$

E viene generalizzato per le diverse geometrie: cone-beam, helicoidal.

Richiede che le immagini siano equi-spaziate. Cambiamento di base. Difficoltà a gestire le peculiarità locali (artefatti da metallo, movimento...).



Tomography – discrete case



- Object discretized into a set of voxels;
- Image discretized into a set of pixels;

$$\ln\left(\frac{I_B}{I_0}\right) = - \int_A^B \mu dx$$

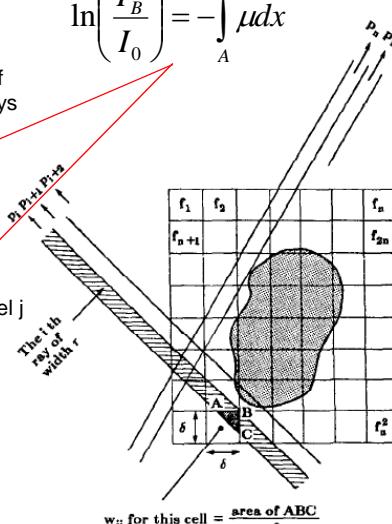
For each measured X-ray intensity (for each pixel of the sensor and for each position), the group of X-rays that hit the pixel are considered together.

Through logarithms:

$$\ln(I_B/I_0) = - \sum_j w_{ij} \cdot f_j \rightarrow$$

- p_i measured value at pixel i .
- w_{ij} length of the path of the X-ray that crosses voxel j and hit pixel i at its center.
- f_j voxel absorption

$$p_i = w_{ij} \cdot f_j$$



$$w_{ji} \text{ for this cell} = \frac{\text{area of ABC}}{\delta^3}$$

A system of linear equations results.



Strategie di ricostruzione



$w_i \cdot f = p_i$ è un sistema di dimensioni enormi, molto sparso. La sparsità non è ancora stata ben esplorata e promette accelerazioni interessanti con schede GPU.

CT Cone beam.

512 immagini su 360 gradi, ciascuna di 512 x 512 pixel su 12bpp (8bpp) = **128MB**
Volume di: 512 x 512 x 512 voxel = **128 M word** (float variables).

N incognite = M misure (equazioni)

Volume di: 256 x 256 x 256 voxel = **16 M word** (float variables).
N incognite << M misure (equazioni)

L'inversione diretta del sistema non è pensabile e algoritmi di ottimizzazione iterativa sono stati introdotti.

**Algoritmi di ricostruzione algebrici
(recentemente diventati popolari)**

$$p_i = w_i \cdot f$$

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Metodo delle proiezioni



S. Kaczmarz, "Angenaherte auflosung von systemen linearer gleichungen," Bull. Acad. Pol. Sci. Lett. A, vol. 6-8A, pp. 355-357, 1937.

K. Tanabe, "Projection method for solving a singular system," Numer. Math., vol. 17, pp. 203-214, 1971

$$w_{11}f_1 + w_{12}f_2 + w_{13}f_3 + \cdots + w_{1N}f_N = p_1$$

$$w_{21}f_1 + w_{22}f_2 + \cdots + w_{2N}f_N = p_2$$

:

$$w_{M1}f_1 + w_{M2}f_2 + \cdots + w_{MN}f_N = p_M.$$

$$p_i = w_i \cdot f$$

Ciascuna misura rappresenta un iper-piano nello spazio N-dimensionale delle N-incognite.

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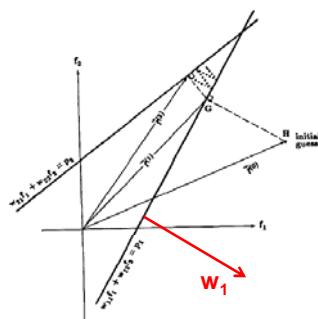
Metodo delle proiezioni - 2



2 incognite -> problema piano -> 2 rette.

$$w_{11}f_1 + w_{12}f_2 = p_1$$

$$w_{21}f_1 + w_{22}f_2 = p_2.$$



First starting with an initial guess, projecting this initial guess on the first line, re-projecting the resulting point on the second line, and then projecting back onto the first line, and so forth. **If a unique solution exists, the iterations will always converge to that point.**

"Kak & Slaney, 2002"

$$\vec{f}^{(i)} = \vec{f}^{(i-1)} - \frac{\vec{w}_i \cdot \vec{p}_i}{\|\vec{w}_i\|^2} \vec{w}_i$$

Residual Direction Length

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Algebraic Reconstruction Technique (ART)



ART iterates through the following steps:

- 1) Choose randomly projection image,
- 2) Choose randomly a pixel in the image,
- 3) Back project the ray in the volume.
- 4) Compute the current absorption along the ray.
- 5) Compute the error on the image (projection error), and smear the error back along the ray.

- It has been applied first in the domain of CAT.

Algebraic representation of the update equations:

$$\mathbf{f}^{(i)} = \mathbf{f}^{(i-1)} - \alpha \frac{(\mathbf{f}^{(i-1)} \cdot \mathbf{w}_i - p_i)}{\|\mathbf{w}_i\|^2} \mathbf{w}_i$$

It is an iterative optimization algorithm of first order!!
 $0 < \alpha < 1$ is a damping factor.

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Problemi con ART



Il calcolo on-the-fly dei coefficienti w diventa proibitivo. Si sostituiscono con valori binari: attraversa non attraversa il voxel i -esimo.

(Damping piccolo -> tempi lunghi) R. Gordon, "A tutorial on ART (algebraic reconstruction techniques)," IEEE Trans. Nucl. Sci., vol. NS-21, pp. 78-93, 1974.

Supersampling (K. T. Smith, D. C. Salmon, and S. L. Wagner, "Practical and mathematical aspects of the problem of reconstructing objects from radiographs," Bull. Amer. Math. Soc., vol. 83, pp. 1227-1270, 1977).

Sostituzione del termine $w \cdot w$ con la lunghezza del raggio che attraversa il tessuto (A. Kak and M. Slaney, Principles of Computerized Tomographic Imaging, Classics in Applied Mathematics, SIAM edition, IEEE Press, New York, 2001).

Inoltre....

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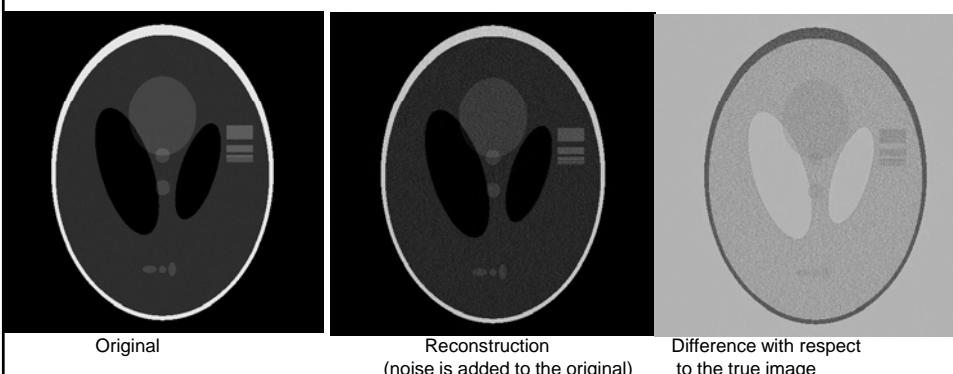
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ART does produce a noisy reconstruction



Results obtained from 300 images equally spaced over 300 degrees. Phantom.



The solution would be to "regularize" the solution considering more than a single ray at a time.

-SIRT (Simultaneous Iterative Reconstruction Technique) updates the absorption coefficients after considering the rays from all the pixels from all the projection images.
Time becomes prohibitive.

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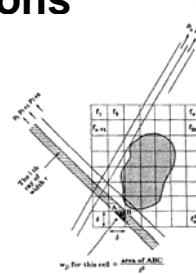


SIRT algebraic interpretations



$$\mu^{k+1} = \mu^k + \lambda \cdot W^T \cdot D \cdot (p - W \cdot \mu^k) \quad SIRT$$

- 1) $W \cdot \mu^k$ projection
- 2) $(p - W \cdot \mu^k)$ projection error
- 3) $W^T \cdot D \cdot (p - W \cdot \mu^k)$ retro - projection of the error



$$p_i = d_{ii} w_i f + v$$

SIRT è un metodo del gradiente (primo ordine).

Si può dimostrare che SIRT minimizza l'errore quadratico pesato sulle proiezioni: $\|D(p - W\mu^k)\|^2 = \min v^2$ pesato dalla matrice diagonale D che contiene gli inversi delle lunghezze al quadrato dei raggi.

Si può anche vedere SIRT come un metodo del gradiente scalato con la matrice D .

Video Matlab sul confronto con i 3 metodi.

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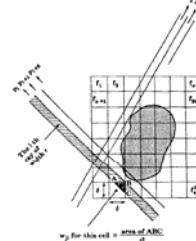


SIRT statistical interpretation



$$\mu^{k+1} = \mu^k + \lambda \cdot W^T \cdot D \cdot (p - W \cdot \mu^k) \quad SIRT$$

- 1) $W \cdot \mu^k$ projection
- 2) $(p - W \cdot \mu^k)$ projection error
- 3) $W^T \cdot D \cdot (p - W \cdot \mu^k)$ retro - projection of the error



$$p_i = w_i f + v$$

SART è un metodo del gradiente (primo ordine). E' riconducibile alla stima a massima verosimiglianza dei parametri, sotto l'ipotesi di rumore Gaussiano a media nulla.

Le osservazioni (misure) vengono pesate con D_{ii} .

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Costo di SIRT e Ruttiman per tomografia



$$\mu^{k+1} = \mu^k + \lambda \cdot W^T \cdot (p - W \cdot \mu^k) \quad SIRT$$

$$\mu^{k+1} = \mu^k + \lambda \cdot (W^T p - W^T W \cdot \mu^k) \quad Ruttiman$$

Ruttiman esplicita l'effetto di un piano ricostruito sugli altri piani.

Il costo della singola iterazione dipende dalla dimensione della matrice W , pari a M righe (numero di dati acquisiti) per N colonne (numero di voxel), con $N = NL \times NL \times NL$, $256^3 = 2^{24}$ voxel.

Sia NI il numero di proiezioni acquisite, di dimensione $NR \times NC$, $M = NI \times NR \times NC$ ($256 \times 512 \times 512 = 2^{26}$ pixel).

Costo di SIRT (si suppone campionato il raggio nei voxel con passo $NL/2$).

Proiezione, $2NL \times \text{pixel}$, costo totale: $2xNL \times M$. **Errore**. Costo pari ad M .

Retropreiezione, $2xNL \times M$

Costo totale. $M \times (4NL + 1) = O(M \times NL) = 2^{36}$

Costo di Ruttiman

Calcolo blurring (effetto di ciascun pixel su ciascun piano): $N \times NL \times NI$

Calcolo errore, aggiornamento: N ciascuno. **Costo totale**: $N \times (NL \times NI + 2) = 2^{42}$

Si preferisce quindi implementare ed utilizzare SIRT, invece di tomosintesi + Ruttiman

La tomosintesi pura (retropreiezione) può essere utilizzata come **inizializzazione** di SIRT.

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SART - Simultaneous Algebraic Reconstruction Technique (Anderson&Kak, 1984)



Three main characteristics:

- 1) Use of a better approximation of the coefficient matrix.
- 2) Updating with a “per epoch” style, analogous to that of SIRT: all the ray in one image at a time are considered.
- 3) Hamming windowing in the longitudinal dimension to emphasize the correction coefficients at the volume center.

In our implementation, we will consider only 1) and 2), as in the most recent literature.

From ART

to:

SART

$$\mu^{(t+1)} = \mu_j^{(t)} + \lambda(p_i - \mathbf{a}_i \cdot \boldsymbol{\mu}) \quad \mu_j^{(t+1)} = \mu_j^{(t)} + \lambda \frac{\sum_i a_{ij} \frac{p_i - \mathbf{a}_i \cdot \boldsymbol{\mu}}{\sum_i a_{ij}^2}}{\sum_i a_{ij}}$$

Where a_{ij} is a weight that measures the “closeness” of the voxel center and the ray through the pixel center. In ART $a_i = \{0, 1\}$

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I coefficienti di SART



Interpolazione dei valori vicini: interpolazione trilineare nel volume e bilineare sulle immagini.
Dato un punto $P=(X,Y,Z)$, definendo (x_k,y_k,z_k) le coordinate del vettore distanza da P al centro del voxel V_k , normalizzate tra 0 e 1, il coefficiente a_k relativo a V_k viene calcolato come:

$$a_k = (1 - x_k)(1 - y_k)(1 - z_k)$$

con:

$$\mu^{(t+1)} = \mu_j^{(t)} + \lambda \frac{\sum_i a_{ij} \frac{p_i - \mathbf{a}_i \cdot \boldsymbol{\mu}}{\sum_i a_{ij}}}{\sum_i a_{ij}}$$

$$\boldsymbol{\mu}^{k+1} = \boldsymbol{\mu}^k + \lambda \cdot \mathbf{W}^T \cdot \mathbf{D} \cdot (\mathbf{p} - \mathbf{W} \cdot \boldsymbol{\mu}^k) \quad \mathbf{D}_{ii} = \frac{1}{\sum_j a_{ij}^2}$$

Sceglieremo di interpolare nel **volume** (in CUDA viene realizzata in HW) e quindi “**pettineremo**” le immagini, tracciando i raggi sui quali calcolare l’assorbimento solamente in corrispondenza dei centri dei pixel.



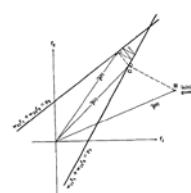
OS-SART



Invece di aggiornare i coefficienti dopo la singola immagine (SART puro) o dopo tutte le immagini (SIRT), si sceglie un compromesso aggiornando dopo avere esaminato K immagini < M . K rappresenta un sub-set delle immagini.

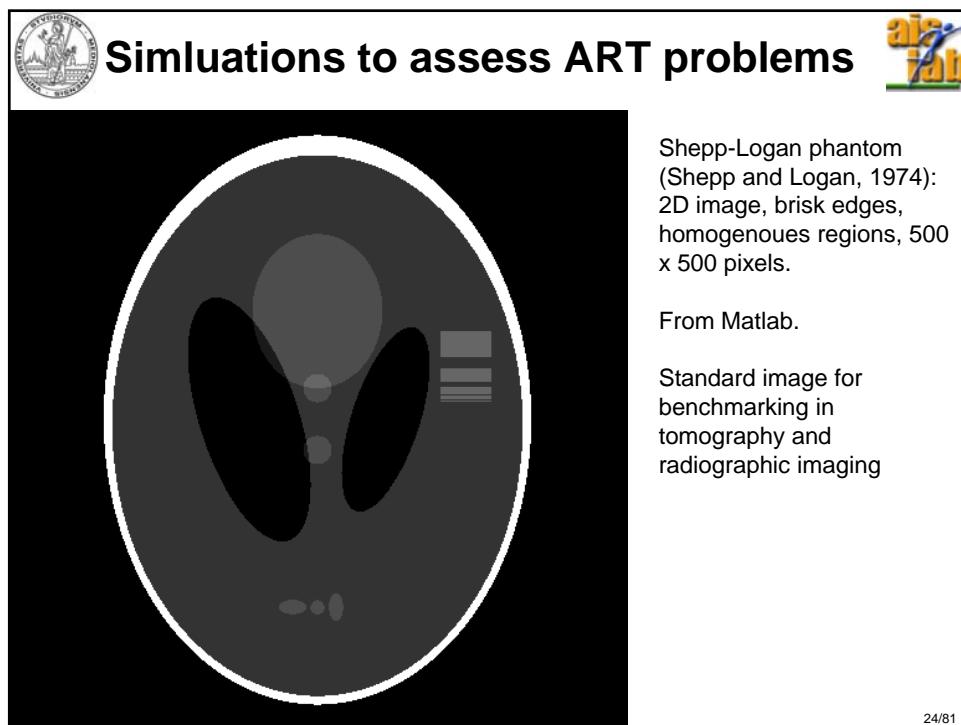
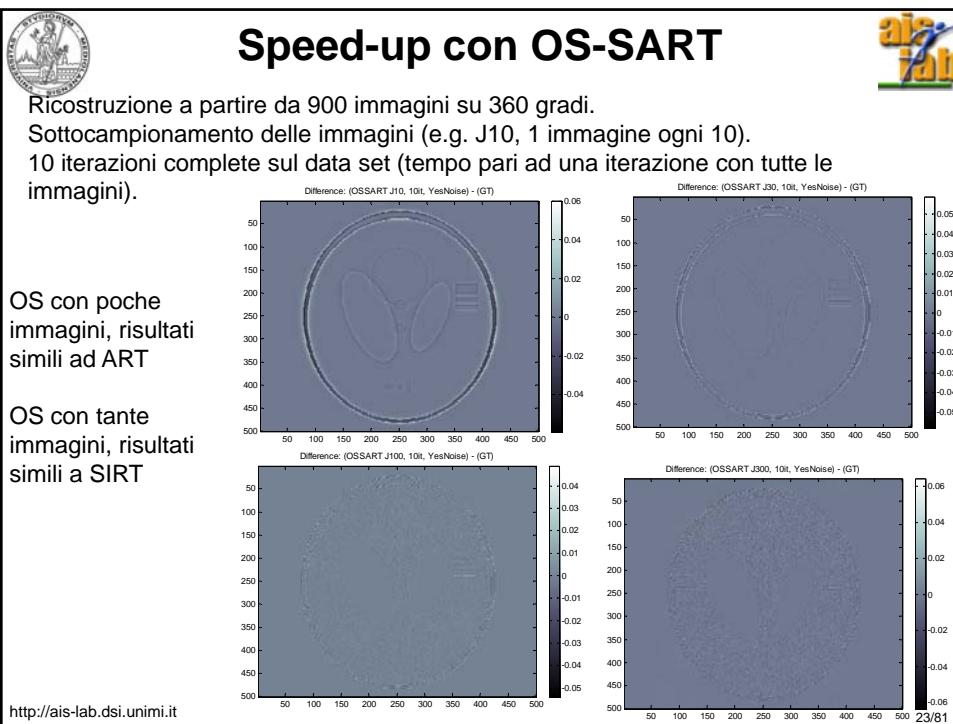
che può essere scelto in diverso modo (ordered-subset, OS):

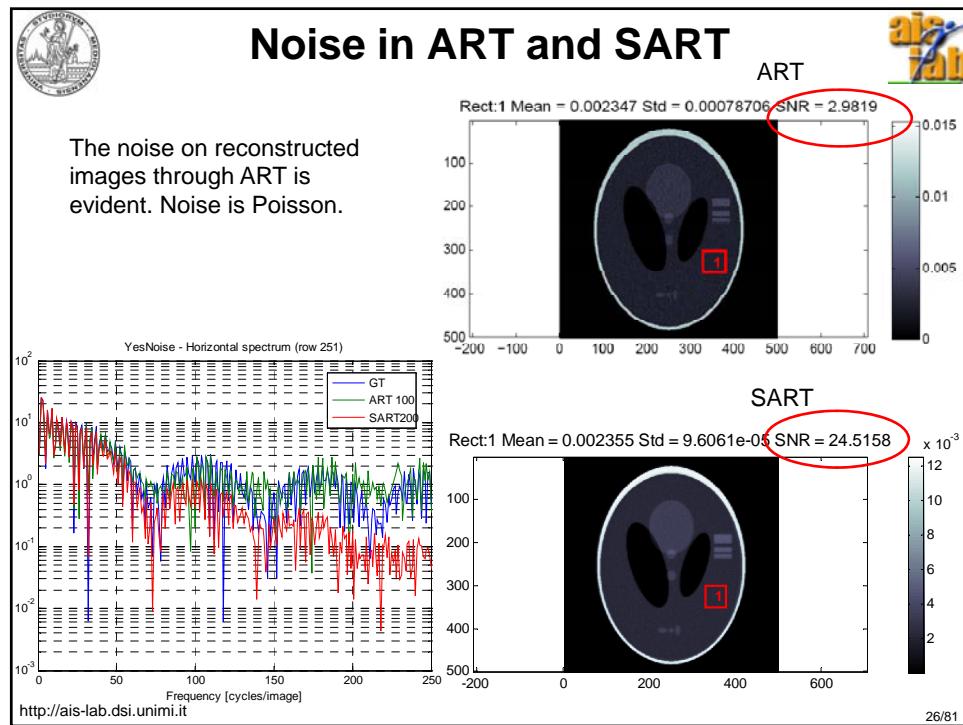
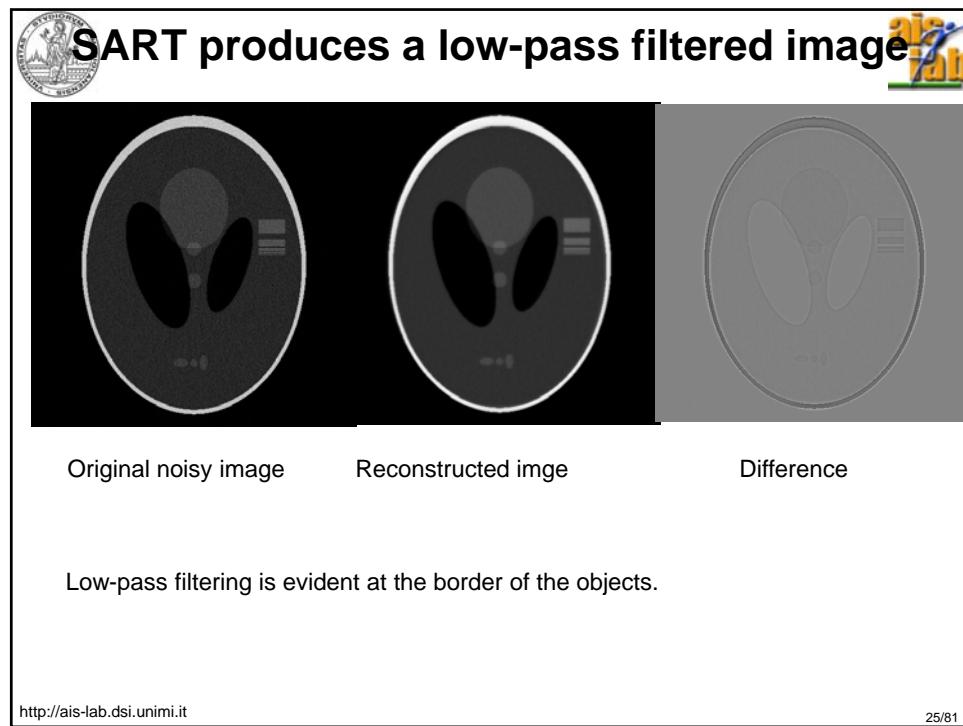
- Sequenziale.
- Massima ortogonalità tra le immagini (projection algorithm)
- Random.

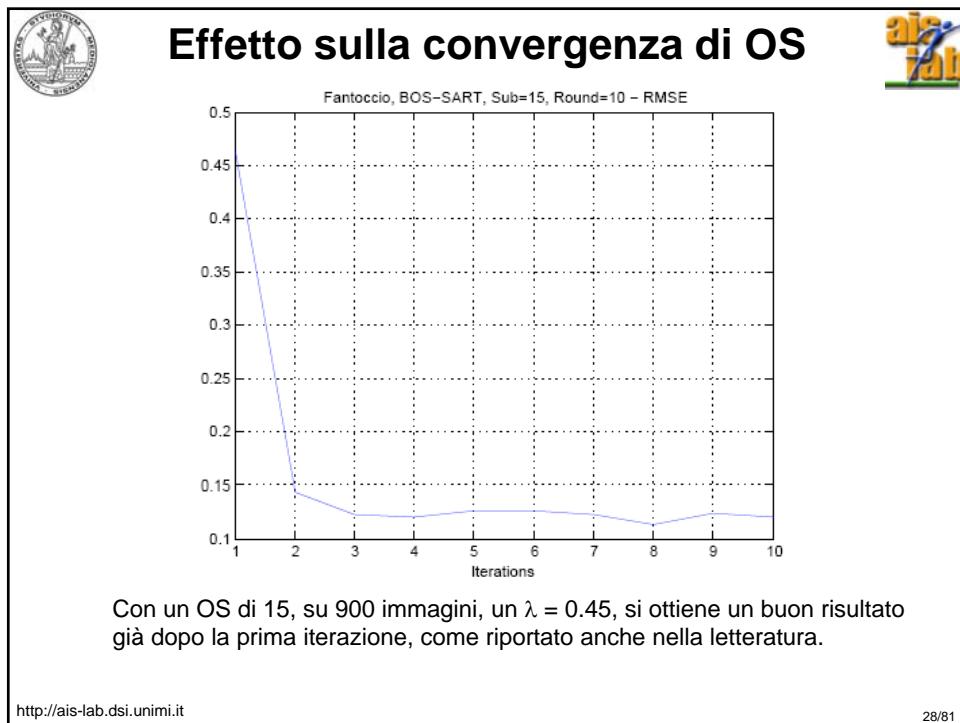
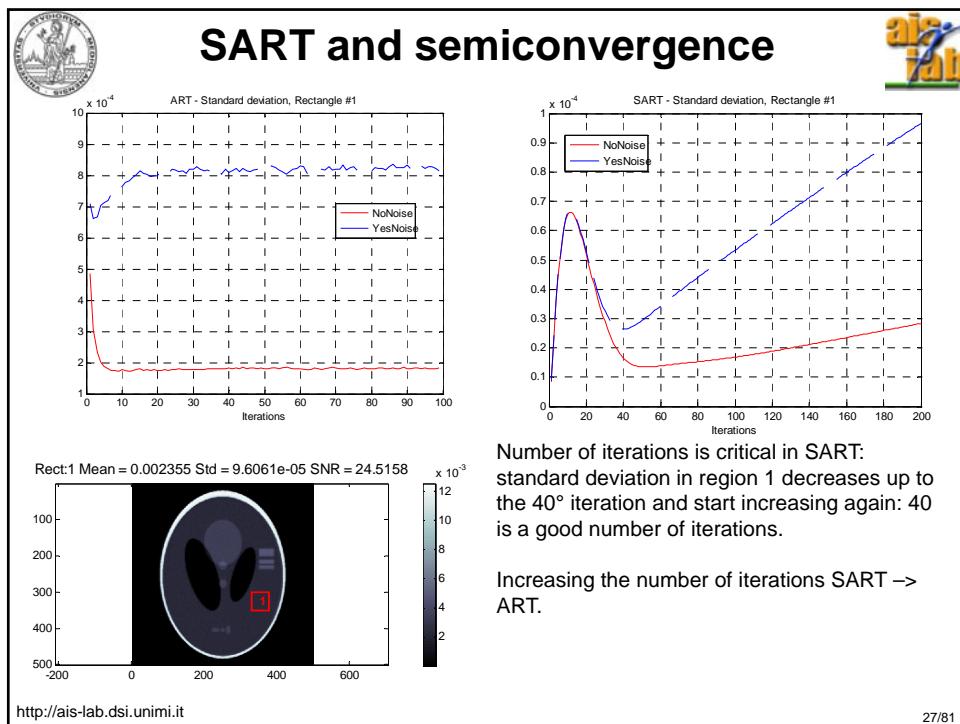


Il parametro di damping può rivelarsi critico, da impostare sperimentalmente.

$$\mu^{(t+1)} = \mu_j^{(t)} + \lambda \frac{\sum_i a_{ij} \frac{p_i - \mathbf{a}_i \cdot \boldsymbol{\mu}}{\sum_i a_{ij}}}{\sum_i a_{ij}}$$









Summary on SART



- SART produces a more homogenous image, with noise filtering.
- Its weak points are the convergence time (number of iterations) and the low-pass filtering effect.
- Acceleration can be achieved through OS techniques.
- GPU processing is mandatory to obtain results in a reasonable amount of time.
- Low-pass filtering can be limited by limiting the number of iterations or adopting a regularization approach (that can be computationally heavy).
- Parameters to be optimized empirically:
 - Number of images in the OS
 - Value of damping factor, λ
 - Voxel dimension.

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Overview



- Algebraic reconstruction
- **Limited angle tomography**
- Bayesian Restoration of X-ray images



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 **Motivation for limited angle tomography** 

Dose reduction
Costs reduction

MAMMOMAT - Siemens



Hyperion CEFLA (MyRay)

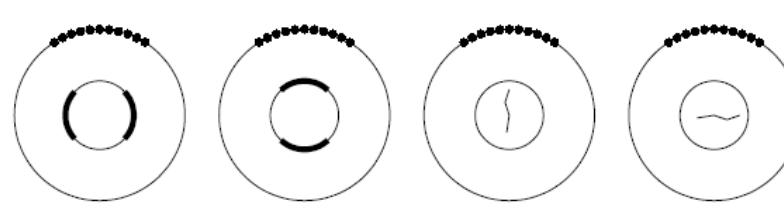


Multiple images over 180 degrees -> 3D volumetric images
<http://ais-lab.dsi.unimi.it> Multiple images over a smaller angle -> 2.5D volumetric images

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 **Which are the limitations?** 

- Per effetto dell'allargamento dell'angolo di acquisizione è possibile allargare l'intervallo delle frequenze osservabili (diversi orientamenti) nel volume ricostruito.
- Per contro, un allargamento dell'angolo di acquisizione porta ad una diminuzione della risoluzione nei piani ortogonali rispetto alla direzione di acquisizione principale.
- E' necessario trovare il giusto compromesso...



Visible edge Indetectable edge Visible crack Indetectable crack

Figure 3. Illustration of discontinuities that can and cannot be recovered based on limited-angle projection data. The black dots denote the locations of the x-ray source in the acquisition of the projection data. For each such location, the detector is thought to be located opposite to the x-ray source as in figure 1. [Somersalo et al., 2003](#)

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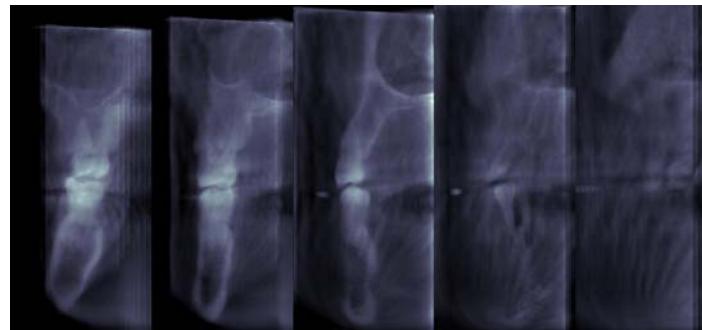


Which are the results of using limited angle?



Limited resolution on depth -> parallel sections

- Mammografia (sezioni parallele al torace)
- Dentale (sezioni parallele al canale mandibolare)



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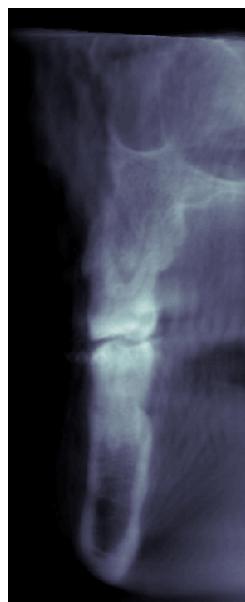
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Standard configuration



- Acquisition angle = -30° ... +30°
- Acquisition time = 60s
- Xray duration = 26s (1,5s x 11)
- Reconstruction time =
 - 118s (OS-SART, 5 iterations)
 - Real time (a few s, tomosynthesis)
- Number of projections = 11
- KV 60-84, mA 1-10, set from panoramic image
- Enlarging factor = 1.43
- Images size = 1536 x 662 pixel
 - Useful images size = 1536x**562** pixel
- Pixel size 0,096mm.
- Volume = 5cm x 4cm x 10.5cm
- Volume dimension in voxels = 50 x 267 x 700
- Voxel dimension: 1mm x 0.15mm x 0.15mm
- Automatic identification of the volume in the panoramic image.
Volume is centered slightly inside for frontal teeth.

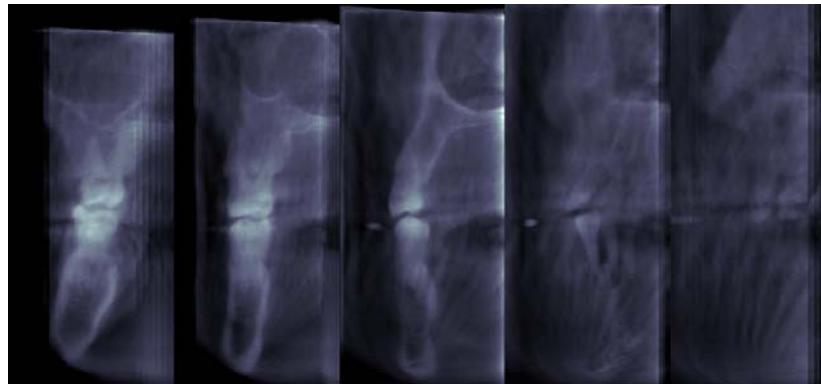


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Artefacts with band shape and truncation



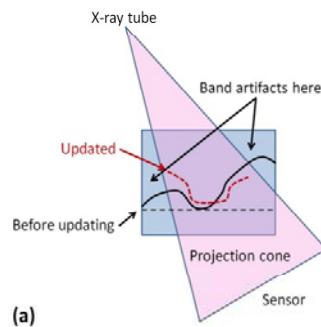
Five sections from SART. Notice vertical lines on the right of the image and the whitening of the image.



Problems analysis

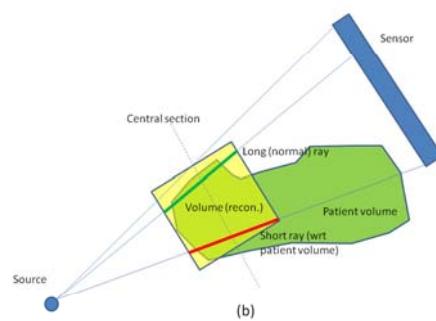


Band shape artifacts due to the cone associated to each projection only partially crosses the volume



(a)

X-ray absorption outside the volume



(b)

We can solve these problems modifying the optimization engine used in SART



Solution for band artifacts

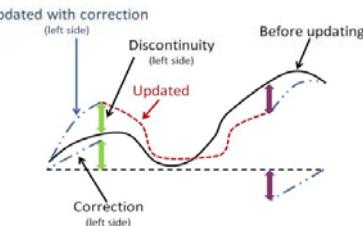
$$\mu^{k+1} = \mu^k + \lambda \cdot W^T \cdot D \cdot (p - W \cdot \mu^k) \quad SART$$

Local equalization (inspired by W. Zbijewski, F. J. Beekman, 2004)

Linear correction of the absorption outside the projecting cone:

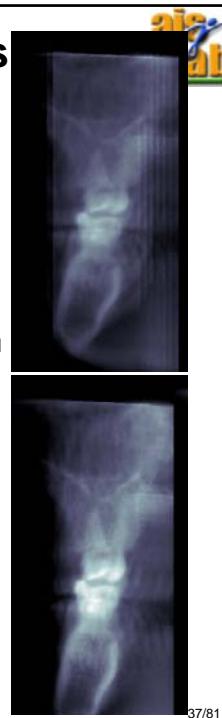
$$x_{ijk} \leftarrow x_{ijk} + \Delta_k \cdot (1 - \delta_{ijk}/\Delta_k)$$

δ_{ijk}/Δ_k ration between 1) the distance of voxel ijk (outside the projecting cone) and the volume border and 2) between the projecting cone and the volume border



(b)

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Solution for band artifacts

$$\mu^{k+1} = \mu^k + \lambda \cdot W^T \cdot D \cdot (p - W \cdot \mu^k) \quad SART$$

$$D_{ii} = \frac{1}{\sum_j a_{ij}^2}$$

Estimate of absorption outside volume in voxels with abnormal response. (Zhang et al., J. Comput. Ass. Tomography 2009)

Modify the weights:

The longer the ray, the more it receives error, the more reliable is the update.

→ In least square sense, the pixel associated to that ray is more reliable → higher weight.

$$\mu^{k+1} = \mu^k + \lambda \cdot W^T \cdot D^{-1} \cdot (p - W \cdot \mu^k)$$

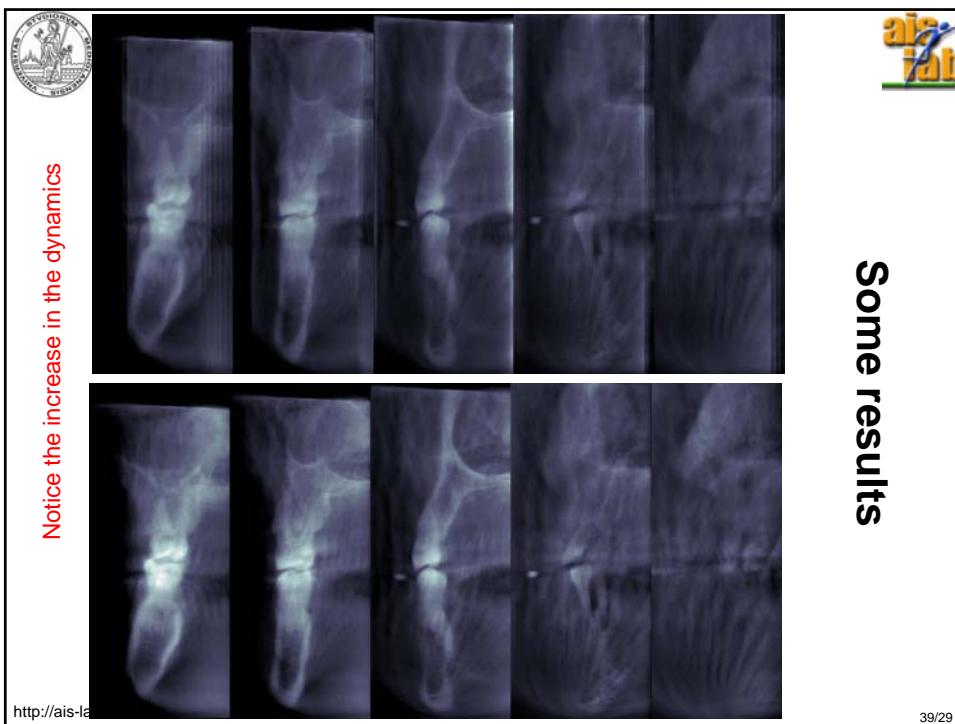
A new weighting scheme for limited angle arises
(Frosio and Borghese, in preparation)

$$p_i = w_i \cdot f$$

In this situation the voxels are updated more strongly for a ray that crosses a thicker volume region because it is more reliable that the measured absorption is due to the voxel crossed and not to outside ones (it is a different philosophy).

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**Ottimizzazione del sistema
patient-based**

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lab

- Utilizzo di un modello anatomico per il calcolo dell'esposizione ottimale immagine per immagine.
- Modello parametrizzato automaticamente a partire dalla panoramica del paziente.
- Considerazione dell'effetto del metallo (amalgama, protesi) sia nell'esposizione che nella formulazione della traiettoria.

Risultato: Optimized images (higher contrast and higher dynamics).

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Compensazione del movimento (eliminazione blurring da movimento)

Identificazione del movimento mediante identificazione di feature sull'immagine radiografica.

Ricerca di feature adatte per i vari distretti mandibolari.

Determinazione del movimento. Affidabilità della stima.

Studio di fattibilità (video? X-ray? Modifiche alla macchina?)

Obiettivo: Correzione del "blurring da movimento"

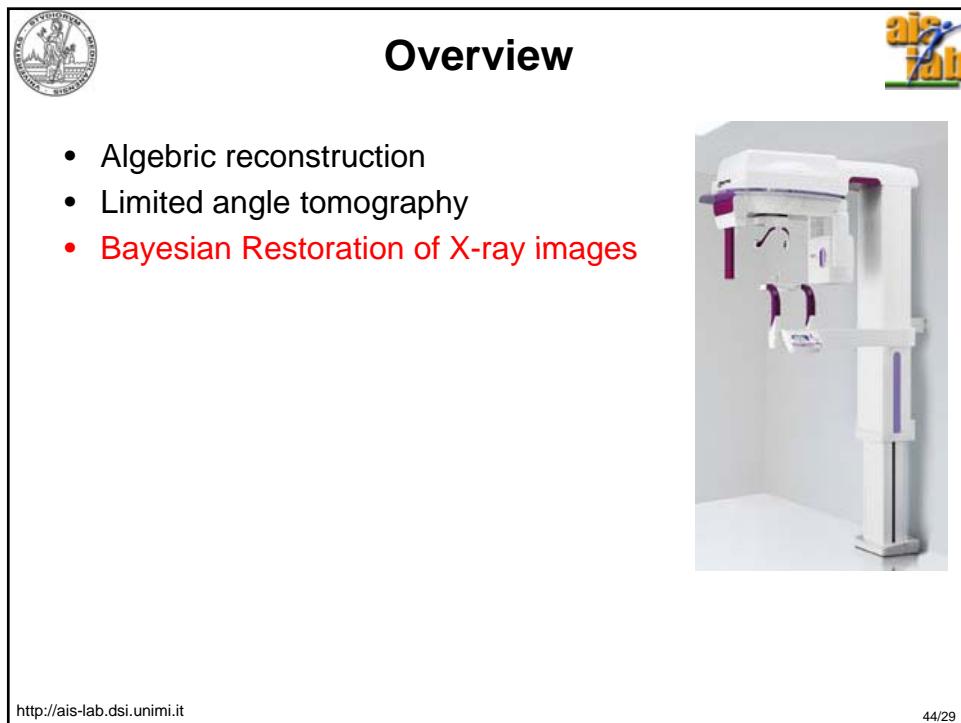
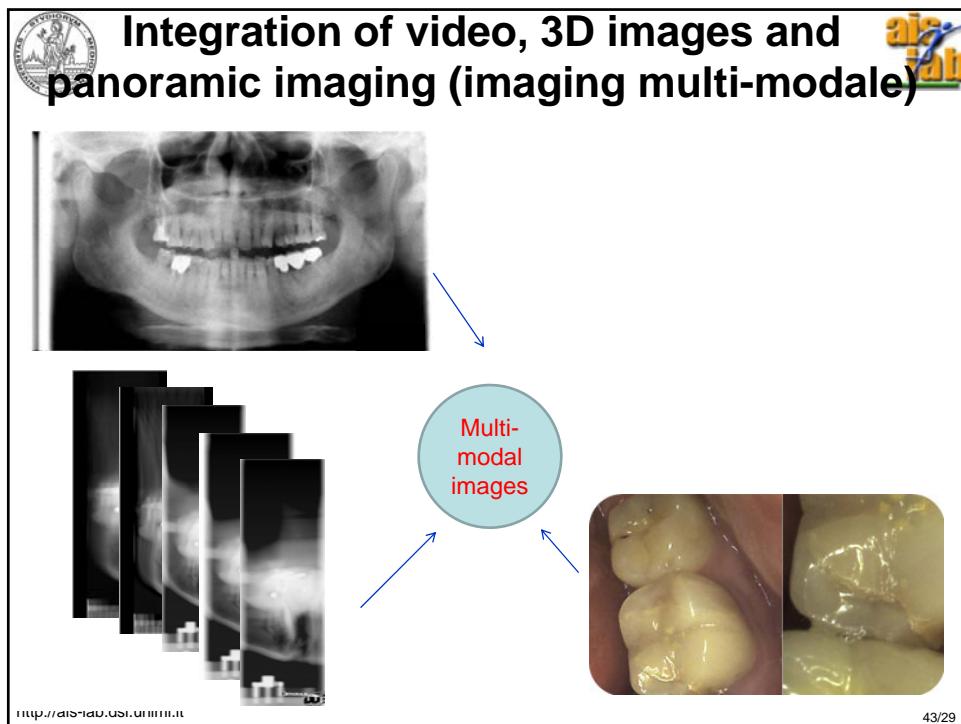
Possibilità di correggere anche errori di posizionamento macchina / taglio immagini.

FIG 4

Borghese, Nanni, Rotondo
US Patent 0008559, Jan 2010

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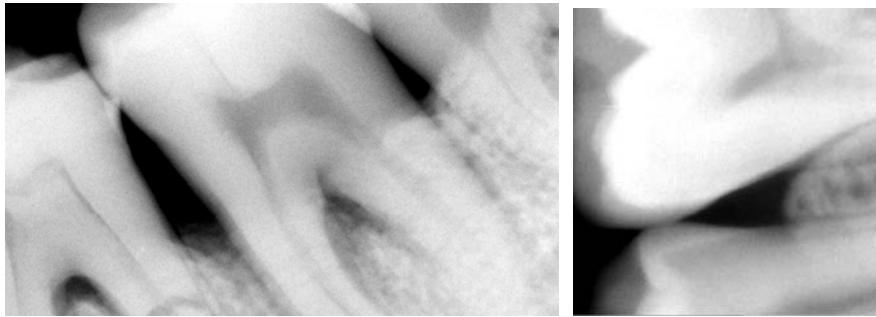




Denoising of a radiographic image



Typical images (digital radiographies @ 12bpp, 1.5Mpixels – negative images), Poisson noise.



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Statistical formulation of image restauration



Measuring an image g_{noisy} taken from an object, f , (e.g. perspective image, transmission image): $g = Af + \text{noise} \rightarrow f?$

(A may or may not contain the blurring of the measuring tool, point spread function)

Each pixel is considered an independent process (white noise). For each pixel therefore, we want to maximize:

$$p(g_{noisy}; f)$$

Being the pixels independent, the total probability can be written in terms of product of independent probabilities:

$$L(g_{noisy,i}; f_i) = \prod_{i=1}^N p(g_{noisy,i}; f_i)$$

$L(\cdot)$ is called likelihood function.

Determine $\{f_i\}$ such that L is maximized. Negative log-likelihood is usually considered to deal with sums:

$$J_0(g_{n,i}; f_i) = -\sum_{i=1}^N \ln(p(g_{n,i}; f_i))$$

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Gaussian & Poisson cases



$$\text{Noise}_i = \|Af_i - g_{ni}\|$$

We know the statistical distribution of the noise -> we now the statistical distribution of the second term.

In case of (normalized) Gaussian distribution therefore:

$$J_0 = -\ln(p(g_n, f)) = -\ln \left\{ \left(\frac{1}{\sqrt{2\pi\sigma^2}} \right)^m e^{-\frac{1}{2\sigma^2} \|g_n - Af\|^2} \right\} = \text{cost} + K \|g_n - Af\|^2$$

The minimization of this function leads to:

$A^T A f = A^T g_n$ **That is the least squares solution.**

The same line of reasoning for Poisson noise leads to the KL divergence:

$$J_0 = -\ln(p(g_n, f)) = D_{KL}(g_n, Af) = \sum_i g_{n,i} \ln \left(\frac{g_{n,i}}{Af_i} + Af_i - g_{n,i} \right)$$

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Can we solve any problem?



$$g = Af + noise \Rightarrow g = g_o + noise$$

Going in the frequency domain: $G(w) = G_o(w) + W(w) = K(w)F_0(w) + W(w)$

If we suppose that $W(w) \neq 0$ outside the support of $K(w)$, the problem is ill-posed.

And if we solve the linear system above, in Fourier space, the following nasty things happen:

$$G(w) - W(w) = K(w)F_0(w) \rightarrow$$

No solution exists as noise $W(w)$ cannot be reproduced by $F_0(w)$.

Moreover the solution is not unique as any function defined outside the support of $F(w)$ can be added.

As $K(w)$ decreases to zero with a transition bandwidth of finite width, noise components tend to be largely amplified, that produce spurious oscillations.

In case of filtering $A = I$, the restored image would be equal to the noisy one.

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Observations



Solution has a single global minimum in both discrete and continuous cases (Shepp and Vardi, 1982).

Solution has many local minima in both Poisson and Gaussian cases, in the continuous case (cf. Malthei, 1993 for Poisson case).

Is the solution interesting? No, as it also incorporates noise. Therefore, semi-convergence has been proposed (Bertero and Boccacci, 1998).



The Bayesian framework



We assume that the object f is a realization of the “abstract” object F that can be characterized statistically.

The probability $p(g_n | f)$ becomes a conditional probability:

$$J_0 = p(g_n | f = f^*)$$

Under this condition, the probability of observing f and g_n (joint probability) can be written as the product of the probabilities:

$$p(g_n, f) = p(g_n | f)p_f$$

As we are interested in determining f , we have to write the conditional probability of f given g_n through the Bayes theorem:

$$p(f | g_n) = \frac{p(g_n | f)p_f}{p_{g_n}} = L(g_n; f) \frac{p_f}{p_{g_n}}$$



MAP estimate



$$p(f | g_n) = \frac{p(g_n | f)p_f}{p_{g_n}} = L(g_n; f) \frac{p_f}{p_{g_n}}$$

We can find f here by maximizing $p(f | g)$, with respect to f . **MAP estimate**. We explicitly observe that the marginal distribution of p_{g_n} is not dependent on f .

Again, it is more efficient to go to the logarithms:

$$f = \operatorname{argmax}\{p(f | g_n)\} =$$

$$\operatorname{argmin}_f -\{\ln(p(g_n | f)p_f)\} = \operatorname{argmin}_f -\{\ln(p(g_n | f)) + \ln(p_f)\}$$

Likelihood = adherence to the data A-priori



Tikhonov regularization



$$f = \operatorname{argmin}_f -\{\ln(p(g_n | f)p_f)\} = \operatorname{argmin}_f -\{\ln(p(g_n | f)) + \ln(p_f)\}$$

Gaussian noise model
Squared shape for the log a-priori term

$$K \sum_i \|g_{n,i} - Af_i\|^2$$

$\|Pf\|^2$

$$f = \operatorname{argmin}_f \sum_i \|g_n - Af\|^2 + \lambda \|Pf\|^2$$

It is a quadratic cost function

This leads to the classical Tikhonov formulation

$$f = A^T A g_n - A^T A f + \lambda P^T P f \quad \text{Poggio and Girosi, 1990}$$

What happens when noise is Poisson



$$f = \arg \min_f -\{\ln(p(g_n | f)p_f)\} = \arg \min_f -\{\ln(p(g_n | f)) + \ln(p_f)\}$$

Poisson noise model
Squared shape for the a-priori term

$$\sum_i g_{n,i} \ln\left(\frac{g_{n,i}}{Af} + Af_i - g_{n,i}\right)$$

$$\|\lambda Pf\|^2$$

$$f = \arg \min_f \sum_i g_{n,i} \ln\left(\frac{g_{n,i}}{Af} + Af_i - g_{n,i}\right) + \lambda \|Pf\|^2 \quad \text{Regularization}$$

No analytical solution -> non-linear optimization

Non-quadratic a-priori



$$f = \arg \min_f -\{\ln(p(g_n | f)p_f)\} = \arg \min_f -\{\ln(p(g_n | f)) + \ln(p_f)\}$$

Gaussian noise model

$$\sum_i \|g_{n,i} - Af_i\|^2$$

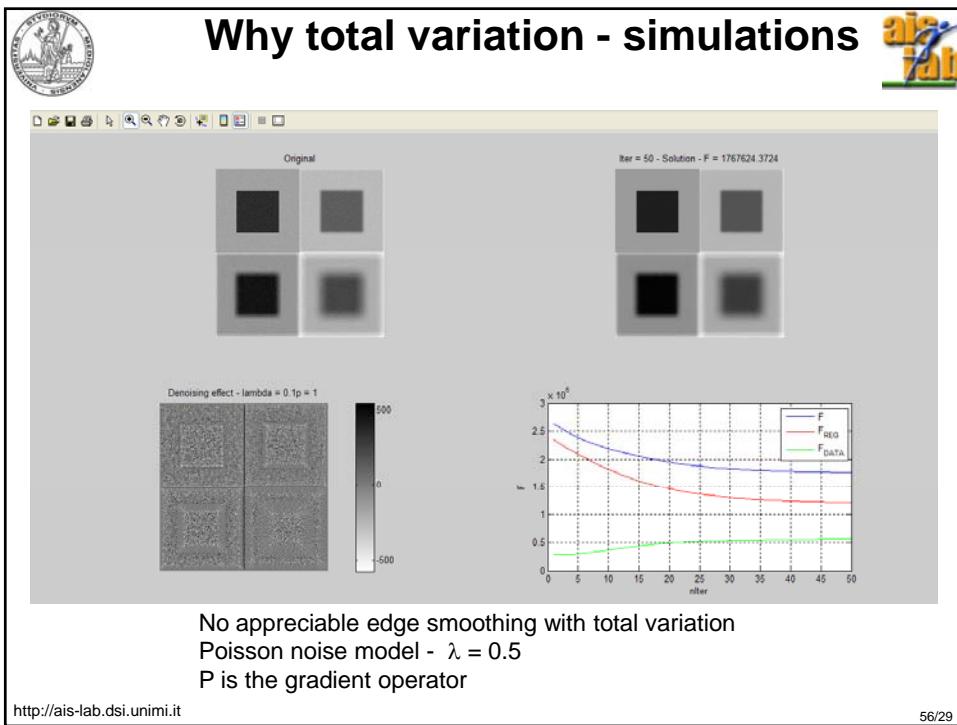
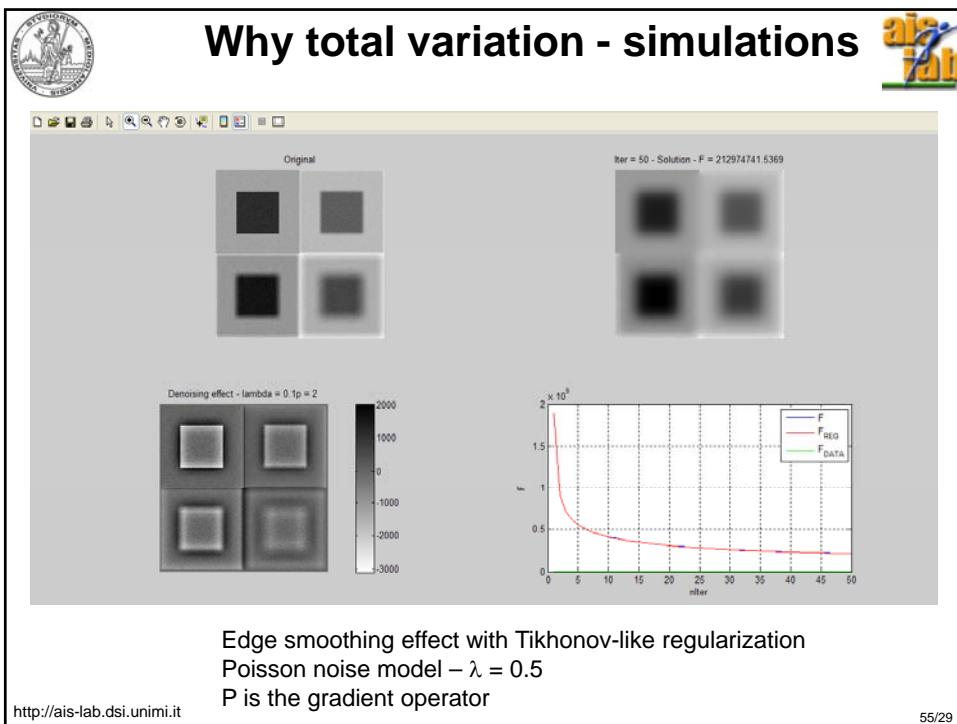
Total variation

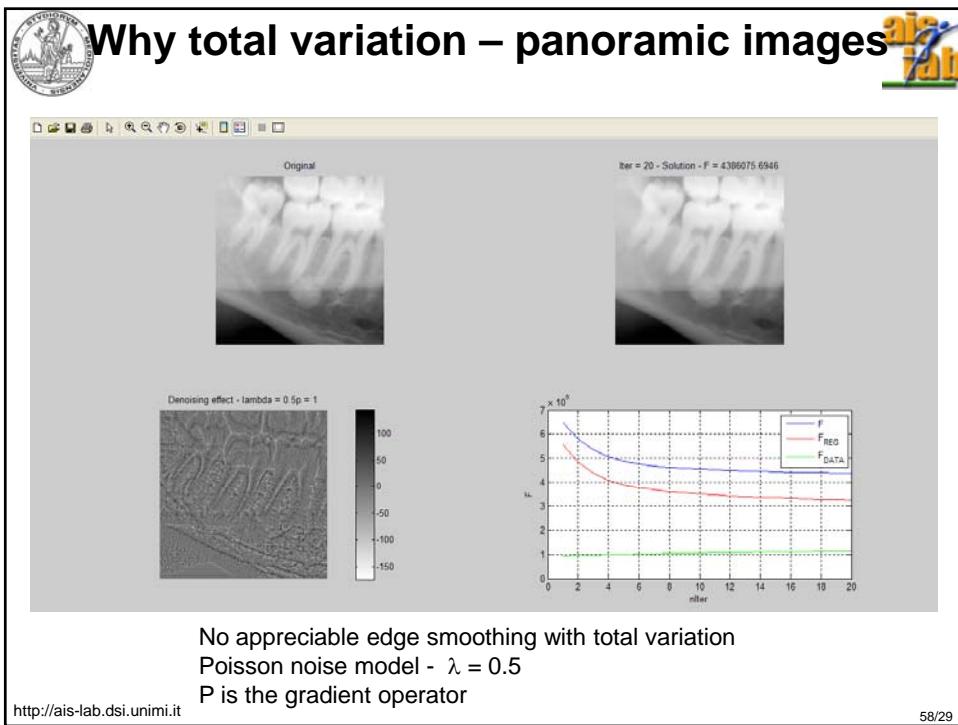
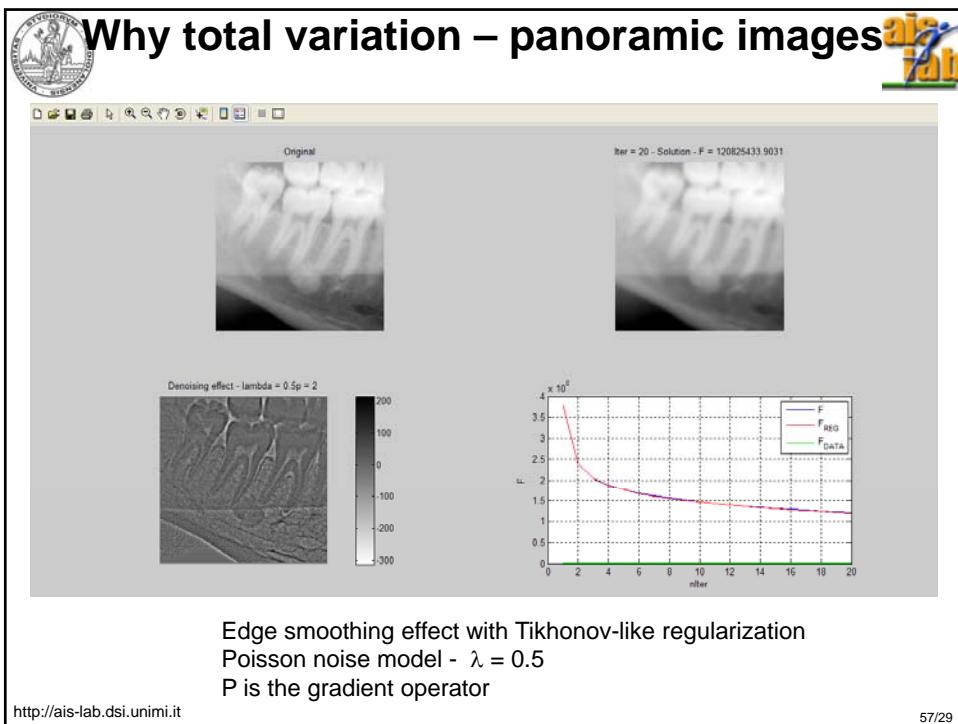
$$\sum_i \sqrt{\sum_{w=1}^W p_{f_w}^2}$$

A-priori is a gradient and it is expressed in ℓ_2 norm

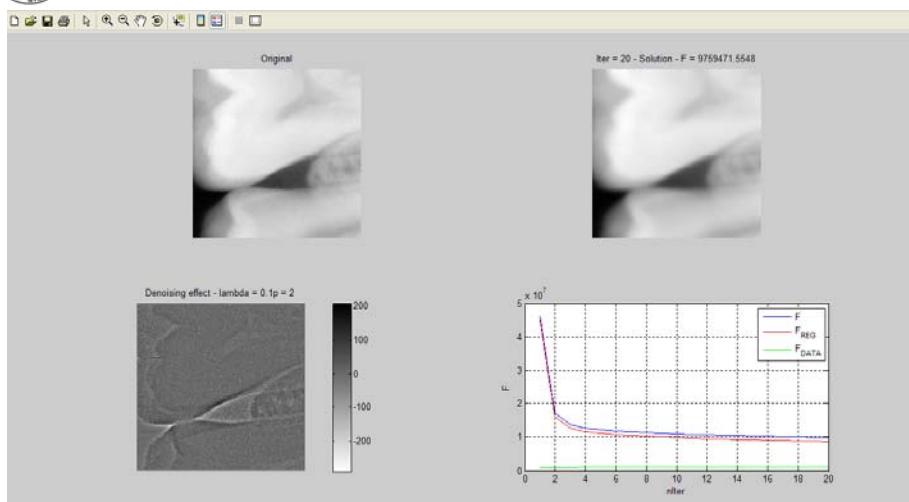
$$p_f = \sqrt{\sum_i (f_{x,i}^2 + f_{y,i}^2 + f_{z,i}^2)}$$

$$f = \arg \min_f \sum_i \left(\|g_n - Af\|^2 + \lambda \sqrt{\sum_1^W p_{f_w}^2} \right)$$





Why total variation- endo-oral images



Edge smoothing effect with Tikhonov-like regularization

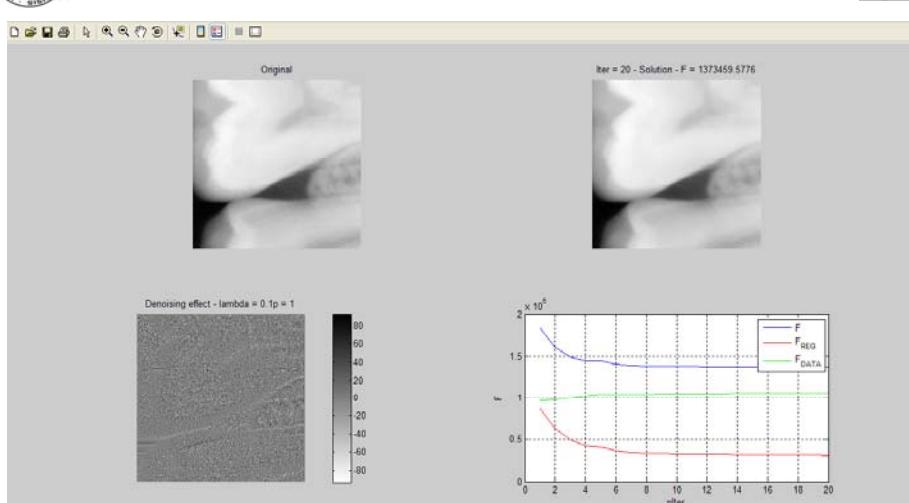
Poisson noise model - $\lambda = 0.1$

P is the gradient operator

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Why total variation – endo-oral images



No appreciable edge smoothing with total variation

Poisson noise model - $\lambda = 0.1$

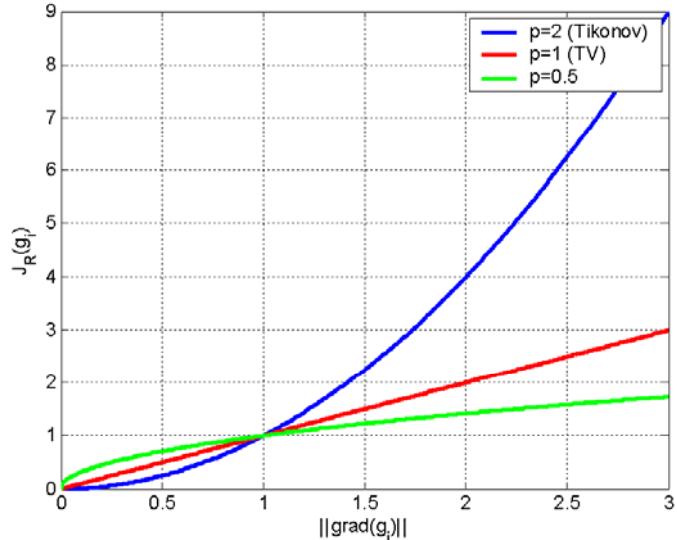
P is the gradient operator

<http://ais-lab.dsi.unimi.it>

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Cost introduced by the regularization term



Cost increases quadratically with the local gradient in Tikhonov

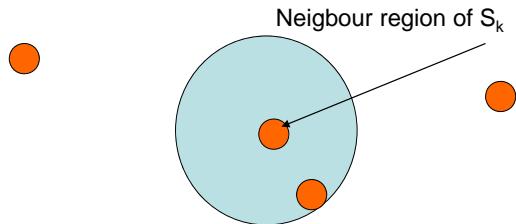


A-priori



We can insert in the a-priori term all the desirable characteristic of the image: local smoothness, edges, piece-wise constancy,....

The idea of defining a neighboring system is a natural one:



Images have a natural neighbouring system: the pixels structure. We want to consider the local properties of the image considering neighboring pixels (in particular differential properties - our vision system is particularly tuning to gradients both spatial and temporal). Ideas have been borrowed from physics.



Neighboring System



Let P be the set of pixels of the image: $P = \{p_1, p_2, \dots, p_P\}$

The neighboring system defined over P, S , is defined as $H = \{\mathcal{N}_p | p, \forall p \in P\}$, that has the following properties:

An element is not a neighbour of itself: $p_k \notin \mathcal{N}_{pk}$

Mutuality of the neighboring relationship: $p_k \in \mathcal{N}_{pj} \leftrightarrow p_j \in \mathcal{N}_{pk}$

(S, P) constitute a graph where P contains the nodes of the graph and S the the links.

Depending on the distance from p , different neighboring systems can be defined:

	o	
o	x	o
	o	

First order neighboring System

4-neighboring System

o	o	o
o	x	o
o	o	o

Second order neighboring System

8-neighboring System

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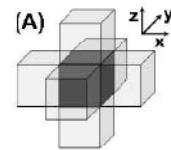
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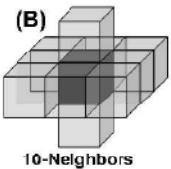
Clique



Borrowed from physics.



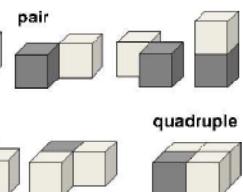
6-Neighbors System



10-Neighbors System

single

pair



triple

quadruple

A clique C , for (S, P) , is defined as a subset of sites in S .

We can consider ordered sets of voxels, that are connected to p through S .

Types of cliques: single-site, pairs of neighboring sites, triples of neighboring sites,... up to the cardinality of \mathcal{N}_p

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Markov Random Field



Given a neighbouring system, S , and a set of pixels, P , we can define a set of random values, $\{f_k(p)\}$ for each element defined by S , that is in \mathcal{N}_p . Therefore we define a **random field**, \mathcal{F} , over S :

$$\mathcal{F}(\mathcal{N}_p) = \{f_k(m) \mid m \in \mathcal{N}_p\} \quad \forall p$$

Under the Markovian hypotheses:

- | | |
|--|--------------|
| 1) $P(f(p)) \geq 0 \quad \forall p$ | Positivity |
| 2) $P(f(p) \mid g(P - \{p\})) = P(f(p) \mid g(\mathcal{N}_p))$ | Markovianity |

2 expresses the fact that the probability of p assuming a certain value f (e.g. a certain gradient), is the same considering all the pixel of P but p , or only the neighbor pixels, that is the value of f depends only on the gray value of the pixels in \mathcal{N}_p .

The random field \mathcal{F} is named **Markov Random Field**.



Energy in a Markov Random Field



A “potential” function, $\phi(f)$, can be defined for a MRF. This is a scalar value that is a function of the random value associated to the pixels for all the possible elements of a clique:

$$\phi_c(f) = \sum_{j \in c} f(p_j)$$

If we consider all the possible cliques defined for each element p , we can define a potential energy function associated to the MRF:

$$U(f) = \sum_{c \in C} \phi_c(f)$$

The higher is the potential energy, the lower is the probability that the set of random values of the elements of the cliques is realized, that is the higher is the penalization for the associated configuration.



Gibbs prior



If we consider all the possible cliques defined for each element p, we can define a potential energy function associated to the MRF:

$$U(\mathbf{f}) = \sum_{c \in C} \phi_c(\mathbf{f})$$

The higher is the potential energy, the lower is the probability that the set of random values of the elements of the cliques is realized, that is the higher is the penalization for the associated configuration.

This is well captured by the Gibbs distribution, that describes the probability of a certain configuration to occur. It is a function exponentially decreasing of U:

$$P(\mathbf{f}) = \frac{1}{Z} e^{\left\{-\frac{1}{\beta} U(\mathbf{f})\right\}}$$

$P(\mathbf{f})$ is a Gibbs random field, Hammersley-Clifford theorem (1971). β regulates the decrease in probability and it is associated with temperature in physics. Z is a normalization constant. NB to define Gibbs random fields, $P(\mathbf{f}) > 0$, $P(\mathbf{f}) \rightarrow 0$ $U(\mathbf{f}) \rightarrow \infty$: there are not configurations with 0 probability.



Gibbs priors and Regularization



$$\arg \min_f -\{\ln(p(g_n | f) p_f)\} = \arg \min_f -\{\ln(p(g_n | f)) + \ln(p_f)\}$$

Likelihood = adherence to the data

A-priori

Gaussian

$$K(\sigma) \sum_i \|g_{n,i} - Af_i\|^2$$

Poisson

$$\sum_i g_{n,i} \ln\left(\frac{g_{n,i}}{Af_i} + Af_i - g_{n,i}\right)$$

$$-\ln\left\{\frac{1}{Z} e^{\left\{-\frac{1}{\beta} U(\mathbf{f})\right\}}\right\}$$

$$J(f) = J_o(f) + \lambda J_R(f)$$

$$J_R(f) = U(f)$$



Role of λ



$$K(\sigma) \sum_i \|g_{n,i} - Af_i\|^2$$

$$-\ln \left\{ \frac{1}{Z} e^{\left\{ -\frac{1}{\beta} U(\mathbf{f}) \right\}} \right\}$$

$$J(f) = J_o(f) + \lambda J_R(f)$$

λ incorporates different elements here:

- the standard deviation of the noise in the likelihood
- the “temperature”, that is the decrease in the energy of the configurations with their cost
- the normalized constant Z.

λ has been investigated in the classical regularization theory (Engl et al., 1996), but not as deep in the Bayesian framework $\rightarrow \lambda$ is set experimentally through cross-validation.



Choice of the Gibbs priors



We choosed $\|\lambda Pf\|^2$ as a quadratic functional, but not specified P.

P is often chosen as a smoothing operator. The rationale is that the noise added to the image is often white (both Gaussian and Poisson) over the image as there is no correlation between adjacent pixels. Therefore its spatial content is uniform and with a larger bandwidth than the signal.

As a smoothing operator P is often a differential operator, which penalizes edges.

$$J_R(\mathbf{f}) = \sum_{c \in C} \phi_c(d^k c \mathbf{f})$$

k is the order of the derivative

ϕ_c can be L_2 norm (total variation), squared (Tikhonov)

k = 2 difference of gradients \rightarrow piecewise linear areas.

k = 3 difference of Hessian \rightarrow piecewise squared.

Neighbor of order higher than 2.



Quadratic Priors with k = 0



k = 0 – No derivative, the same gray level – single site cliques.

$$J_R(\mathbf{f}) = \sum_{c \in C} \phi(d^k c \mathbf{f}) = \sum_{c \in C} (d^0 c \mathbf{f})^2 = \sum_{p \in P} \mathbf{f}(p)^2$$

It has been applied to both Poisson and Gaussian noise models

Reduces bright spots and biases the solution to low intensity values.



Quadratic Priors with k = 1



k = 1 – First order derivatives – pair-sites cliques.

$$J_R(\mathbf{f}) = \sum_{c \in C} \phi(d^1 c \mathbf{f}) = \sum_{p \in P} \sum_{m \in \mathcal{N}_p} \phi(d^1 c \mathbf{f})^2 = \sum_{p \in P} \sum_{m \in \mathcal{N}_p} \phi\left(\frac{f(p) - f(m)}{d(p, m)}\right)$$

d(p,m) takes into account anisotropies in computing the distance.

If we consider $\phi(\cdot)$ a squared function, we have another form of Tikhonov regularization:

$$J_R(\mathbf{f}) = \sum_{p \in P} \sum_{m \in \mathcal{N}_p} \left(\frac{f(p) - f(m)}{d(p, m)} \right)^2 \quad ||P\mathbf{f}||^2$$



Quadratic Priors with k = 1



k = 1 – First order derivatives – pair-sites cliques.

$$J_R(\mathbf{f}) = \sum_{p \in P} \sum_{m \in \mathcal{N}_p} \left(\frac{f(p) - f(m)}{d(p, m)} \right)^2$$

If we consider $\phi(\cdot)$ a squared function, we have another form of Tikhonov regularization:

$$\| Pf \|_2^2$$

P is the convolution with the Laplacian operator:

$$\begin{bmatrix} 0 & -1 & 0 \\ -1 & 4 & -1 \\ 0 & -1 & 0 \end{bmatrix}$$

First order neighboring System
4-neighboring System

$$\begin{bmatrix} -\frac{\sqrt{2}}{2} & -1 & -\frac{\sqrt{2}}{2} \\ -1 & 4+2\sqrt{2} & -1 \\ -\frac{\sqrt{2}}{2} & -1 & -\frac{\sqrt{2}}{2} \end{bmatrix}$$

Second order neighboring System
8-neighboring System



Non-quadratic potential functions, k = 1



Quadratic functions priors imposes smoothness everywhere. Large true gradients of the solution are therefore penalized → smoothing sharp edges.

In imaging objects tend to be piecewise smooth, but different pieces of objects are separated by more or less sharp edges. We want to smooth inside the object but not the edge. A parallel worthwhile to be investigated is with anisotropic diffusion (Koenderink, 1987; Perona&Malik, 1990).

We search different potential functions (Geman&McClure, 85; Charbonnier et al., 1994, 1997; Hebert&Lehay, 1989).



Non-quadratic potentials (Charbonier et al., 1997)



1. $\phi(t) \geq 0 \quad \forall t \quad \phi(0) = 0$ Derives from the definition of potential
2. $\Phi(t) \geq 0 \quad \forall t$
3. $\phi(t) = \phi(-t)$ Positive and negative gradients are equally considered
4. $\phi(t) \in C^1$ This is to avoid instability.

Up to now quadratic potentials are OK

5. $\frac{\varphi'(t)}{2t}$ The potential increase rate should decrease with t.
6. $\lim_{t \rightarrow \infty} \frac{\varphi'(t)}{2t} = 0$ The potential increase rate should decrease for all t (at least for large values of t)
7. $\lim_{t \rightarrow 0} \frac{\varphi'(t)}{2t} = \cos t > 0$ The potential increases at least linearly for t = 0.



Few non-quadratic potentials (Vicedomini 2008)

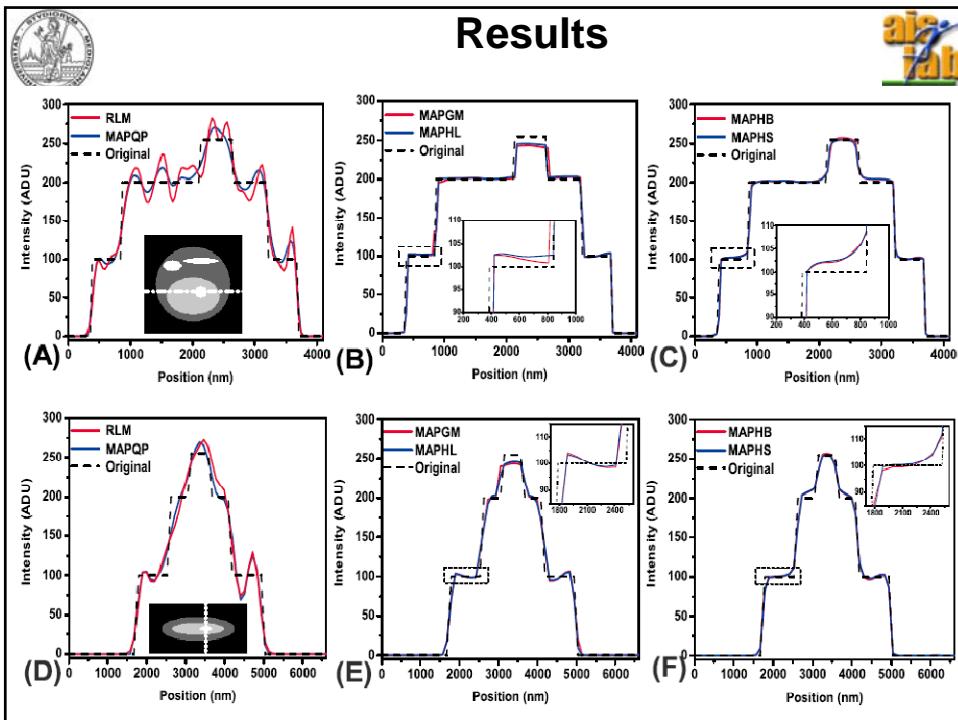


Regularization name	Potential function	Expression of $\varphi(t)$	Expression of $\psi(t) = \varphi'(t)/2t$	Convex
Quadratic-Potential	φ_{QP}	t^2	1	yes
Geman-McClure	φ_{GM}	$\frac{t^2}{1+t^2}$	$\frac{1}{(1+t^2)^2}$	no
Hebert-Leahy	φ_{HL}	$\log(1+t^2)$	$\frac{1}{1+t^2}$	no
Huber	φ_{HB}	$\begin{cases} t^2, & t \leq 1 \\ 2 t -1, & t > 1 \end{cases}$	$\begin{cases} 1, & t \leq 1 \\ 1/ t , & t > 1 \end{cases}$	yes
Hyper-Surface	φ_{HS}	$2\sqrt{1+t^2} - 2$	$\frac{1}{\sqrt{1+t^2}}$	yes

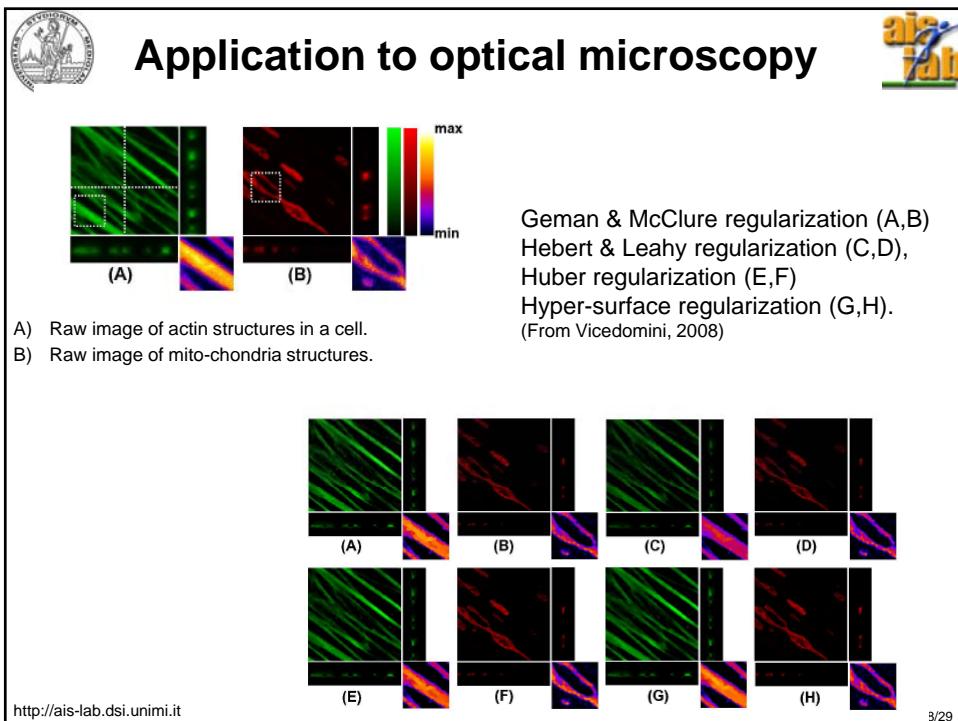
Asymptotic linear behavior

Asymptotic log-like behavior

Results



Application to optical microscopy





Summary



MAP estimate can be seen as a statistical version of regularization.

The regularization term can be derived from the potential energy associated to an adequate neighbor system defined over the object (e.g. over the image).

Under this hypothesis the value assumed by the elements of the object to be reconstructed (e.g. restored or filtered image) represent a MRF.

Different neighbor systems and different potential functions allow defining different properties of the object.

For quadratic potential functions, Tikhonov regularizer are derived.

The discrepancy term for the data represents the likelihood and can accommodate different statistical models: Poisson, Gaussian or even mixture models.



Issues to be investigated



- Is the a-priori term really suitable? There is a difference between the metric used in evaluating the distance between the image and the projection and the gradient. Which is the best metric?
- How to set the regularization parameter? We have introduced a generalized discrepancy principle, but can we do something better?
- Optimization and parallelization of the code using CUDA. Real-time reconstruction and filtering.



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