



Algorithms for Tomographic Reconstruction of Non-stationery Targets: A Scientific Computing Project

Debasis Mitra

Department of Computer Sciences

Florida Institute of Technology

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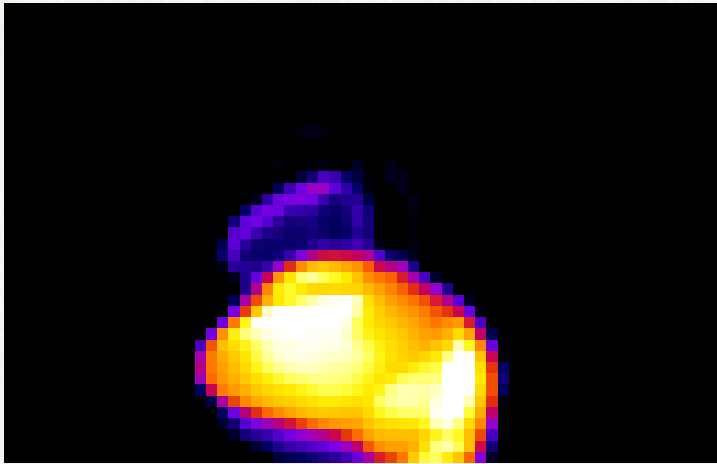
Debasis Mitra

ABSTRACT: Imagine taking pictures of an object V from different angles and then reconstructing the 3D image of the surface of V computationally. Now, further imagine that the source of the radiation is not the reflected light from the surface of O , but actually the source is inside V . This latter technique is used in medical imaging over the decades for non-invasively probing diseases, and is called tomography.

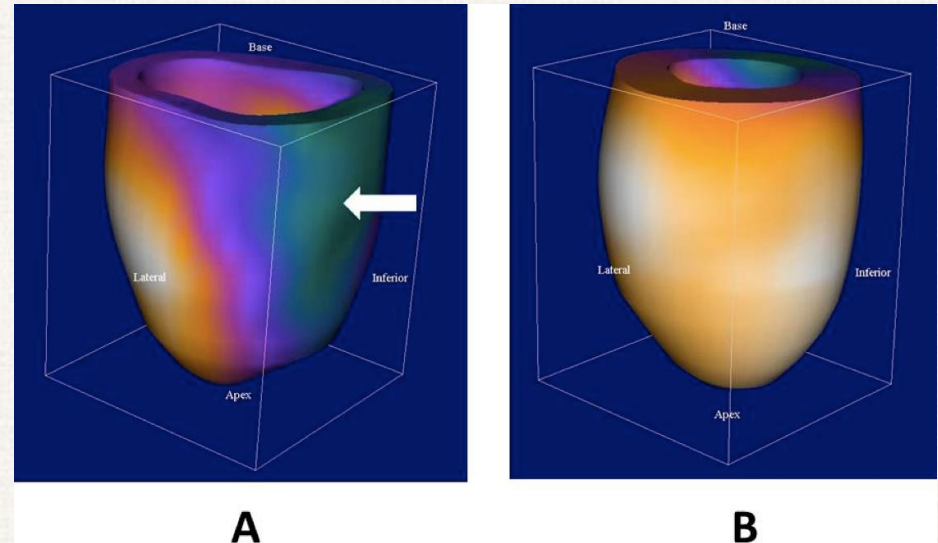
Tomographic imaging is normally done under stationery conditions. In our current project we address dynamic tomography problems, where the sources of radiation inside a live object are the metabolites. In this talk I will introduce the basics of the tomography problem from an algorithmic point of view, and some of our results.

Tomography: Non-invasive Probing of Human Body

Views from a rotating camera: Sinogram

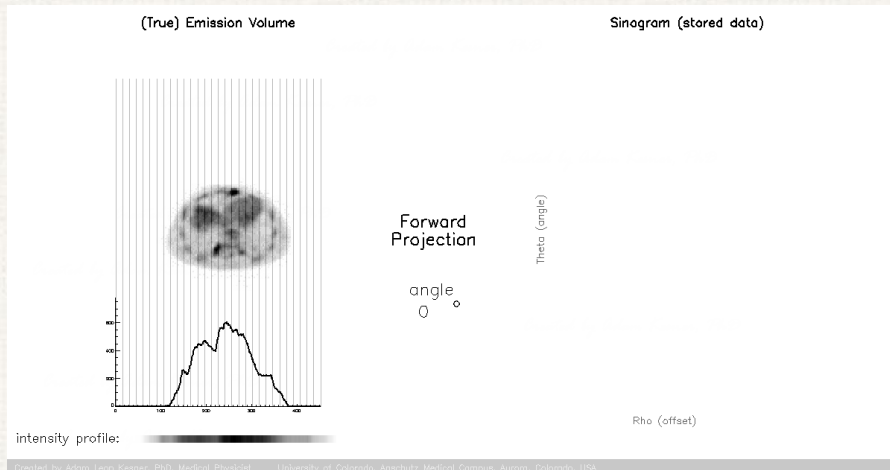


Computed 3D Reconstructed Image



Cardiac reversible ischemia: stressed(A), rest(B)
<http://www.aipes-eeig.org/white-paper-spect-spect-ct.html>

Tomography: Non-invasive Probing of Human Body



Forward Problem:

This is what the imaging system does

$$P = S.V$$

P: Camera Views - input

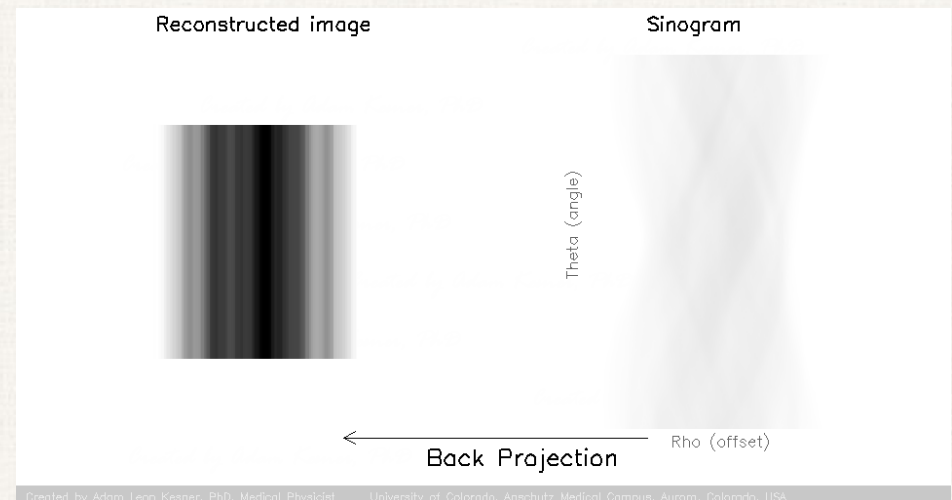
S: Camera model / System Matrix - computed

V: Target object - unknown

Inverse Problem:

This is what a reconstruction algorithm does

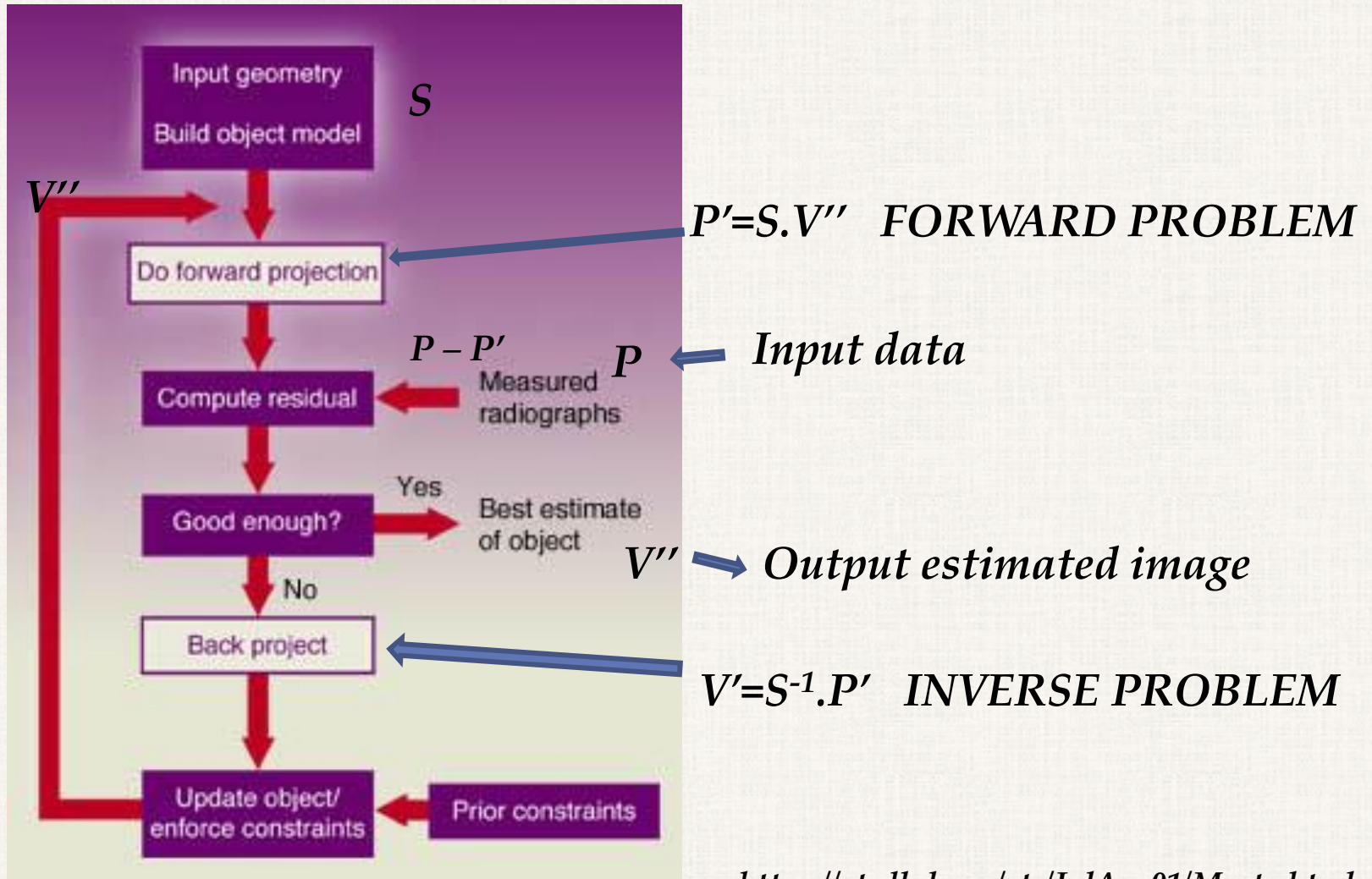
$$V = S^{-1}.P$$



Typical dimensions of the Problem

- $V = 64 \times 64 \times 64$ voxels $\Rightarrow 262,000 \Rightarrow \times 4$ bytes $\Rightarrow 8$ Mb
- $P = 64 \times 64$ pixels per view $\times 120$ views $\Rightarrow 500,000$
 $\Rightarrow \times 4$ b $\Rightarrow 20$ Mb
- $S = 8 \times 20 \Rightarrow 160$ Mb
- **Moreover,**
 - P is Very noisy
 - S is not perfect

Iterative Reconstruction



<https://str.llnl.gov/str/JulAug01/Martz.html>

Types of Medical Tomography Systems

Computed Tomography (CT): X-Ray Absorption

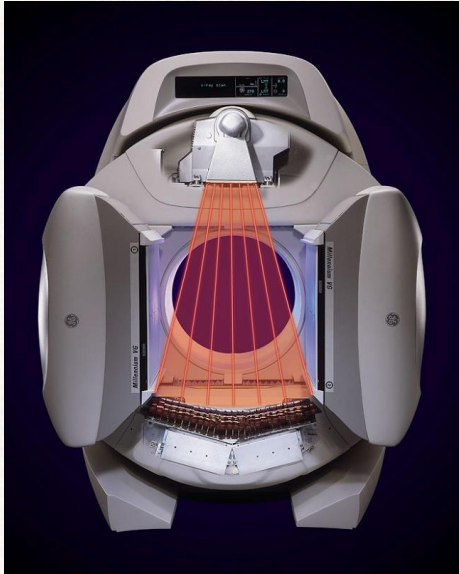
Positron Emission Tomography (PET): positron->gamma ray emission

Single Photon Emission Computed Tomography (SPECT): gamma

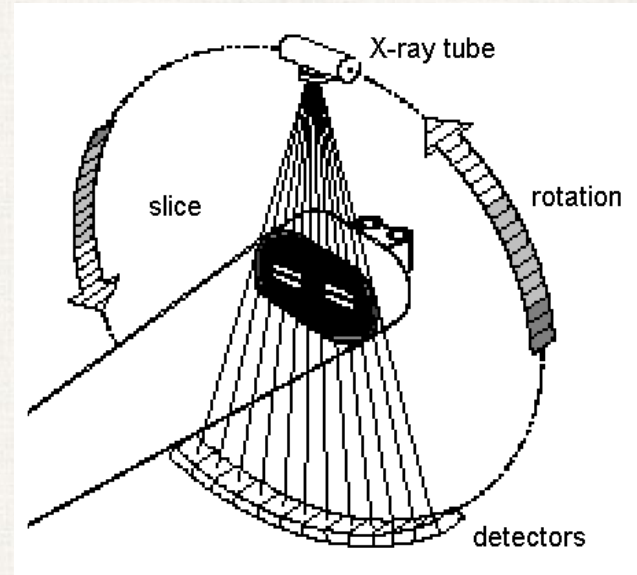
Magnetic Resonance Imaging (MRI)

• • • • •

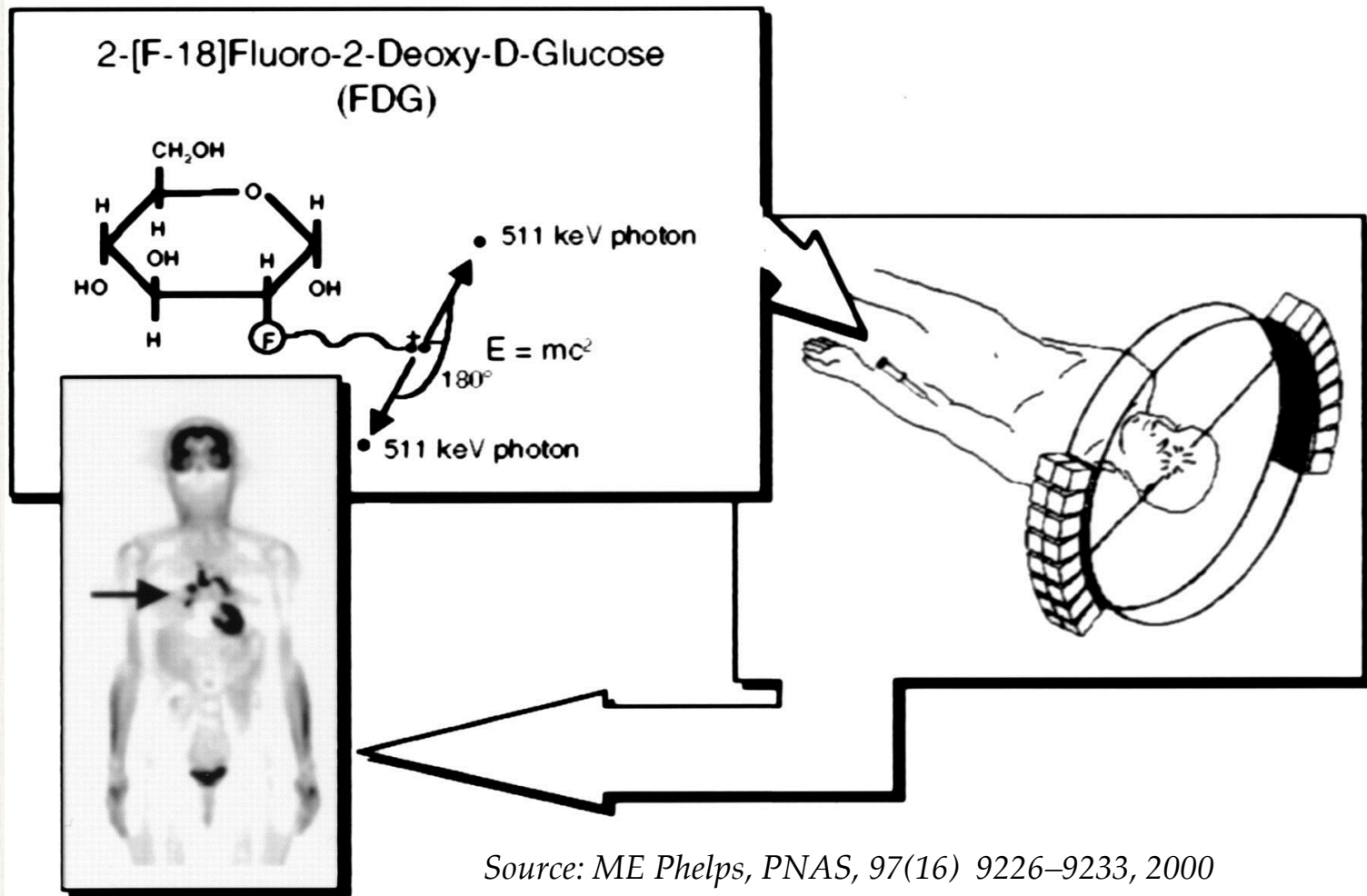
Computed Tomography: Absorption of X-ray Anatomical Imaging



20-40 keV



Emission Tomography: Functional Imaging

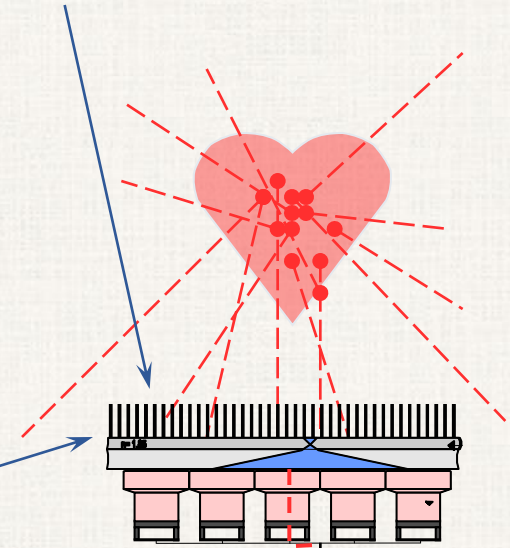


SPECT: Gamma Emission Tomography

(Single Photon Emission Computed Tomography)



collimators



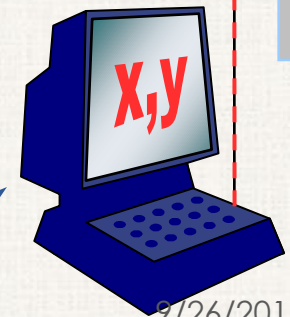
γ -ray detectors

Resolution

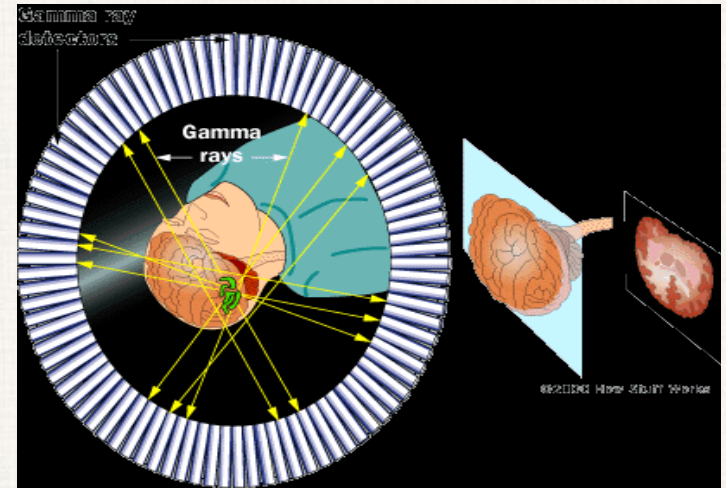
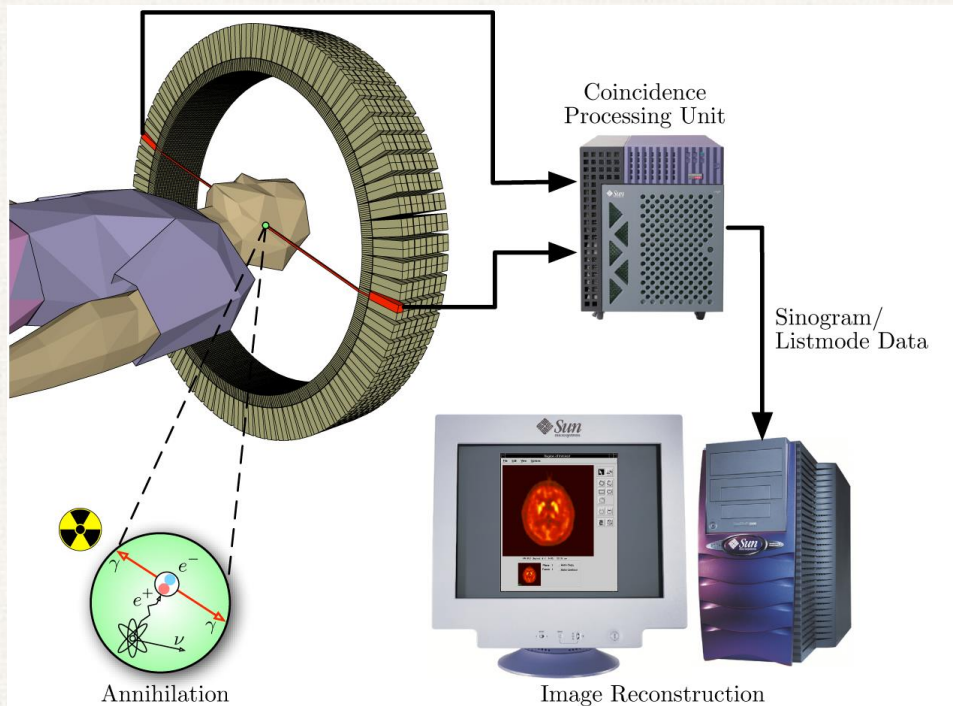
100-200 keV

Sensitivity

Acquisition system



PET: Positron Emission Tomography



- Positron annihilates with electron
⇒ two gamma photons each at 511 keV leave at 180°
- Coincidence detection (“electronic collimation”)

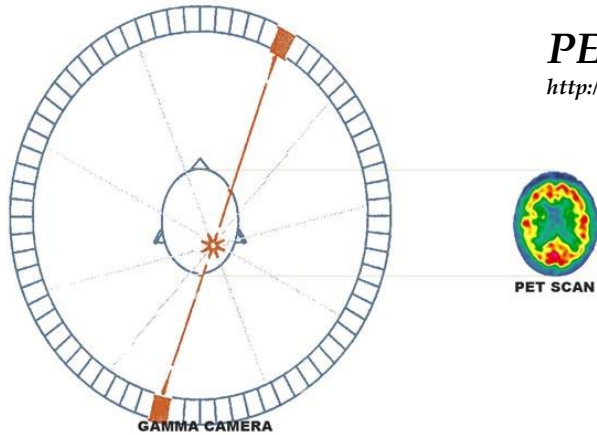
Dynamic Imaging: PET

Project 1

$3D \times t$

Dynamic Imaging: PET

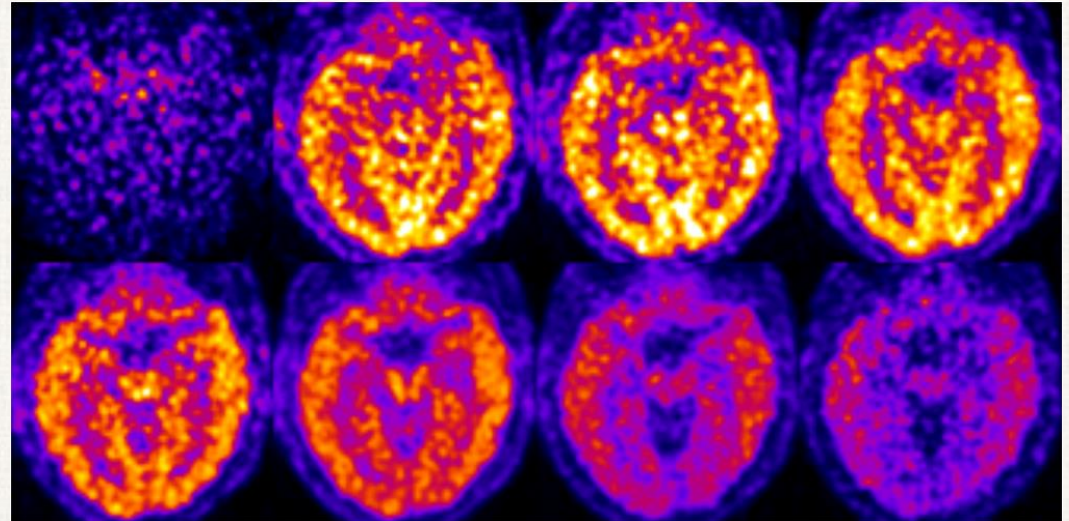
Project 1



PET: views from all angles available at all instances.

<http://www.whatisnuclearmedicine.com/What-is-62-How-does-it-work?&PHPSESSID=4f85d88a38d337434cfa6b2e95401e2>

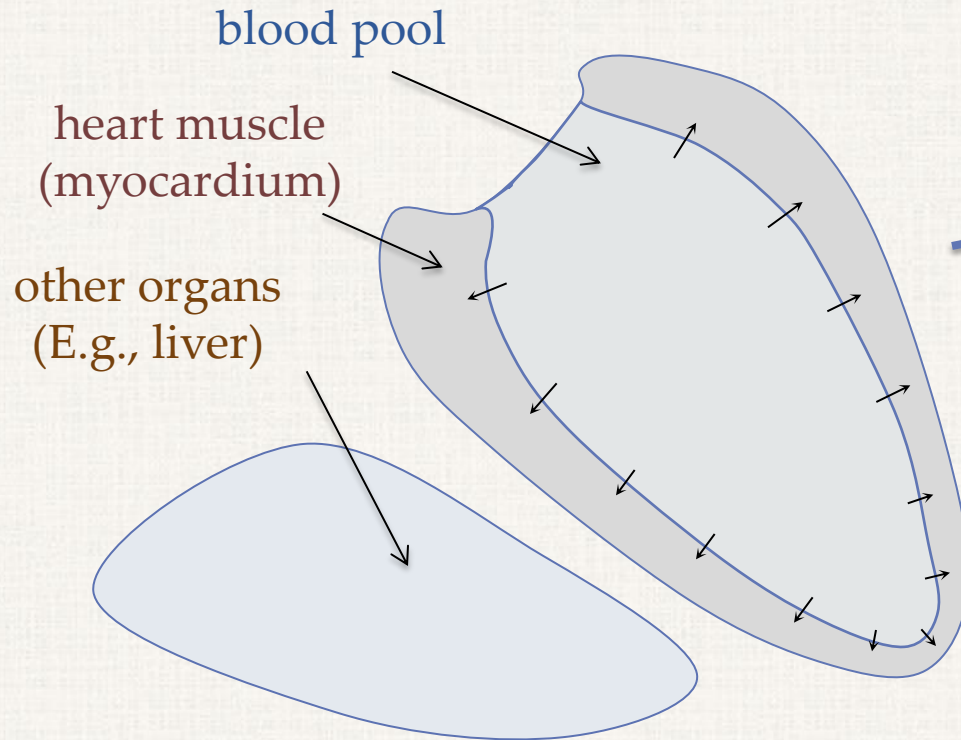
- *Time-lapsed Reconstructed 3D images, a slice through human brain*
- *Tracer concentration is changing with time*



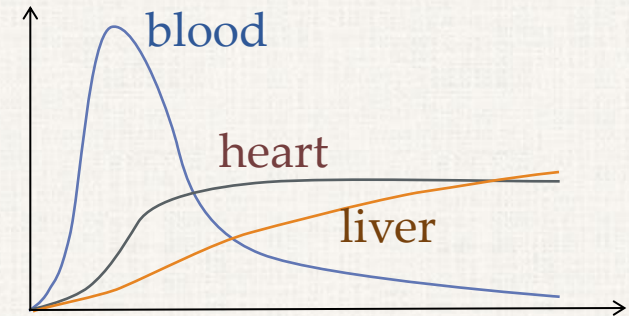
Dynamic Imaging- Challenges

- Low counts – high noise
 - *less time for data acquisition on each view*
 - *Ill-posed Problem*
- Lesser data/information in each time window after binning
 - *Underdetermined Problem*

Diagnostic Value of Dynamic Data



Time activity curves (TACs)



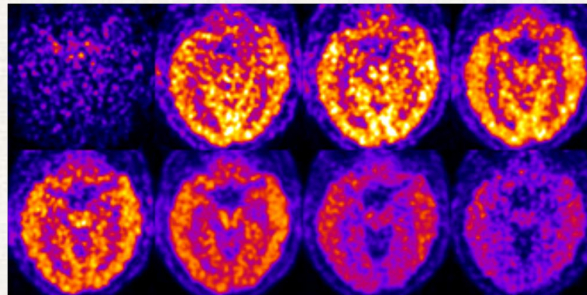
Local tracer exchange (kinetic) rates – important diagnostic parameters

CIFA Algorithm for Dynamic PET

Project 1

CIFA: Cluster-Initialized Factor Analysis

Input: 4D images of possible Alzheimer's patient



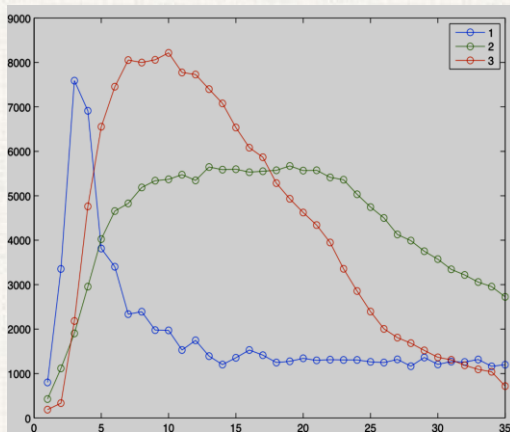
Output: Visualize affected tissues based on their tracer kinetics

*SPIE Medical Imaging Conference, (submitted) February 2015, Orlando,
R Bouthcko, D Mitra, H Pan, W Jagust, and GT Gullberg*

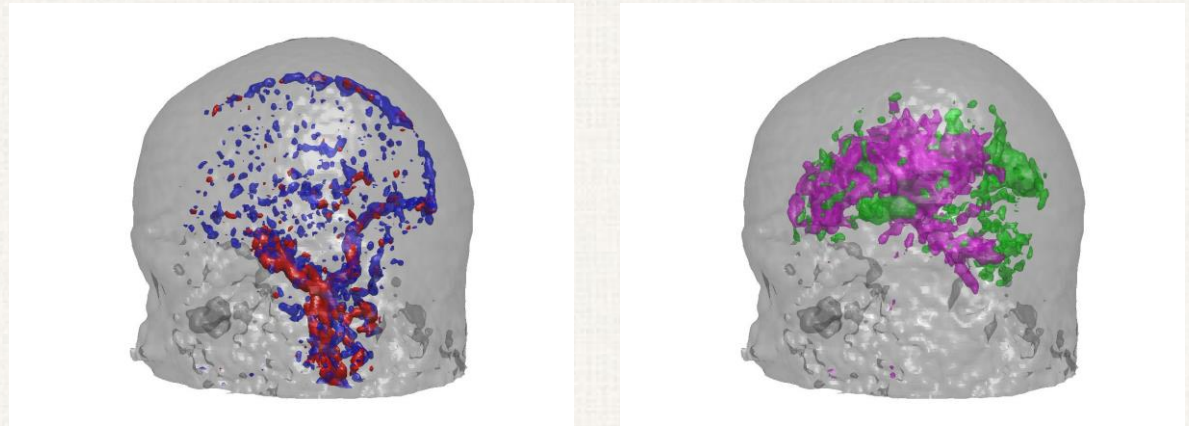
CIFA Algorithm for Dynamic PET

Project 1

*Output(1): Time-activity curves :
Carotid artery, Normal tissue, and Alzheimers affected tissue;*



Output (2): Corresponding segments



3D views of above tissues

*SPIE Medical Imaging Conference, (submitted) February 2015, Orlando,
R Bouthcko, D Mitra, H Pan, W Jagust, and GT Gullberg*

Dynamic Imaging: **SPECT**

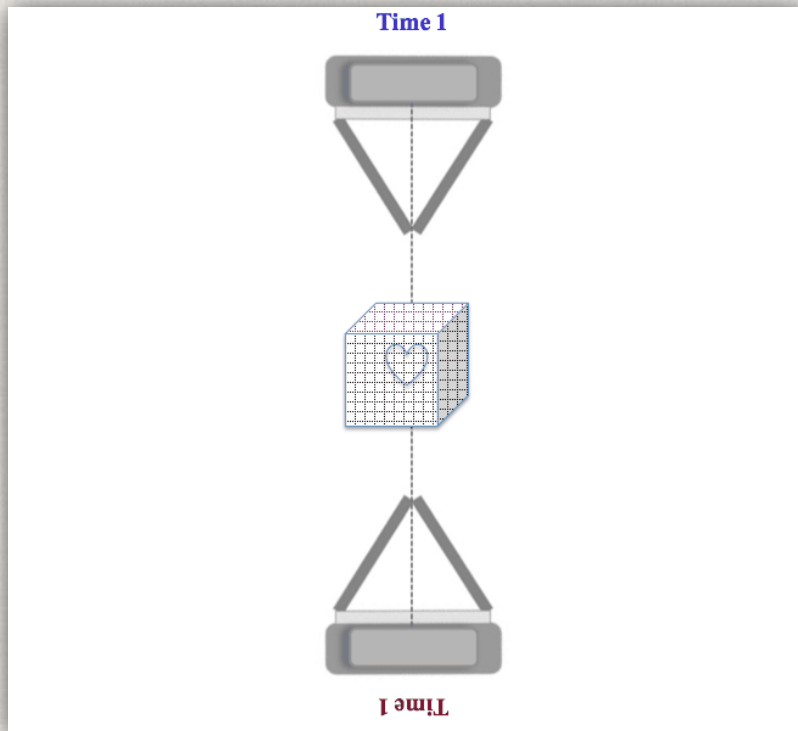
Project 2

$2D \times \theta \times t$

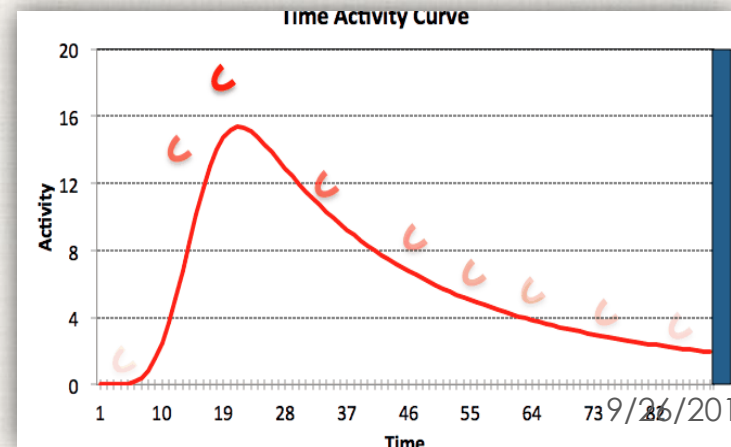
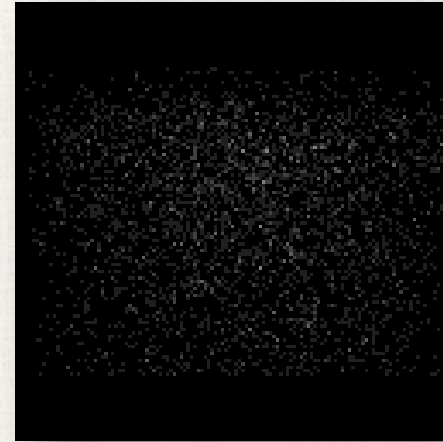
Dynamic Imaging: SPECT

Project 2

Only two projections for each time point:
more difficult than PET

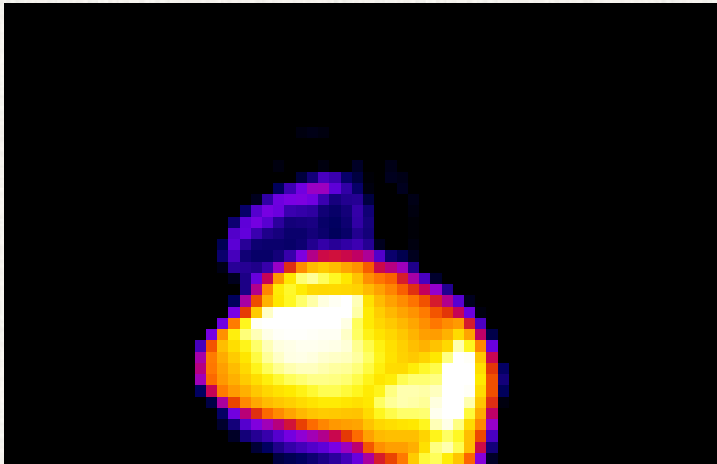


First Rotation Sinogram
Immediately after injection



Dynamic Vs. Static Projections

Static Sinogram



Dynamic Sinogram



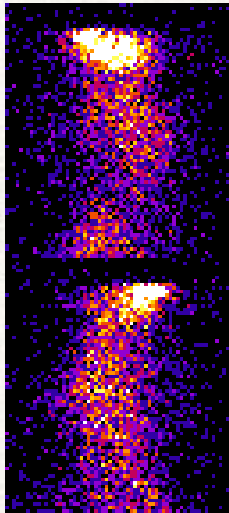
Dynamic SPECT – Additional Challenges

- Low counts – less time for data acquisition:
 - *Ill-posed Problem*
- Few projections for each time point:
 - *Underdetermined Problem*
- First rotation data, only two views per rotation:
 - *inconsistent*
- Small animal imaging:
 - *Low resolution & motion*

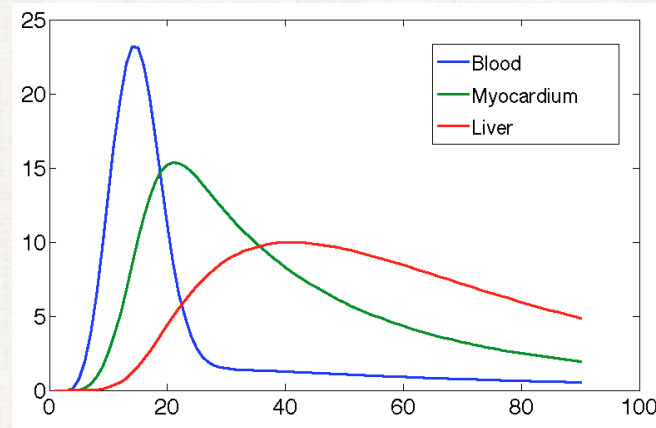
Dynamic SPECT: Task

- Goal: Estimation of tracer's **temporal** distribution in the imaged tissues *directly* from inconsistent projections

Input: Dynamic Sinogram



Output: Time Activity Curves (TACs)



Our contributions



- *SIFADS* (Spectral Initialized Factor Analysis of Dynamic Structures): sparsification with adaptable basis-functions
- Conditional Regularization for Constrained Optimization

Dynamic SPECT Model

- Dynamic SPECT is modeled by:
- 4D volume is factored with J time basis functions:

$$P_n(t) = \mathop{\mathring{a}}_{k=1}^K S_{n,k} V_k(t)$$

$$V_k(t) = \mathop{\mathring{a}}_{j=1}^J C_{k,j} f_{j,t}$$



$$P_n(t) = \mathop{\mathring{a}}_{k=1}^K S_{n,k} \mathop{\mathring{a}}_{j=1}^J C_{k,j} f_{j,t}$$



$$P = SCf$$

Space Time

- P : Sinogram as function of time
- S : System Matrix
- V : 4D Imaged volume, as function of time.
- n : pixel index on the detector
- k : voxel index on the volume
- f : Time basis functions
- C : Coefficients of time basis functions
- J : Number of time basis functions.

Existing Methods

- **Spectral Methods:**

- Select a set of representative time basis functions (Typically cubic b-splines). **Problem: what is the best set of basis functions?**

$$\underset{c}{\operatorname{arg\,min}} \left\{ \|SCf - P\|_w^2 \right\}$$

- **Factor Analysis of Dynamic Structures (FADS):**

- Initialize both time basis functions and coefficients with proper values. **Problem: what to initialize with?**

$$\underset{c,f}{\operatorname{arg\,min}} \left\{ \|SCf - P\|_w^2 + \text{Regularization} \right\}$$

Our Approach

Enhancements:

- Imposed Data-driven Prior information as constraints in optimization
- Combined two types of optimization techniques

Consequence:

Reduced dependence on initialization

Proposed Methods 1- Impose Prior information

- Reconstruction of later frames is segmented
- Segments are used to impose regularization functions:

1. An anisotropic total variation $Q(c) = |ATV(c)|_1$

2. Coefficients mix prevention $W(C) = |\vec{C}_j \cdot \vec{C}_i|_1 \quad j \neq i$

3. Curves' smoothness constraint $F(f) = |\nabla f|_1$

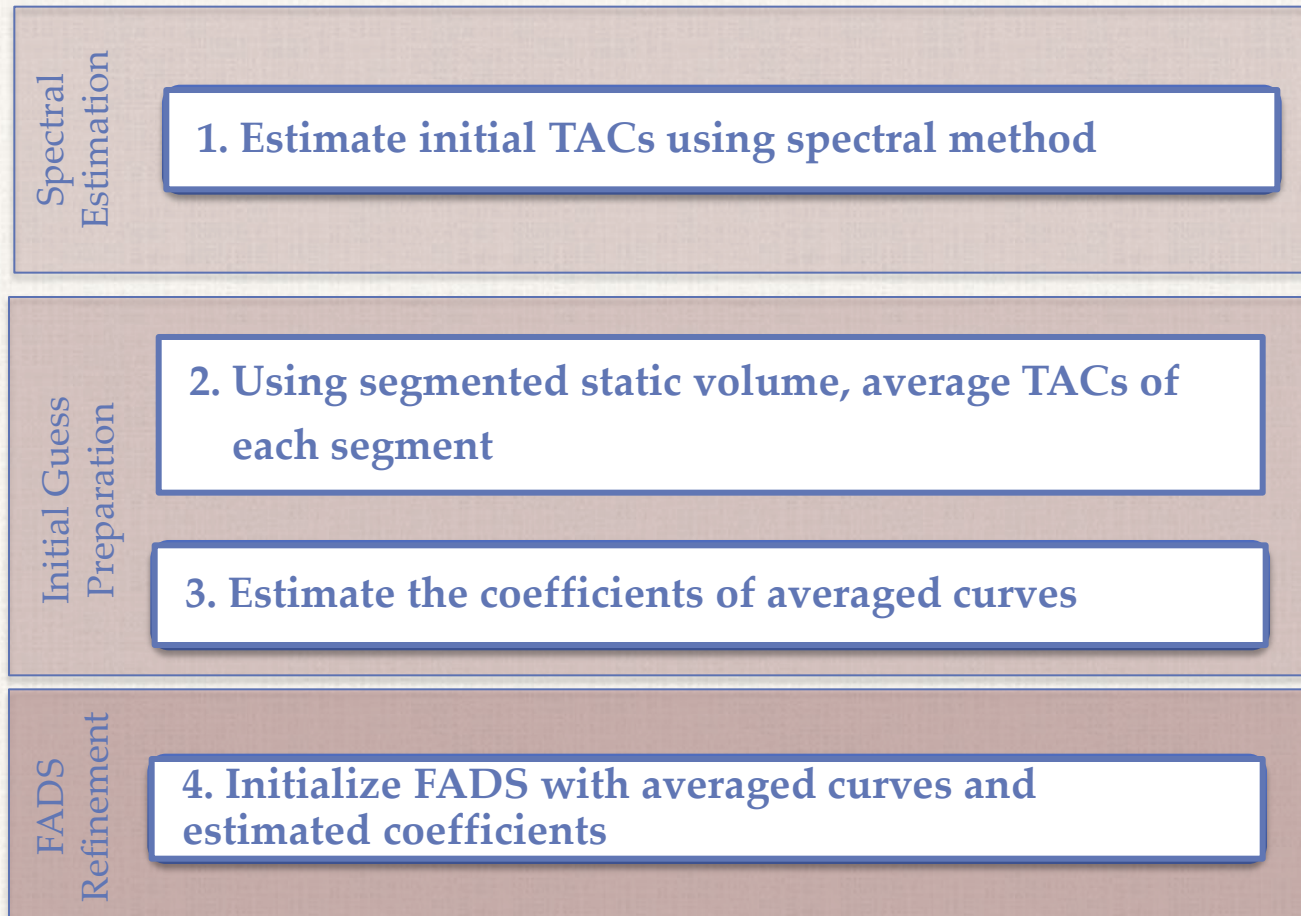
$$\operatorname{argmin} \left\{ \|SCf - p\|_w^2 + \underbrace{I_1 Q(c) + I_2 W(C)}_{\text{Spatial Regularization}} + \underbrace{I_3 F(f)}_{\text{Temporal Regularization}} \right\}$$

Spatial
Regularization

Temporal
Regularization

Proposed Methods 2– Hybrid Optimization

- Spectral Initialized FADS (**SIFADS**) algorithm



SIFADS Algorithm

SIFADS Algorithm

//STEP 1: Initialization:

//B-Splines fitting

- 1: $f^0 \leftarrow \{B - \text{Spline functions}\};$
- 2: $C^0 \leftarrow 0;$
- 3: $C^1 \leftarrow \arg \min_c \left\{ \|SC^0 f^0 - p\|_w^2 \right\}$

// Estimating initial Curves and Coefficients:

- 4: $V(t) \leftarrow C^1 f^0;$
- 5: $f^1 \leftarrow \text{Ave}(\text{segment}(V(t)));$
- 6: $C^2 \leftarrow \arg \min_c \left\{ \|SC^1 f^1 - p\|_w^2 + \lambda_1 \Theta(C^1) + \lambda_2 \Omega(C^1) + \lambda_3 \Phi(f^1) \right\}$

// STEP 2: FADS Refinement

- 7: $(C^*, f^*) \leftarrow \arg \min_c \left\{ \|SC^2 f^1 - p\|_w^2 + \lambda_1 \Theta(C^2) + \lambda_2 \Omega(C^2) + \lambda_3 \Phi(f^1) \right\}$

// Estimate and Output Final Curves:

- 8: $V(t) \leftarrow C^* f^*;$
- 9: $f \leftarrow \text{Ave}(\text{segment}(V(t)));$

MAP Algorithm for coefficients estimation

```

C ← 1;
for i = 1 to N do
    U(C[i]) ← λ1Ω(C[i]) + λ2Θ(C[i]);
    ∇U(C[i]) ←  $\frac{\partial U}{\partial C^{[i]}}$ ;
    C[i+1] =  $\frac{C^{[i]}}{\sum Sf + \nabla U(C^{[i]})} \sum \frac{P}{SC^{[i]}f} Sf;$ 
end for
return C
    
```

MAP Algorithm for coefficients and factors estimation

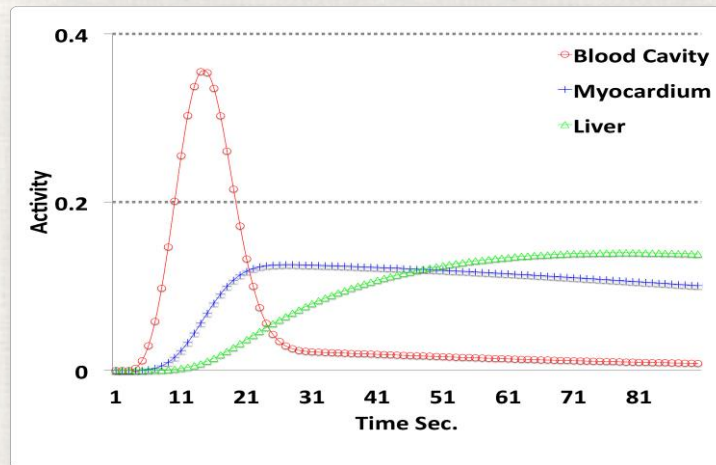
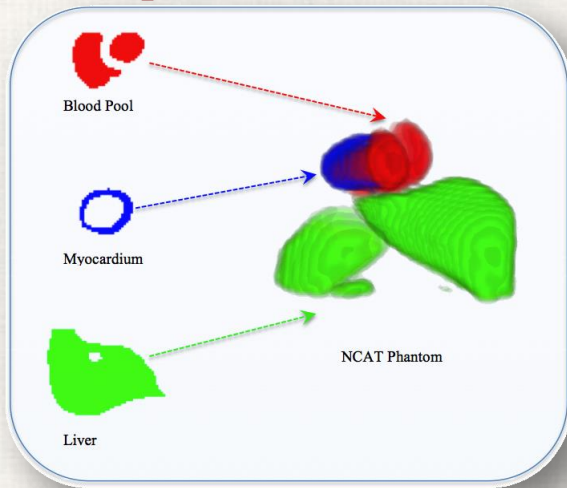
```

for i = 1 to N do
    // Coefficient minimization:
    U1(C[i]) ← λ1Ω(C[i]) + λ2Θ(C[i]);
    ∇U1(C[i]) ←  $\frac{\partial U_1}{\partial C^{[i]}}$ ;
    C[i+1] =  $\frac{C^{[i]}}{\sum Sf^{[i]} + \nabla U_1(C^{[i]})} \sum \frac{P}{SC^{[i]}f^{[i]}} Sf^{[i]};$ 

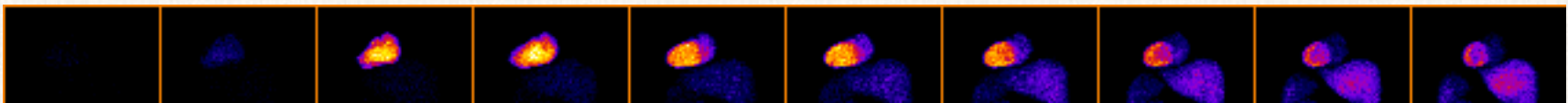
    // Factor minimization:
    U2(f[i]) ← λ3Φ(f[i]);
    ∇U2(f[i]) ←  $\frac{\partial U_2}{\partial f^{[i]}}$ ;
    f[i+1] =  $\frac{f^{[i]}}{\sum SC^{[i+1]} + \nabla U_2(f^{[i]})} \sum \frac{P}{SC^{[i+1]}f^{[i]}} SC^{[i+1]};$ 
end for
return C, f
    
```

Validation with Simulation

Coefficients used for simulation
(NCAT phantom)



Generated projections with Poisson noise



0

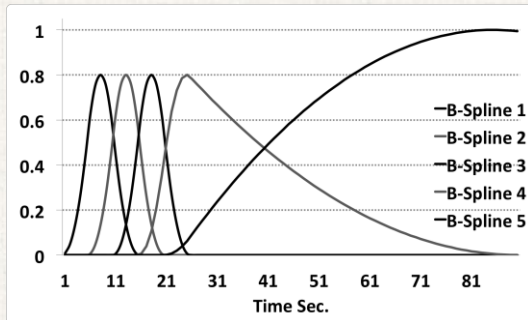
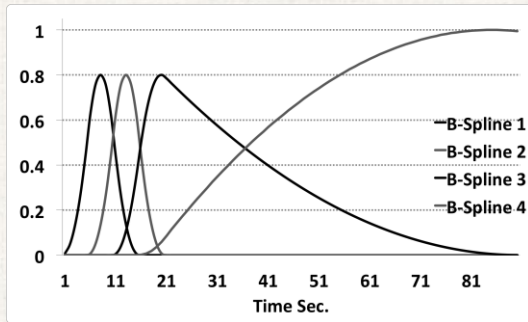
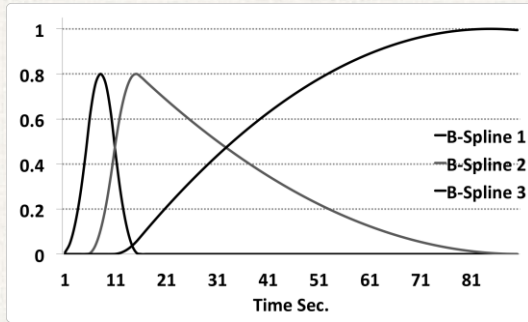
20

40

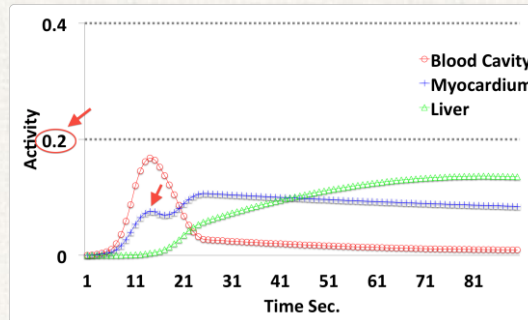
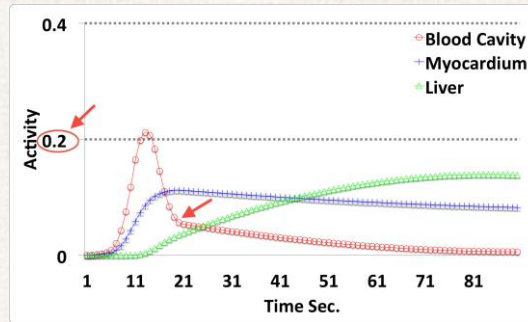
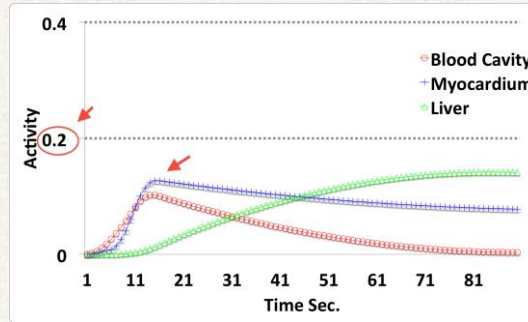
Time Sec.

Spline vs. SIFADS results

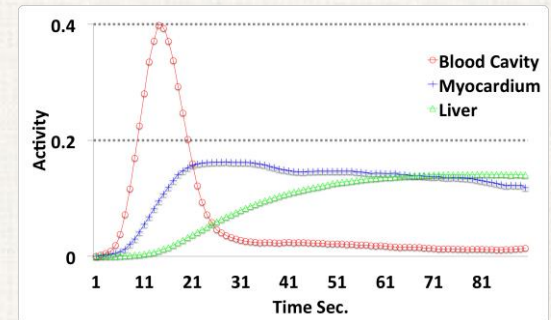
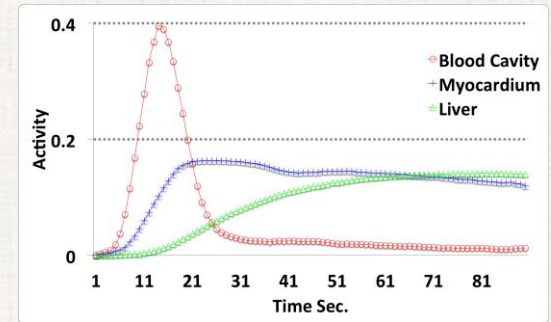
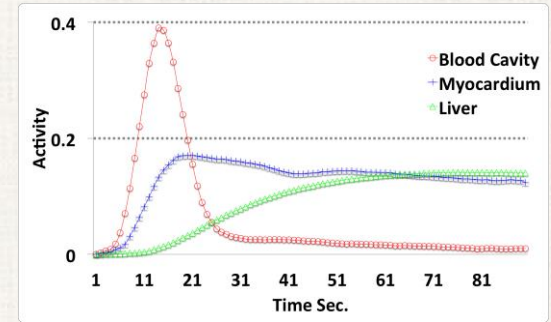
B-splines



Spectral



SIFADS



Real Data: Rat heart

- Dynamic, pinhole SPECT study, rat's heart
- Collimators: 1.5×2 mm tungsten pinholes
- GE VG3 Millennium Hawkeye camera
- Acquisition started with injection of 7 mCi ^{123}I -MIBG
- 30 rotations, 90 one-second views, per rotation
- Detector pixel: 4.42 mm, recon voxel 0.8 mm



Original projections:

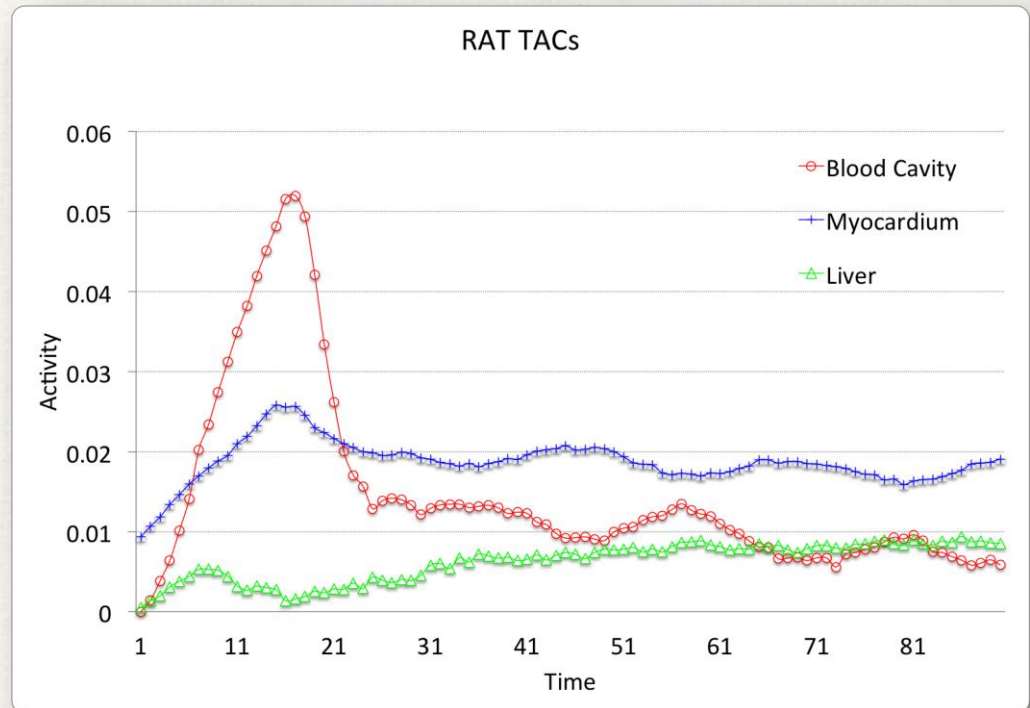


Reproduced projections by forward projecting dynamic reconstruction:



Results from Rat data

Estimated rat TACs from the first inconsistent rotation:



Cell Tracking on Fluorescent Microscopy

Project 3

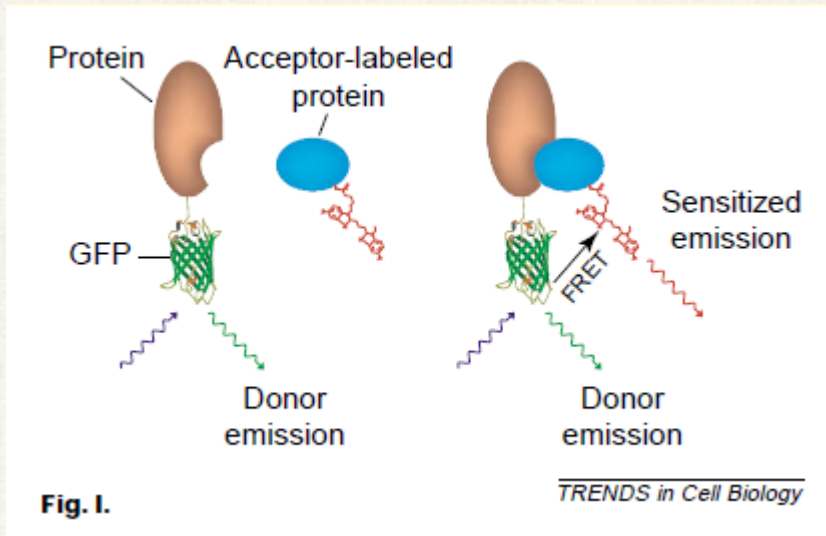
2D x t

Cell Tracking on Fluorescent Microscopy

Project 3

FRET: Fluorescence Resonance Energy Transfer

New Technology for quantifying gene expression in live single cells



$$E_A(i) = \frac{F^{DA}(i) - F^D(i) \cdot R_D - F^A(i) \cdot R_E}{F^A(i)}$$

Three channels for each Time-frame:

Donor emission (F_d), Acceptor Emission (F_a), D-to-A Excitation emission (F_{da})

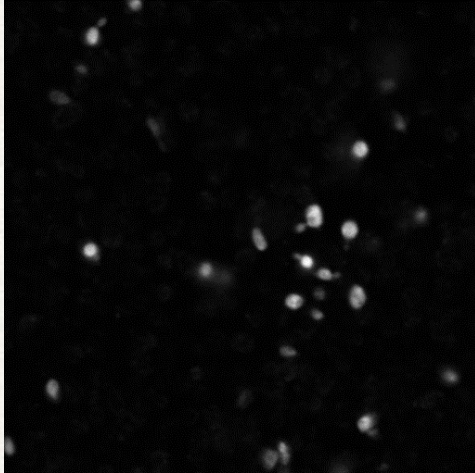
"Imaging biochemistry inside cells"

TRENDS in Cell Biology, 11(5): 203-211, 2011

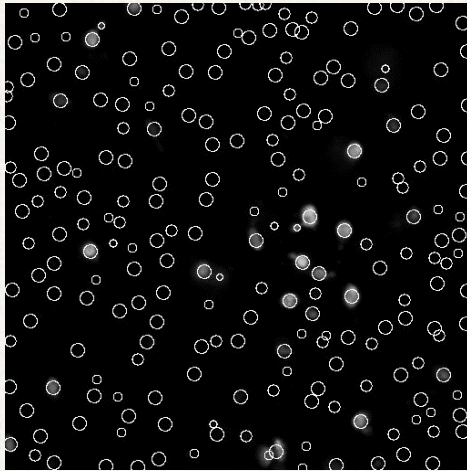
Wouters, Verveer and Bastiaens

Cell Tracking on Fluorescent Microscopy

Project 3



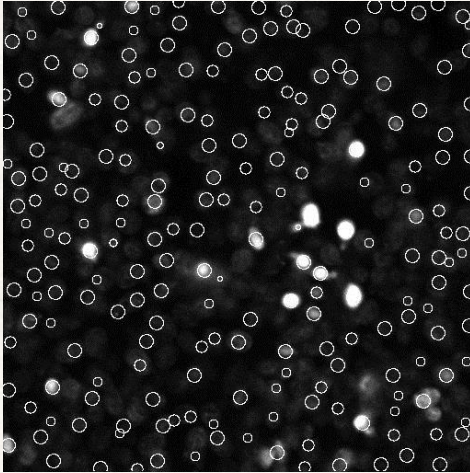
Input: frames of the time-lapsed 2D image from a confocal microscope



Output: same frames after tracking by scale-space segmentation

Cell Tracking on Fluorescent Microscopy

Project 3



Scale Space Algorithm: handles varying sizes of cells

Problem:

Live cells move in 3D – across the frame, in-out of focal plane;

Cells also divide!

How to track a cell from frame to frame?

Semi-solved:

Search around a cell in next frame for similar average intensity

*SPIE Medical Imaging Conference, (submitted) February 2015, Orlando,
Debasis Mitra, Rostyslav Bouthcko, Judhajeet Ray, and Marit Nilsen-Hamilton*

Future Works with SIFADS

- Use different imaging data:
PET, CT, Microscopy
- Use different basis function types:
wavelets or other non-orthogonal bases
- Use different objective functions:
dynamic data is very low intensity – use entropy
- Use different optimization techniques:
primal-dual algorithm shows promise
- Use different parallelization platforms: GPU

Tomography Beyond Medicine: Inverse Problems with similar mathematics - Linear Algebra, Statistics, Numerical Optimization, ...

- *Muon tomography: cosmic ray-generated muon scattered from heavy metals*
- *Electron Microscopy: to “see” 3D view of a virus or molecule*
- *Seismic: Acoustic waves from earthquake or artificial source to study subsurface structure*
- *Cosmology: Structure of the universe from telescopic observations*



Graduate Students

Mahmoud Abdalah

Hui Pan

Bo Li

Shi Chen

Collaborators



Grant T. Gullberg

Rostyslav Boutchko

*Lawrence Berkeley National
Lab, Berkeley, CA*

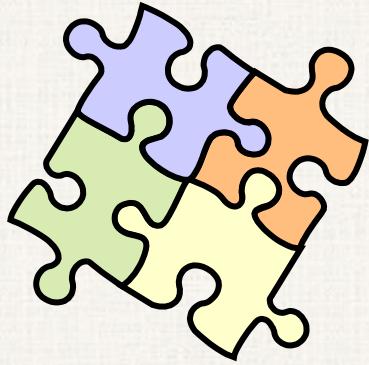


Youngho Seo

Sreejita Bannerjee

*University of California
San Francisco, CA*

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Thanks

Questions



Contact: dmitra@cs.fit.edu