

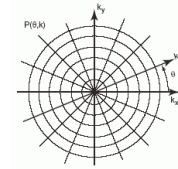
Introduction to Medical Imaging

Iterative Reconstruction Methods

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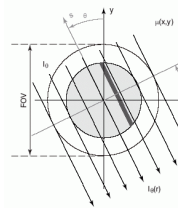
Ideal Assumptions



Dense and regular sampling of the Fourier domain → many projections



Noise free projections



Straight rays

Non-Ideal Scenarios

Projections might be:

- sparse
- acquired over less than 180°
- noisy



20 projections

SNR=10

low-dose CT

high-dose CT

Rays might be non-linear (curved, refracted, scattered,...)

- for example: refraction in ultrasound imaging

Dealing With Non-Ideal Scenarios

Iterative methods are advantageous in these cases

They can handle:

- limited number of projections
- irregularly-spaced and -angled projections
- non-straight ray paths (example: refraction in ultrasound imaging)
- corrective measures during reconstruction (example: metal artifacts)
- presence of statistical (Poisson) noise and scatter (mainly in functional imaging: SPECT, PET)

Specifics

In medical imaging:

- M unknown voxels (depending on desired object resolution)
- N known measurements (pixels in the projection images)
- represent voxels and pixels as vectors V and P , respectively

$$w_{11}v_1 + w_{12}v_2 + \dots + w_{1M}v_M = p_1$$

$$w_{21}v_1 + w_{22}v_2 + \dots + w_{2M}v_M = p_2$$

....

$$w_{N1}v_1 + w_{N2}v_2 + \dots + w_{NM}v_M = p_N$$

- this gives rise to a system $W \cdot V = P$

Solving for V

The obvious solution is then:

- compute $V = W^{-1} \cdot P$

The main problem with this direct approach:

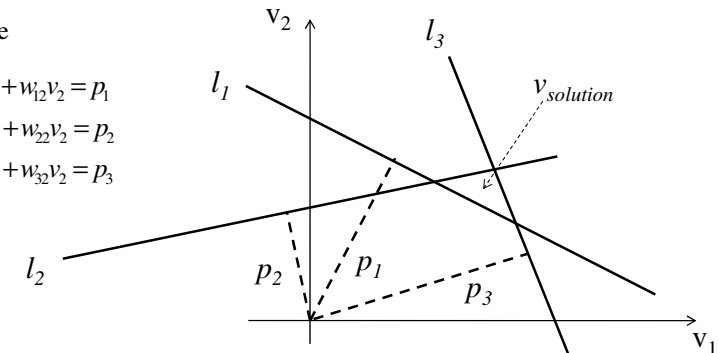
- P is not be consistent due to noise \rightarrow lines do not intersect in solution
- This turns $W \cdot V = P$ into an optimization problem

2D case

$$w_{11}v_1 + w_{12}v_2 = p_1$$

$$w_{21}v_1 + w_{22}v_2 = p_2$$

$$w_{31}v_1 + w_{32}v_2 = p_3$$



Optimization Algorithms

Algebraic methods

- Algebraic Reconstruction Technique (ART), SART, SIRT
- Projection Onto Convex Sets (POCS)

Sparse system solvers

- Gradient Descent (GD), Conjugate Gradients (CG)
- Gauss-Seidel

Statistical methods

- Expectation Maximization (EM)
- Maximum Likelihood Estimation (MLE)

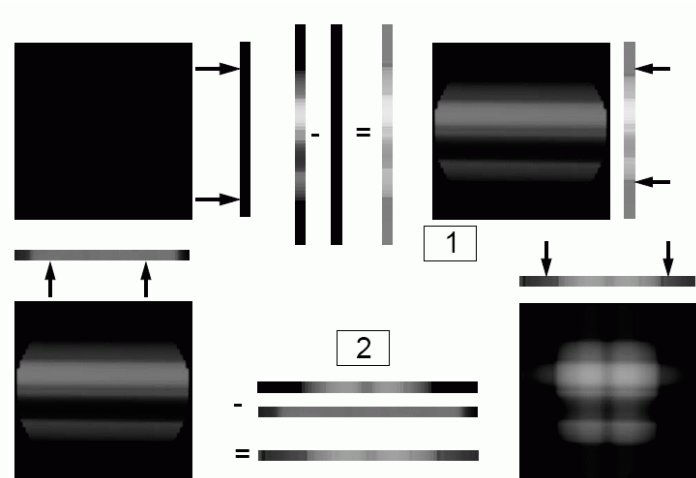
All of these are *iterative* methods:

- predict \rightarrow compare \rightarrow correct \rightarrow predict \rightarrow compare \rightarrow correct ...

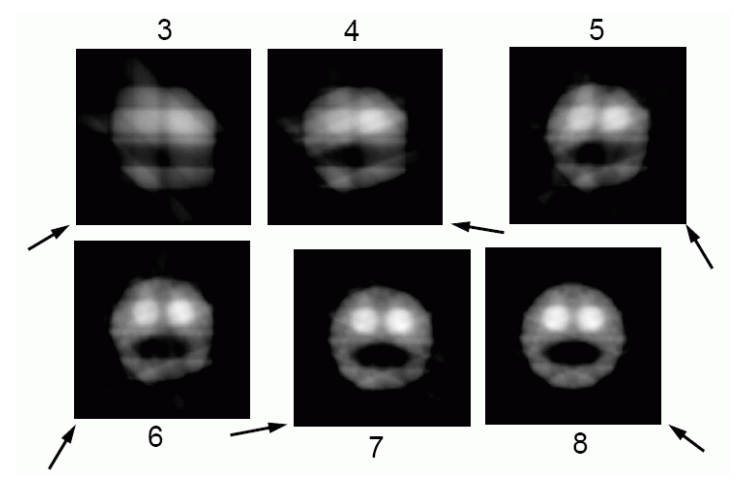
Big Picture: Iterative Reconstruction

Before delving into details,
let's see an iterative scheme at work

Iterative Reconstruction Demonstration: SART

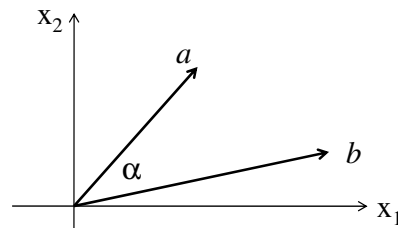


Iterative Reconstruction Demonstration: SART



Foundations: Vectors

Consider two vectors, a and b



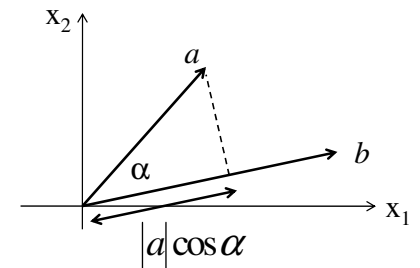
$$a = \vec{a} = [a_1 \ a_2], \quad |a| = \sqrt{a_1^2 + a_2^2}$$

$$b = \vec{b} = [b_1 \ b_2], \quad |b| = \sqrt{b_1^2 + b_2^2}$$

Foundations: Scalar Projection

Scalar projection of a onto b :

$$|a| \cos \alpha = a \cdot \frac{b}{|b|}$$



The dot product:

$$\begin{aligned} a \cdot b &= \vec{a} \cdot \vec{b}^T = [a_1 \ a_2] \cdot [b_1 \ b_2]^T = a_1 b_1 + a_2 b_2 \\ &= |a| \cdot |b| \cos \alpha \end{aligned}$$

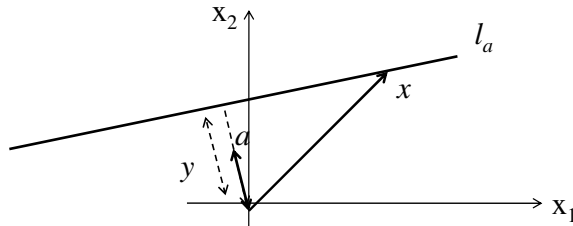
→ the scalar projection is the dot product with $|b| = 1$ (unit vector)

$$|b| = \sqrt{b_1^2 + b_2^2} = 1$$

Foundations: Line Equation

$$a_1x_1 + a_2x_2 = y$$

$$|a| = \sqrt{a_1^2 + a_2^2} = 1$$



The vector a is the unit vector normal to the line l_a

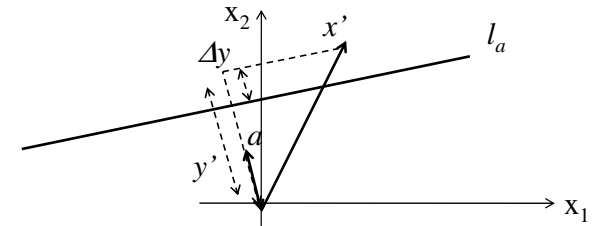
The length y is the perpendicular distance of l_a to the origin

For any point x :

- if x is on l_a then the scalar projection of x onto a will be:

$$x \cdot a = y$$

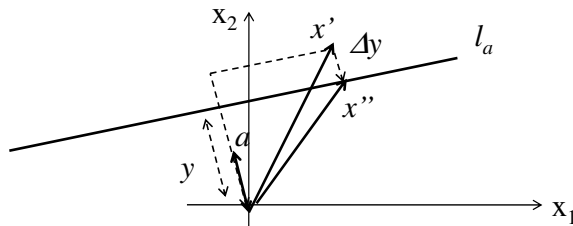
Foundations: Distance From Line



For any other point x' not on l_a the scalar projection of x' onto a will be:

$$x' \cdot a = y' = y + \Delta y$$

Foundations: Closest Point



The closest point to x' on l_a is x'' , computed by:

$$\begin{aligned} x'' &= x' - \Delta y \cdot a \\ &= x' - (x' \cdot a - y) \cdot a \\ &= x' + (y - x' \cdot a) \cdot a \end{aligned}$$

Foundations: Solving an Equation System

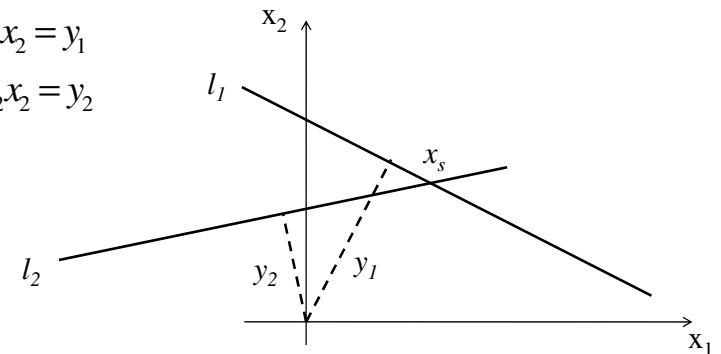
Assume you have two equations to solve for solution point

$$x_s = (x_1, x_2)$$

- the intersection of the two lines

$$a_{11}x_1 + a_{12}x_2 = y_1$$

$$a_{21}x_1 + a_{22}x_2 = y_2$$

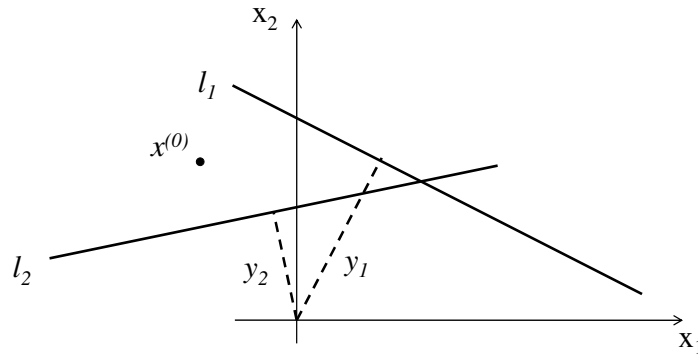


Foundations: Iterating to Solution

Of course, you could solve this equation via Gaussian elimination

- we shall take an iterative approach instead

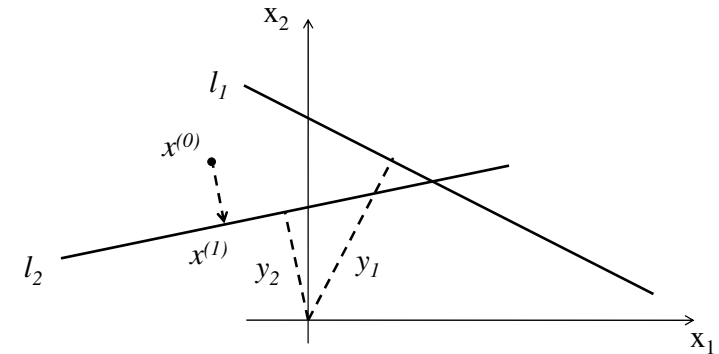
Start with some point $x^{(0)} = (x_1, x_2)$



Foundations: Iterating to Solution

Pick an equation (line, say l_2) and find the closest point to $x^{(0)}$

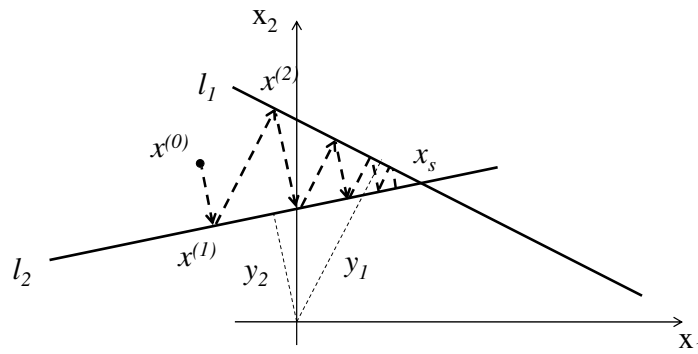
- use the approach outlined before
- this gives a new point $x^{(1)}$



Foundations: Iterating to Solution

Iteratively

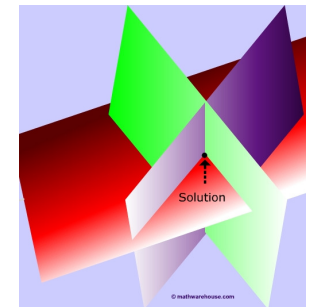
- pick alternate equations (lines) and project
- the solution will *converge* towards x_s
- the more iterations the closer the convergence



Foundations: Extension to Higher Dimensions

Three dimensions:

- 3 equations with 3 unknowns



N dimensions:

- N equations with M unknowns
- M can be less or greater than N
- inconsistent (most often) or not

Specifics to Medical Imaging

In medical imaging:

- M unknown voxels (depending on desired object resolution)
- N known measurements (pixels in the projection images)
- represent voxels and pixels as vectors V and P , respectively

$$w_{11}v_1 + w_{12}v_2 + \dots + w_{1M}v_M = p_1$$

$$w_{21}v_1 + w_{22}v_2 + \dots + w_{2M}v_M = p_2$$

....

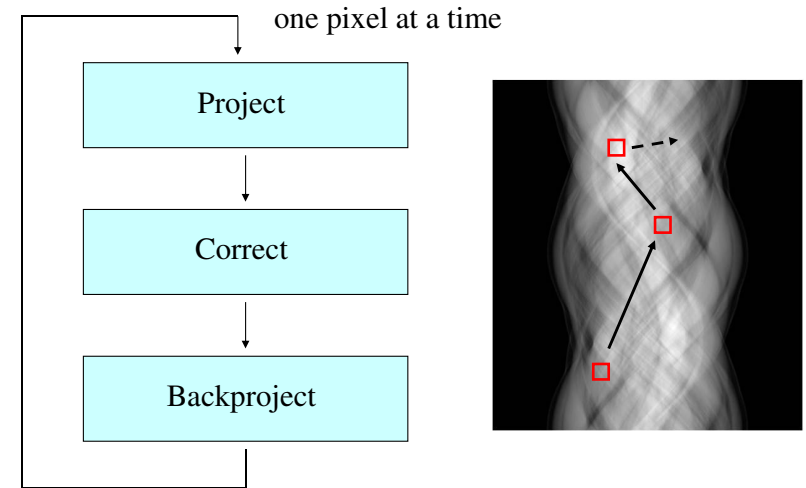
$$w_{M1}v_1 + w_{M2}v_2 + \dots + w_{NM}v_M = p_N$$

- this gives rise to a system $W \cdot V = P$

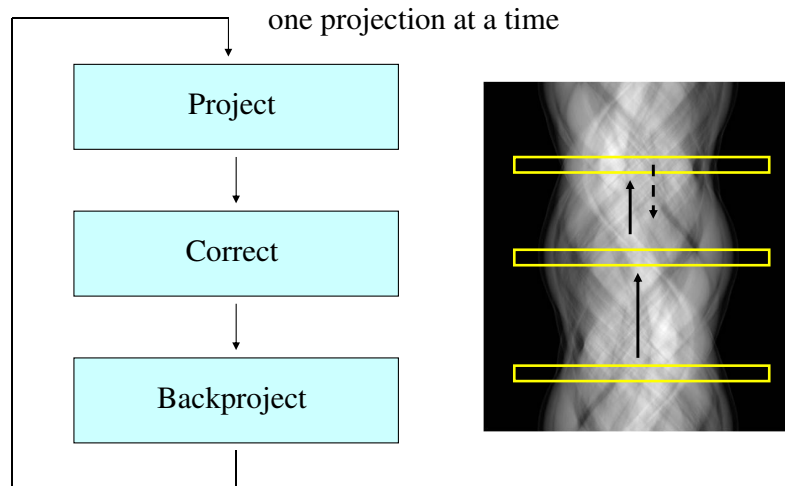
Iterate either by

- ray by ray (Algebraic Reconstruction Technique, ART)
- image by image (Simultaneous ART, SART)
- all data at once (SIRT)

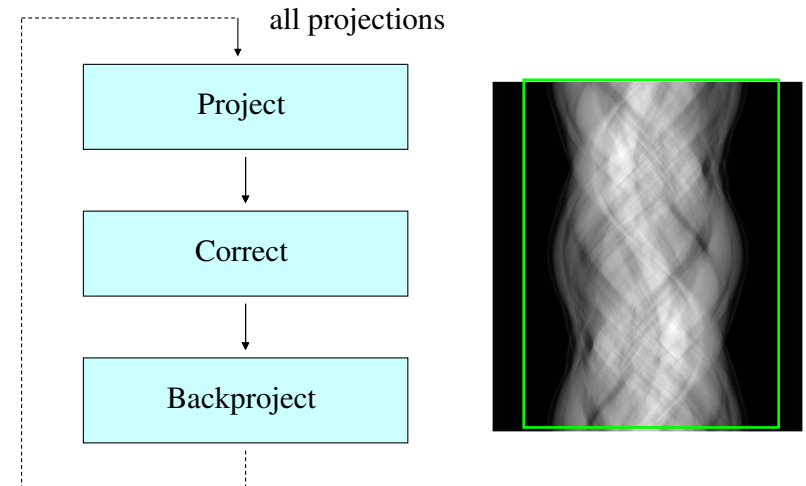
Iterative Update Schedule: ART



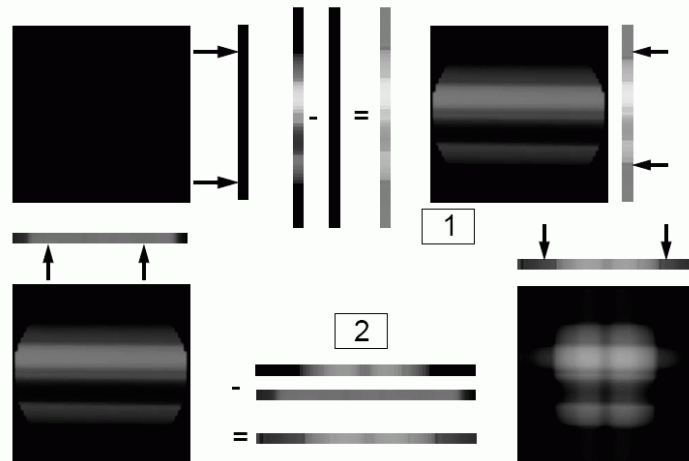
Iterative Update Schedule: SART



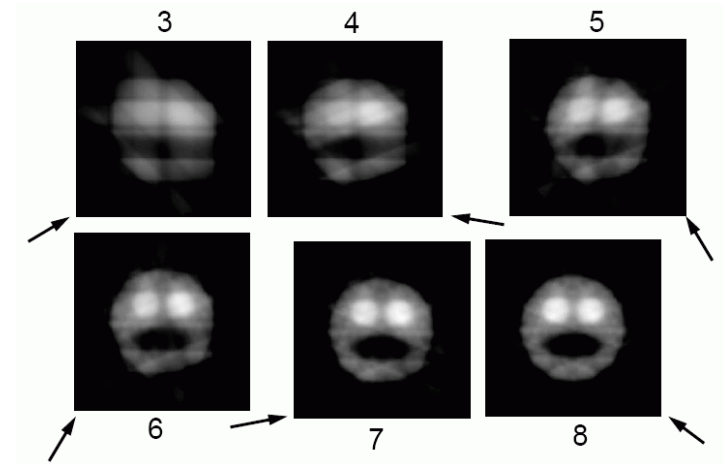
Iterative Update Schedule: SIRT



Iterative Reconstruction Demonstration: SART



Iterative Reconstruction Demonstration: SART



SART

Iteratively solves $W \cdot V = P$

$$v_j^{k+1} = v_j^k + \lambda \frac{p_i - \sum_j v_j^k w_{ij}}{\sum_i \frac{p_i - \sum_j v_j^k w_{ij}}{\sum_j w_{ij}} w_{ij}}$$

SART

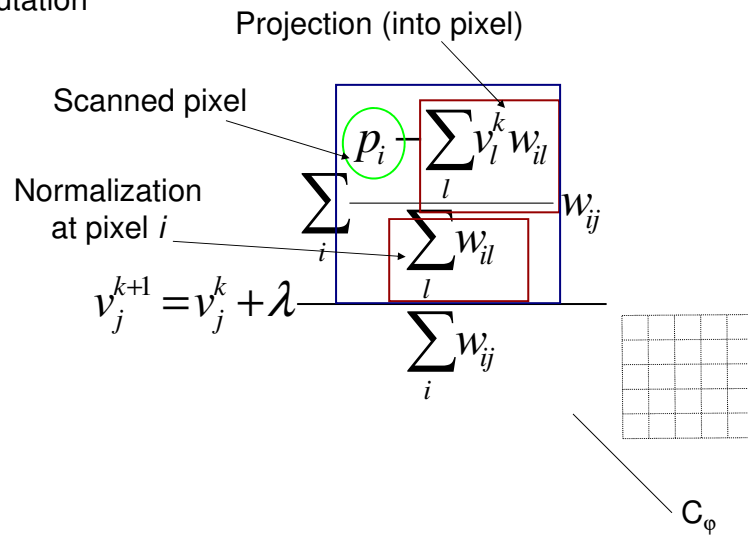
Projection

Projection (into pixel)

$$v_j^{k+1} = v_j^k + \lambda \frac{p_i - \sum_l v_l^k w_{il}}{\sum_l w_{il}} w_{ij}$$

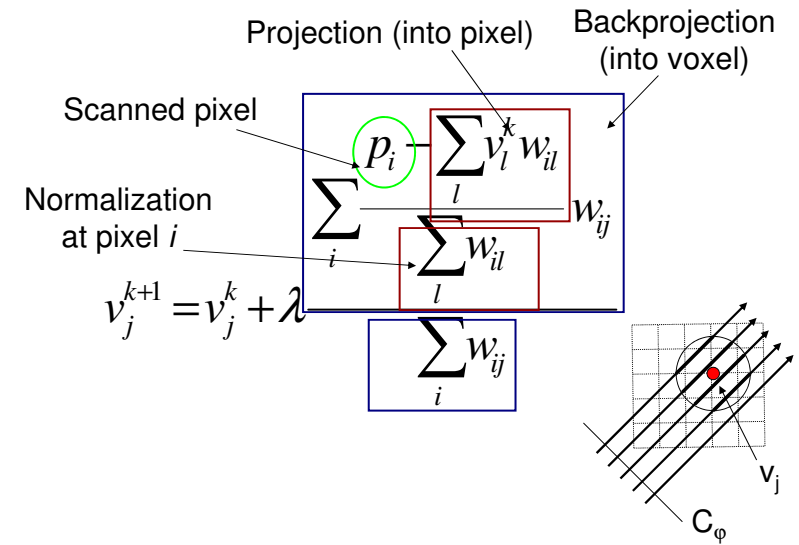
SART

Correction factor
computation



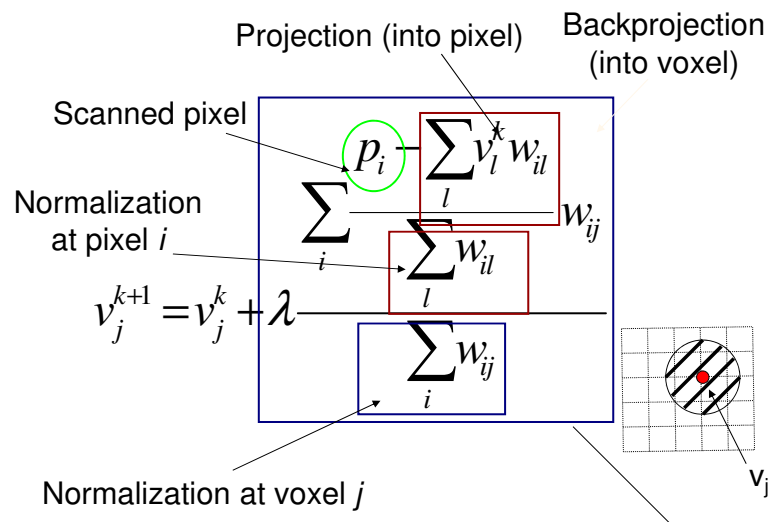
SART

Backprojection



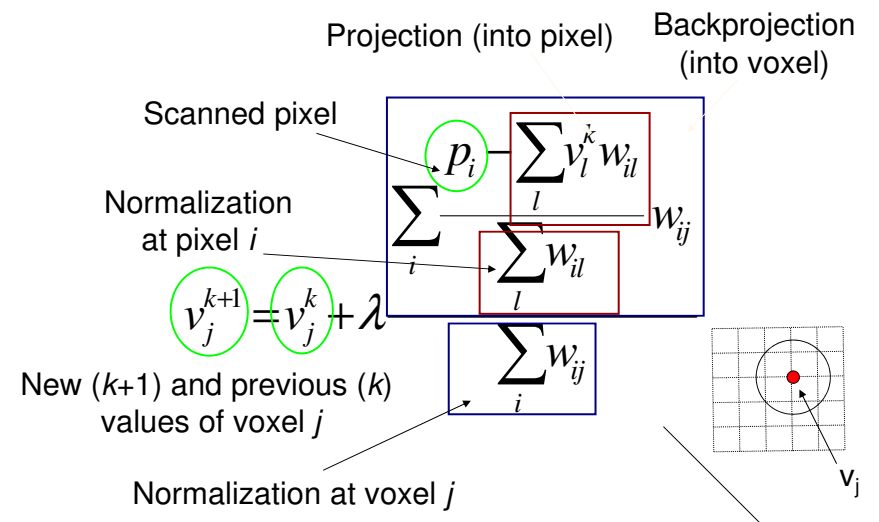
SART

Voxel normalization



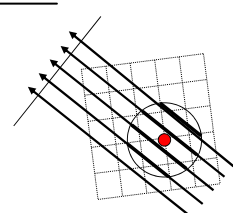
SART

Voxel update



SART

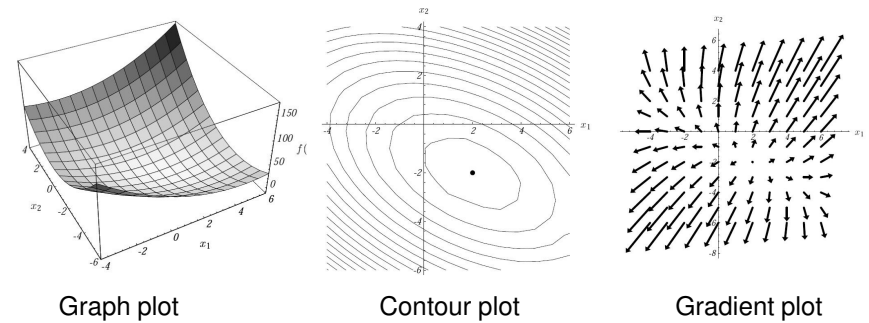
Next projection

$$v_j^{k+1} = v_j^k + \lambda \frac{\sum_i \frac{p_i - \sum_l v_l^k w_{il}}{\sum_l w_{il}} w_{ij}}{\sum_i w_{ij}}$$


Gradient Descent

Quadratic form of a vector: $f(x) = \frac{1}{2} x^T A x - b^T x + c$

- this equation is minimized when $A \cdot x = b$
- this occurs when $f'(x) = 0$
- thus, minimizing the quadratic form will solve the reconstruction problem



Steepest Descent

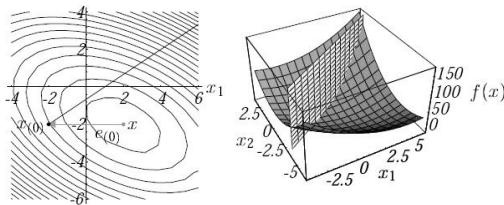
Start at an arbitrary point and slide down to the bottom of the parabola

- in practice this will be a hyper-parabola since x, b are high-dimensional
- choose the direction in which f decreases most quickly

$$-f'(x_{(i)}) = b - Ax_{(i)}$$

where $x_{(i)}$ is the current (predicted) solution

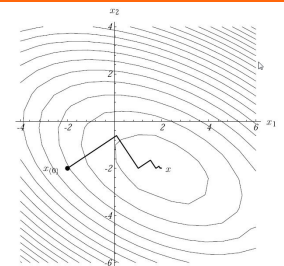
- similar to ART but now looks at all equations simultaneously



Steepest Descent

Start at some initial guess $x_{(0)}$

- this will likely not find the solution
- need to follow $f'(x_{(0)})$ some ways and then change directions
- question is *where* do we change directions



Some basics:

- error: how far are we away from the solution

$$e_{(i)} = x_{(i)} - x$$

- residual: how far are we away from the correct value of b

$$r_{(i)} = b - Ax_{(i)}$$

$$r_{(i)} = Ae_{(i)} \quad \text{A transforms } e \text{ into the space of } b$$

$$r_{(i)} = -f'(x_{(i)})$$

Steepest Descent

Finding the right place to turn directions is called *line search*

$$x_{(1)} = x_{(0)} + \alpha r_{(0)}$$

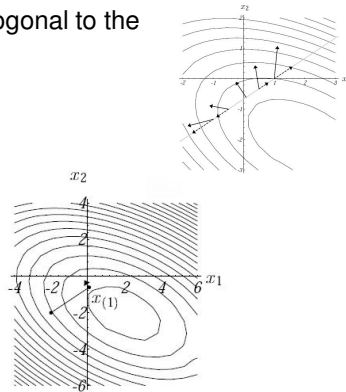
To find α we can use the following requirements:

- the new direction of r must be orthogonal to the previous:

$$r_{(1)}^T r_{(0)} = 0$$

- the residual at $x_{(1)}$ $f'(x_{(1)}) = -r_{(1)}$

- after some math: $\alpha = \frac{r_{(0)}^T r_{(0)}}{r_{(0)}^T A r_{(0)}}$

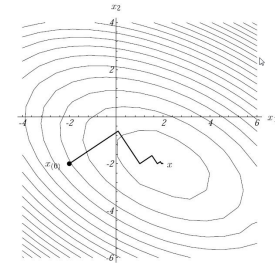


Steepest Descent: Summary

$$r_{(i)} = b - A x_{(i)}$$

$$\alpha = \frac{r_{(i)}^T r_{(i)}}{r_{(i)}^T A r_{(i)}}$$

$$x_{(i+1)} = x_{(i)} + \alpha r_{(i)}$$



Shortcoming:

- sub-optimal since some directions might be taken more than once
- this can be fixed by the method of Conjugate Gradients

Conjugate Gradients

Picks a set of *orthogonal* search directions $d_{(0)}, d_{(1)}, d_{(2)}, \dots$

- take exactly one step along each
- stop at exactly the right length for each to line up evenly with x

$$x_{(i+1)} = x_{(i)} + \alpha_{(i)} d_{(i)}$$

- to determine $\alpha_{(i)}$ use the fact that $e_{(i+1)}$ should be orthogonal to $d_{(i)}$

$$d_{(i)}^T e_{(i+1)} = 0$$

$$d_{(i)}^T (e_{(i)} + \alpha d_{(i)}) = 0$$

$$\alpha_{(i)} = \frac{d_{(i)}^T e_{(i)}}{d_{(i)}^T d_{(i)}}$$

- however, this requires knowledge of $e_{(i)}$ which we do not have

Conjugate Gradients

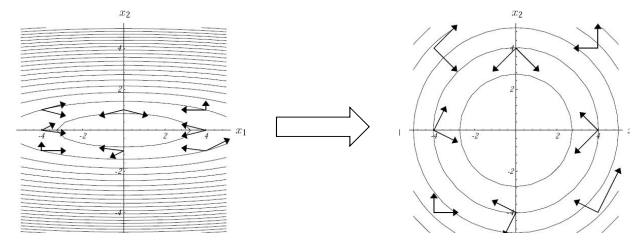
Solution:

- make the search direction A -orthogonal (or, *conjugate*)

$$\alpha_{(i)} = \frac{d_{(i)}^T A e_{(i)}}{d_{(i)}^T A d_{(i)}} = \frac{d_{(i)}^T r_{(i)}}{d_{(i)}^T A d_{(i)}}$$

- A transforms a coordinate system such that two vectors are orthogonal

$$d_{(i)}^T A d_{(j)} = 0 \quad i \neq j$$



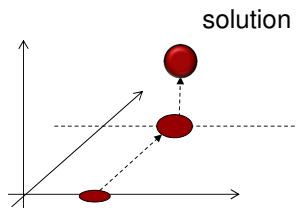
Conjugant Gradients

All directions taken are mutually orthogonal

- each new residual is orthogonal to all the previous residuals and search directions
- each new search direction is constructed (from the residual) to be A-orthogonal to all the previous residuals and search directions

Each new search direction adds a new dimension to the traversed sub-space

- the solution is a projection into the sub-space explored so far
- so after n steps the full space is built and the solution has been reached



Conjugant Gradients: Summary

$$d_{(0)} = r_{(0)} = b - Ax_{(0)},$$

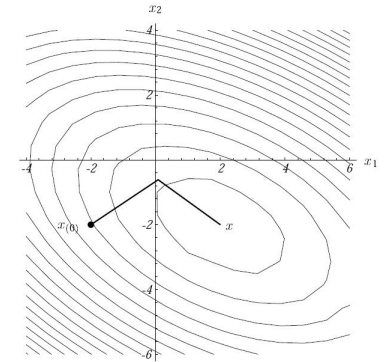
$$\alpha_{(i)} = \frac{r_{(i)}^T r_{(i)}}{d_{(i)}^T A d_{(i)}}$$

$$x_{(i+1)} = x_{(i)} + \alpha_{(i)} d_{(i)},$$

$$r_{(i+1)} = r_{(i)} - \alpha_{(i)} Ad_{(i)},$$

$$\beta_{(i+1)} = \frac{r_{(i+1)}^T r_{(i+1)}}{r_{(i)}^T r_{(i)}},$$

$$d_{(i+1)} = r_{(i+1)} + \beta_{(i+1)} d_{(i)}.$$



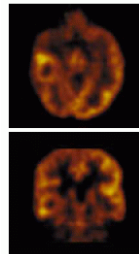
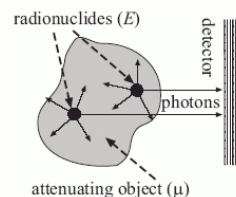
Statistical Techniques

Algebraic/gradient methods do not model statistical effects in the underlying data

- this is OK for CT (within reason)

However, the emission of radiation from radionuclides is highly statistical

- the direction is chosen at random
- similar metabolic activities may not emit the same radiation
- not all radiation is actually collected (collimators reject many photons)
- in low-dose CT, noise is also a significant problem



Need a reconstruction method that can accounts for these statistical effects

- Maximum Likelihood – Expectation Maximization (ML-EM) is one such method

Foundations: The Poisson Distribution

Also called the *law of rare events*

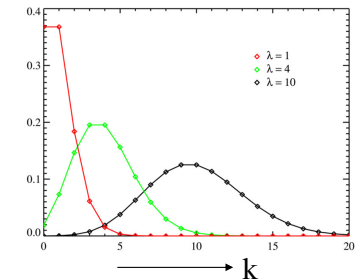
- it is the binomial distribution of k as the number of trials n goes to infinity

$$\lim_{n \rightarrow \infty} \Pr(X = k) = \lim_{n \rightarrow \infty} \binom{n}{k} p^k (1-p)^{n-k}$$

- with $p = \lambda / n$

$$f(k; \lambda) = \frac{e^{-\lambda} \lambda^k}{k!}$$

λ : expected number of events (the mean)
in a given time interval



Some examples for Poisson-distributed events:

- the number of phone calls at a call center per minute
- the number of spelling errors a secretary makes while typing a single page
- the number of soldiers killed by horse-kicks each year in each corps in the Prussian cavalry
- the number of positron emissions in a radio nucleotide in PET and SPECT
- the number of annihilation events in PET and SPECT

Overall Concept of ML-EM

There are three types of variables

#1: The observed data $y(d)$:

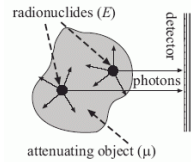
- the detector readings

#2: The unobserved (latent) data $x(b)$:

- the photon emission activities in the pixels (the tissue), $x(b)$
- these give rise to the detector readings
- they follow a Poisson distribution

#3: The model parameters $\lambda(b)$:

- these cause the emissions
- they are the metabolic activities (state) of interest
- the emissions only approximate those
- they represent the expectations (means, λ) of the resulting Poisson distribution causing the readings at the detectors



Overall Concept of ML-EM

There is a many-to-one mapping of parameters → data

Since there is a many-to-one mapping, many objects are probable to have produced the observed data

- the object reconstruction (the *image*) having the highest such probability is the *maximum likelihood estimate* of the original object

Goal:

- estimate the model parameters using the observed data

Solution:

- EM will converge to a solution of maximum likelihood (but not necessarily the global maximum)

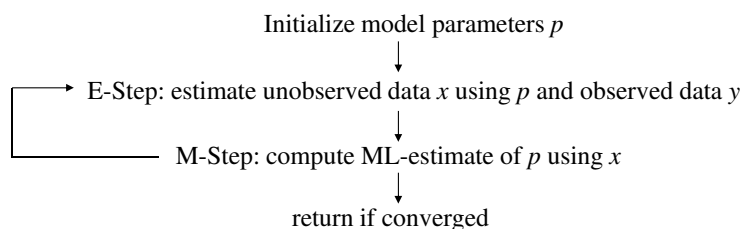
Overall Concept of ML-EM

Initialization step: choose an initial setting of the model parameters

Then proceed to EM, which has two steps, executed iteratively:

- E (expectation) step: estimate the unobserved data from the current estimate of the model parameters and the observed data
- M (maximization) step: compute the maximum-likelihood estimate of the model parameters using the estimated unobserved data

Stop when converged



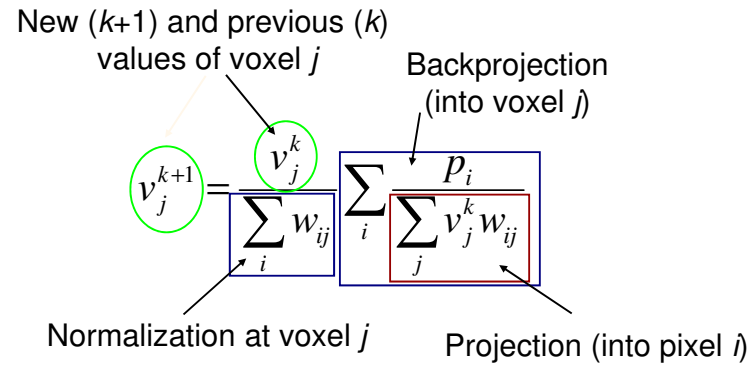
Maximum Likelihood Expectation Maximization (ML-EM)

After combining the E-step and the ML-step:

$$v_j^{k+1} = \frac{v_j^k}{\sum_i w_{ij}} \sum_i \frac{p_i}{\sum_j v_j^k w_{ij}}$$

Maximum Likelihood Expectation Maximization (ML-EM)

Maximizes the likelihood of the values of (object) voxels j , given values at (detector) pixels i



Algorithm Comparison

SART:

- projection ordering important
- ensure that consecutively selected projections are approximately orthogonal
- random selection works well in practice

CG:

- much depends on the condition number of the (system) matrix A
- various pre-conditioning methods exist in the literature
- also, line search can be expensive and inaccurate
- various methods and heuristics for line search have been described in the literature

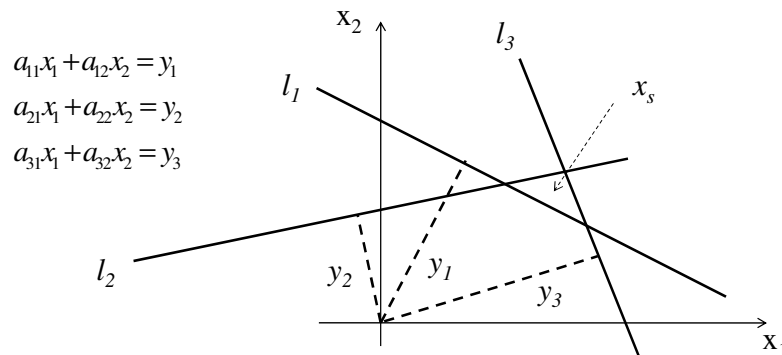
EM:

- convergence slow if all projections are applied before voxel update
- use OS-EM (Ordered Subsets EM): only a subset of projections are applied per iteration

Inconsistent Equations

Real life data (as mentioned earlier)

- typically equations (the data) are not consistent
- you may have more equations (data) than unknowns or not enough
- solution falls within a *convex* shape spanned by the intersection set
- need further criteria to determine the true solution (some *prior model*)



Determining the True Solution

Need further criteria to determine the true solution

Use some *prior model*

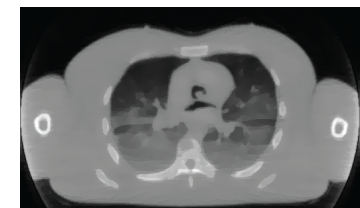
- smoothness, approximate shape, sharp edges, ...
- incorporate this model into the reconstruction procedure

Example:

- enforce smoothness by intermittent blurring
- but at the same time preserve edges



streak artifacts, good edges



smooth, good edges