

Estimation, Detection and Filtering of Medical Images

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Outline

- Part I : Basics of Medical image Filtering and Convolution
- Part II : Estimation Theory and examples
- Part III: Detection Theory and examples

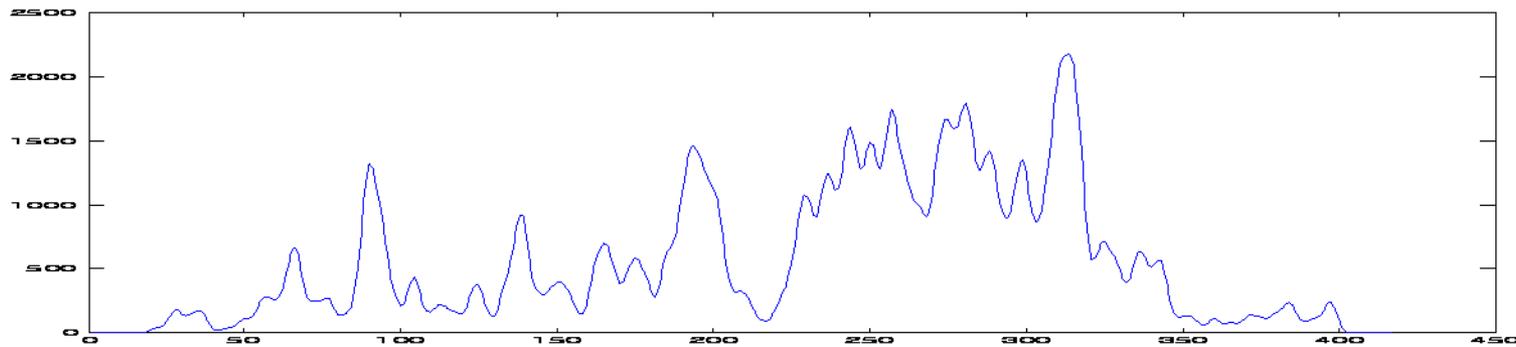
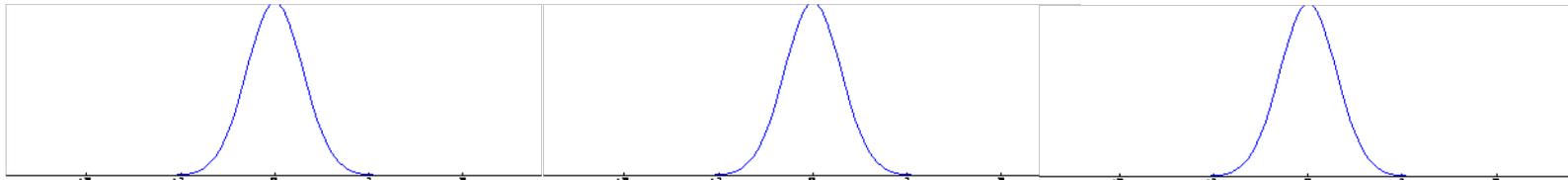
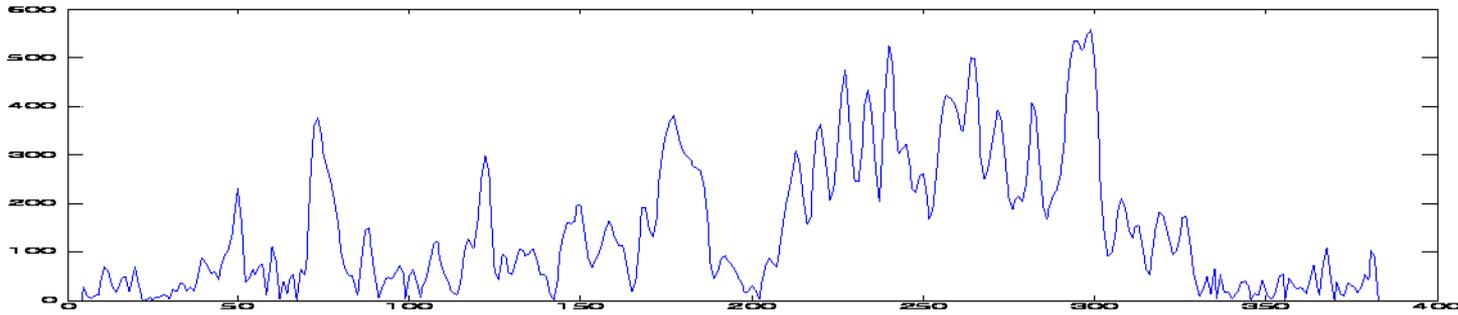
Convolution Basics

- Convolution is defined as

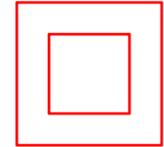
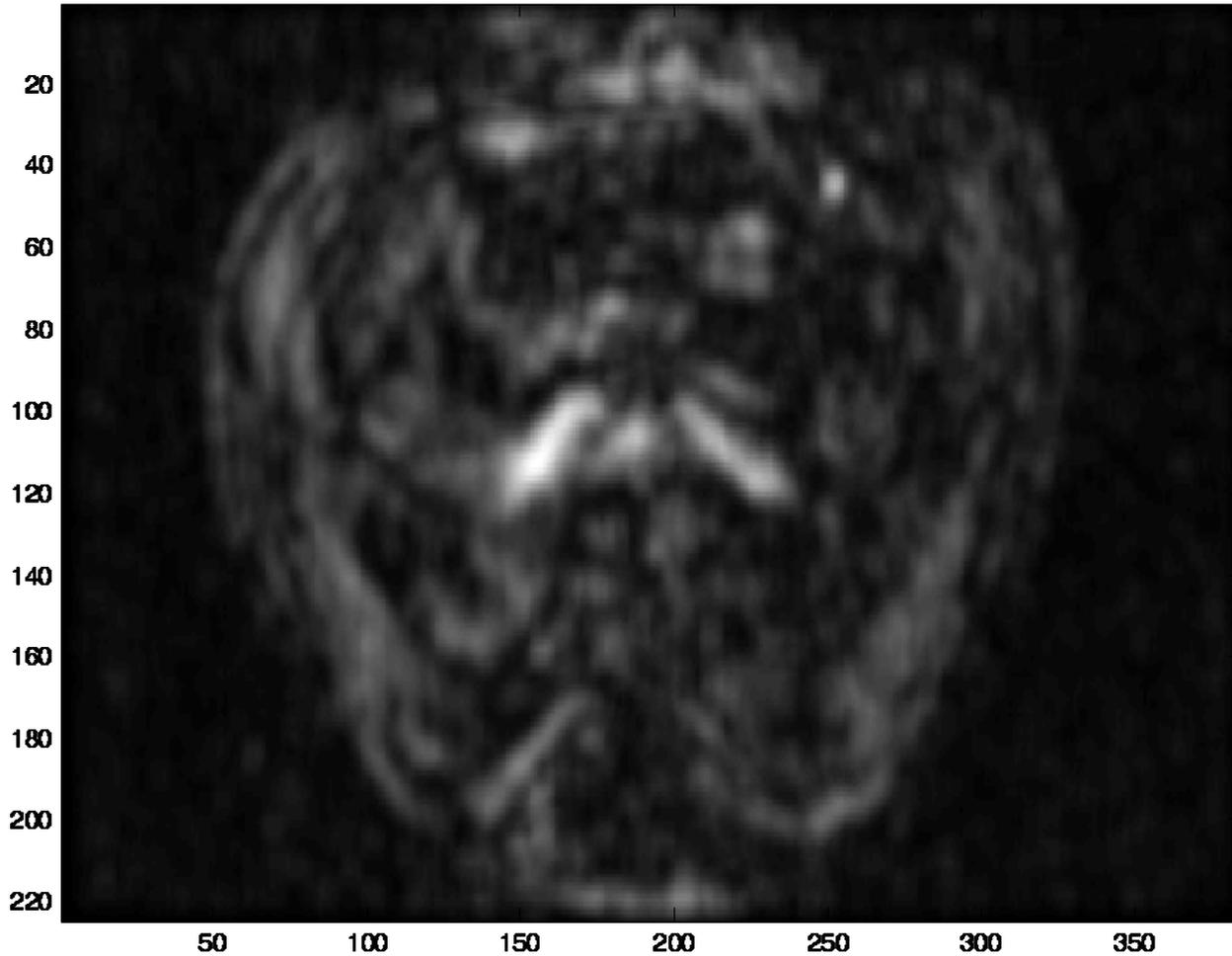
$$y(t) = x(t) * h(t) = \int_{-\infty}^{\infty} x(\tau)h(t - \tau)d\tau$$

- Practically achieved as follows:
 - Flip $h(t)$
 - Slide it into $x(t)$ by amount τ
 - At each position τ , calculate the area of overlap between x and h

Example



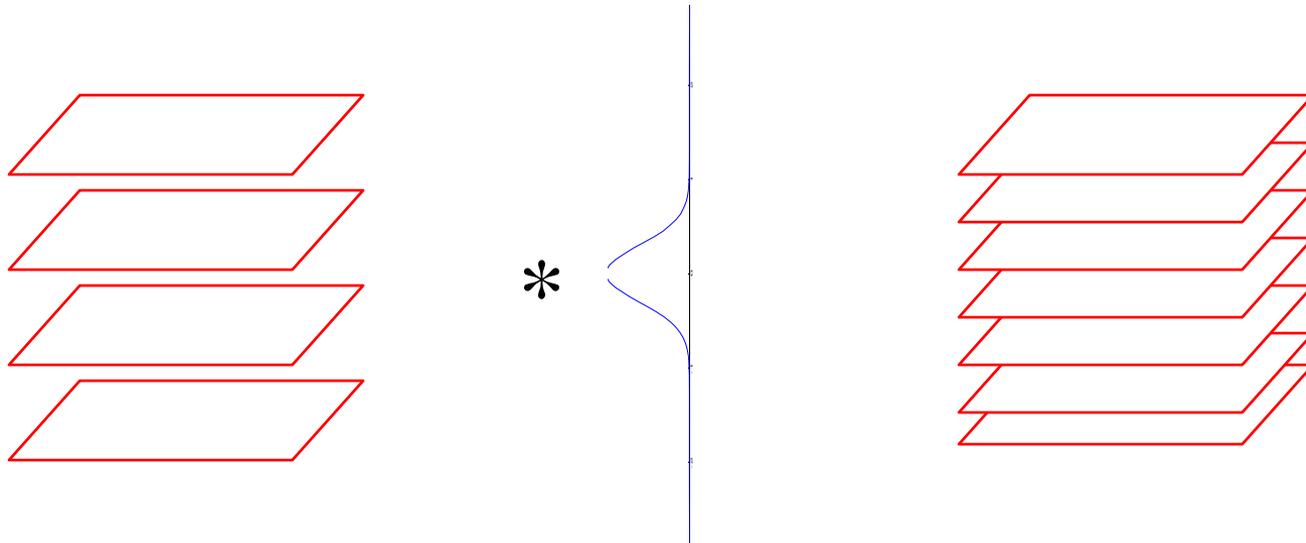
Filtering and Convolution



Convolution For Interpolation and Resampling

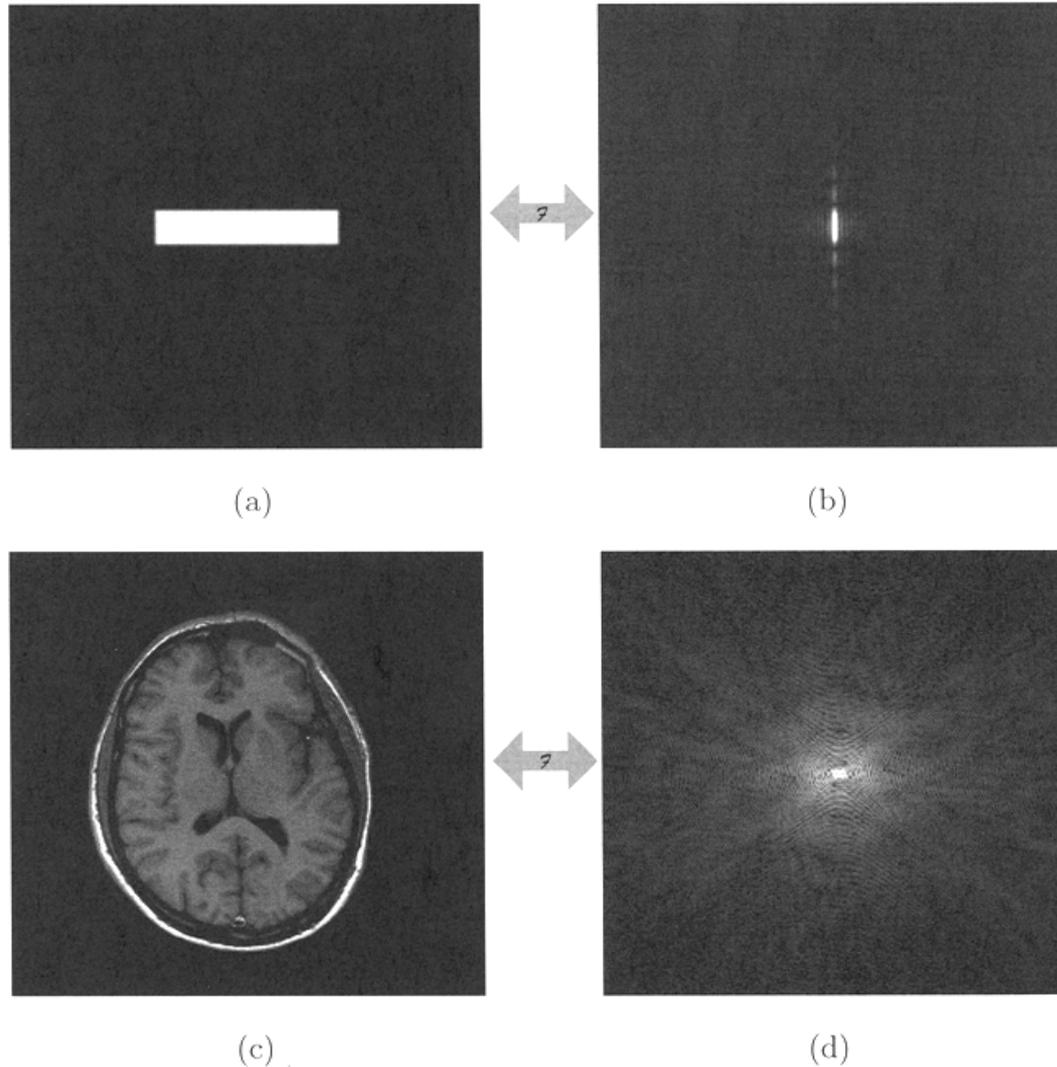
- Sometimes need to “fill in” missing data
- Interpolation – to resample image on finer grid
- Resampling is used to change the “nominal” resolution of images
- Example: if multi-slice images with non-isotropic resolution, resampling can make it isotropic
- **IMPORTANT:** resampling, filtering or interpolation does NOT increase “actual” resolution

Resampling example



k-space and image-space

k-space & image-space are related by the 2D FT



How many points do we need to sample?

- $\Delta k = 1/FOV$
- Why? Due to the Sampling Theorem

“Suppose a signal $I(x)$ is non-zero only within $[-W/2, W/2]$. Then its Fourier transform $F(I)(k_x)$ must be sampled at least as densely as $\Delta k_x = 1/W$.”

- Note this works regardless of direction of transform (Duality property)
- What happens if this is violated? ALIASING

Aliasing Example

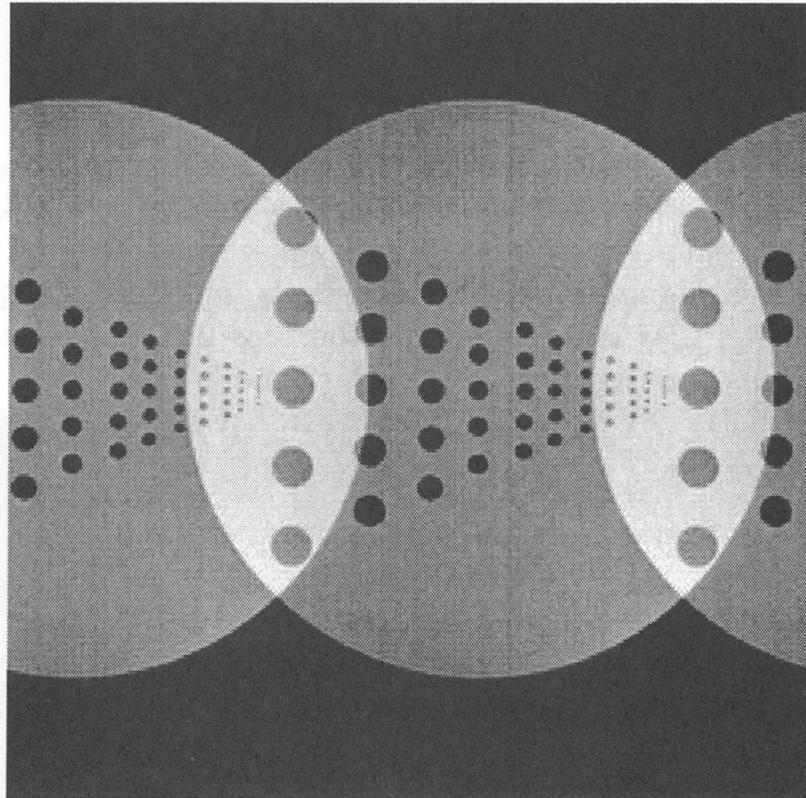
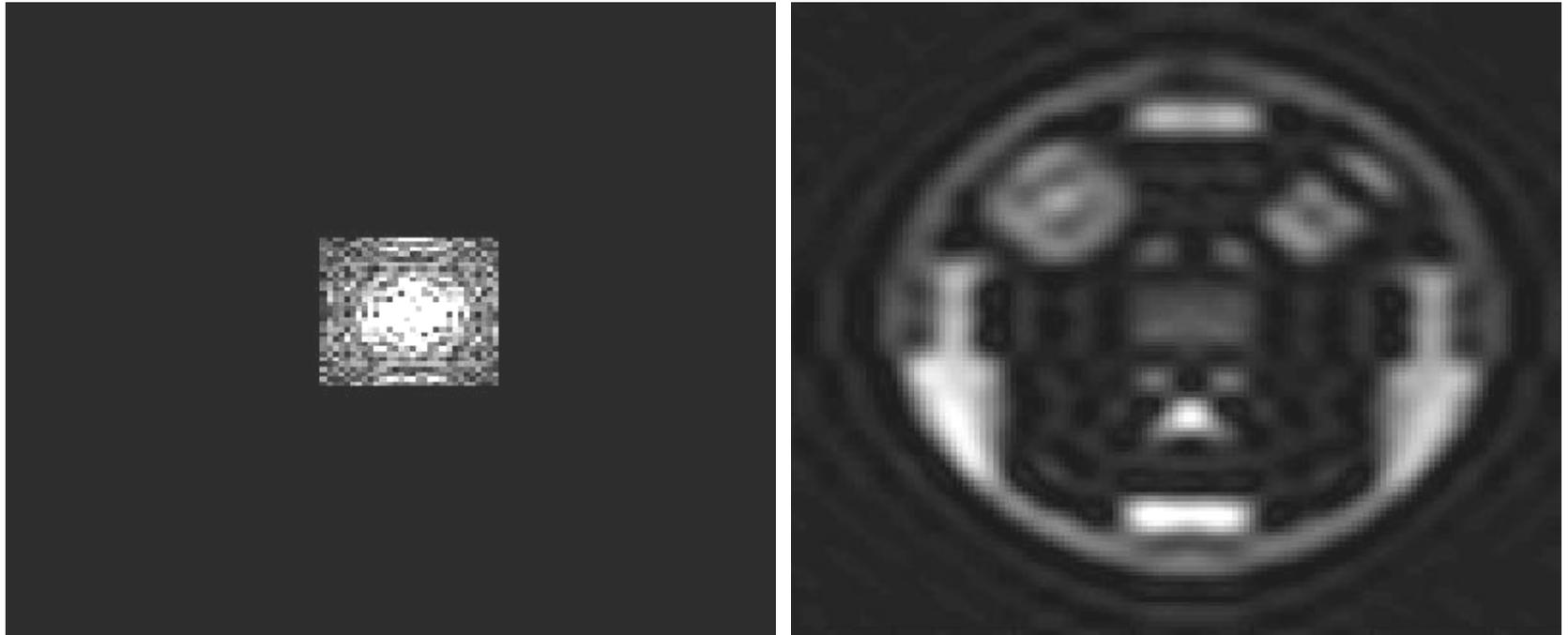


Figure 8.9 Aliasing artifacts due to undersampling along the horizontal direction by a factor of two.

Truncation

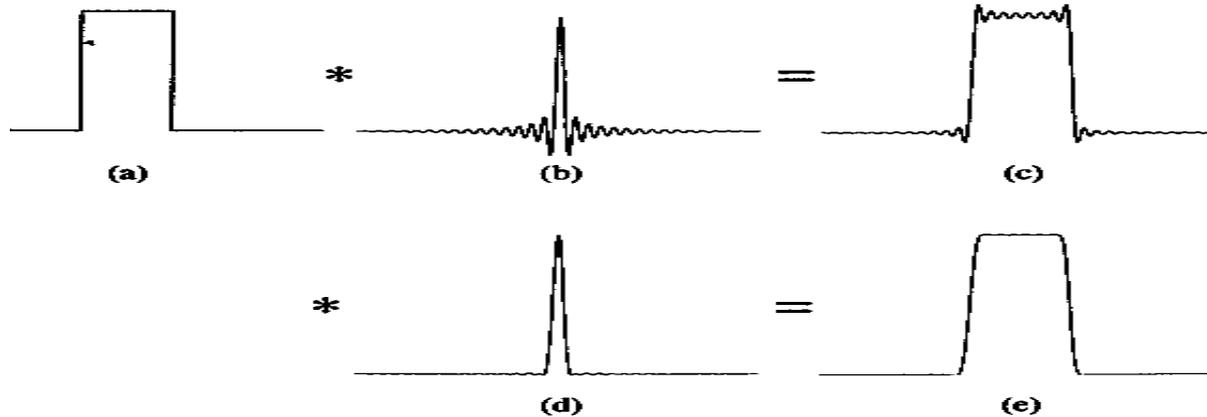
- Truncation = sampling central part of k-space



Truncation

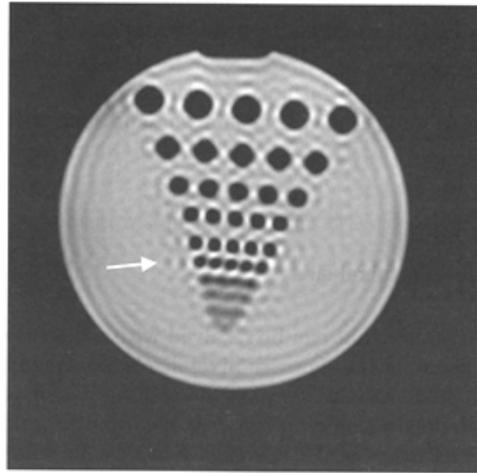
- Both blurring and ringing are a result of truncation
- Intrinsic resolution = size of the blur
- To reduce blur (hence increase resolution) we need to sample up to a larger k-space radius
- Can characterise resolution by the point spread function (PSF) which is simply the blurring kernel
- Note: zero-padding can increase matrix size but can not increase resolution!

Ringling

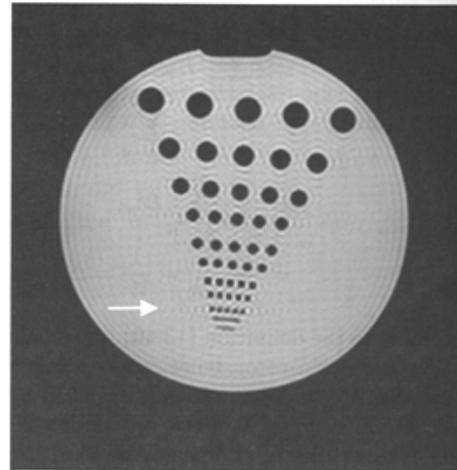


- ringing can be reduced by multiplying the signal by a smooth window - called windowing
- Popular window choices:
 - Kaiser-Bessel
 - Hanning
 - Raised cosine

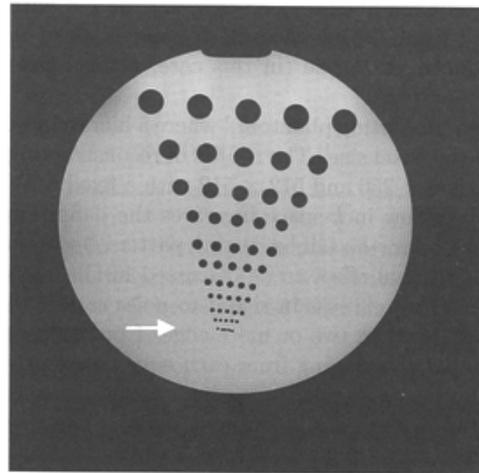
Ringing Example



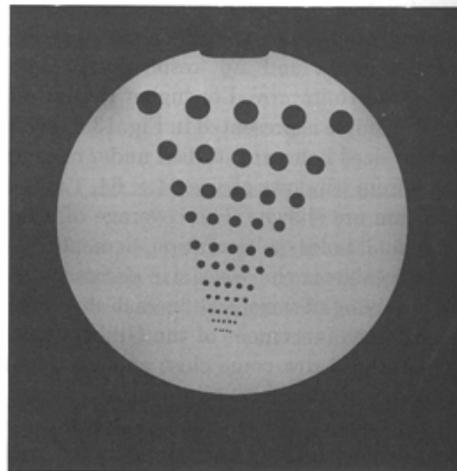
(a)



(b)



(c)



(d)

Part II : Estimation Theory and Examples

- Introduction to optimal estimates
- Different types of optimal estimates
- Estimation examples from MR

Estimation Theory

- Consider a linear process

$$y = H \theta + n$$

y = observed data

θ = set of model parameters

n = additive noise

- Then Estimation is the problem of finding the statistically optimal θ , given y , H and knowledge of noise properties
- MR is full of estimation problems

Different approaches to estimation

- Minimum variance unbiased estimators
- Least Squares
- Maximum-likelihood
- Maximum entropy
- Maximum a posteriori

*has no
statistical
basis*

*uses knowledge
of noise PDF*

*uses prior
information
about θ*

Least Squares Estimator

- Least Squares:

$$\theta_{LS} = \operatorname{argmin} \|y - H\theta\|^2$$

- Natural estimator– want solution to match observation
- Does not use any information about n
- There is a simple solution (a.k.a. pseudo-inverse):

$$\theta_{LS} = (H^T H)^{-1} H^T y$$

What if we know something about the noise?

Say we know $\Pr(n)$...

Maximum Likelihood Estimator

- Simple idea: want to maximize $\Pr(y|\theta)$
- Can write $\Pr(n) = e^{-L(n)}$, $n = y - H\theta$, and
$$\Pr(n) = \Pr(y|\theta) = e^{-L(y, \theta)}$$
- if white Gaussian n , $\Pr(n) = e^{-\|n\|^2/2\sigma^2}$ and
$$L(y, \theta) = \|y - H\theta\|^2/2\sigma^2$$
$$\theta_{ML} = \operatorname{argmax} \Pr(y|\theta) = \operatorname{argmin} L(y, \theta)$$
 - called the likelihood function
$$\theta_{ML} = \operatorname{argmin} \|y - H\theta\|^2/2\sigma^2$$
- This is the same as Least Squares!

Maximum Likelihood Estimator

- But if noise is jointly Gaussian with cov. matrix C
- Recall C , $E(nn^T)$. Then

$$\Pr(n) = e^{-\frac{1}{2} n^T C^{-1} n}$$

$$L(y|\theta) = \frac{1}{2} (y-H\theta)^T C^{-1} (y-H\theta)$$

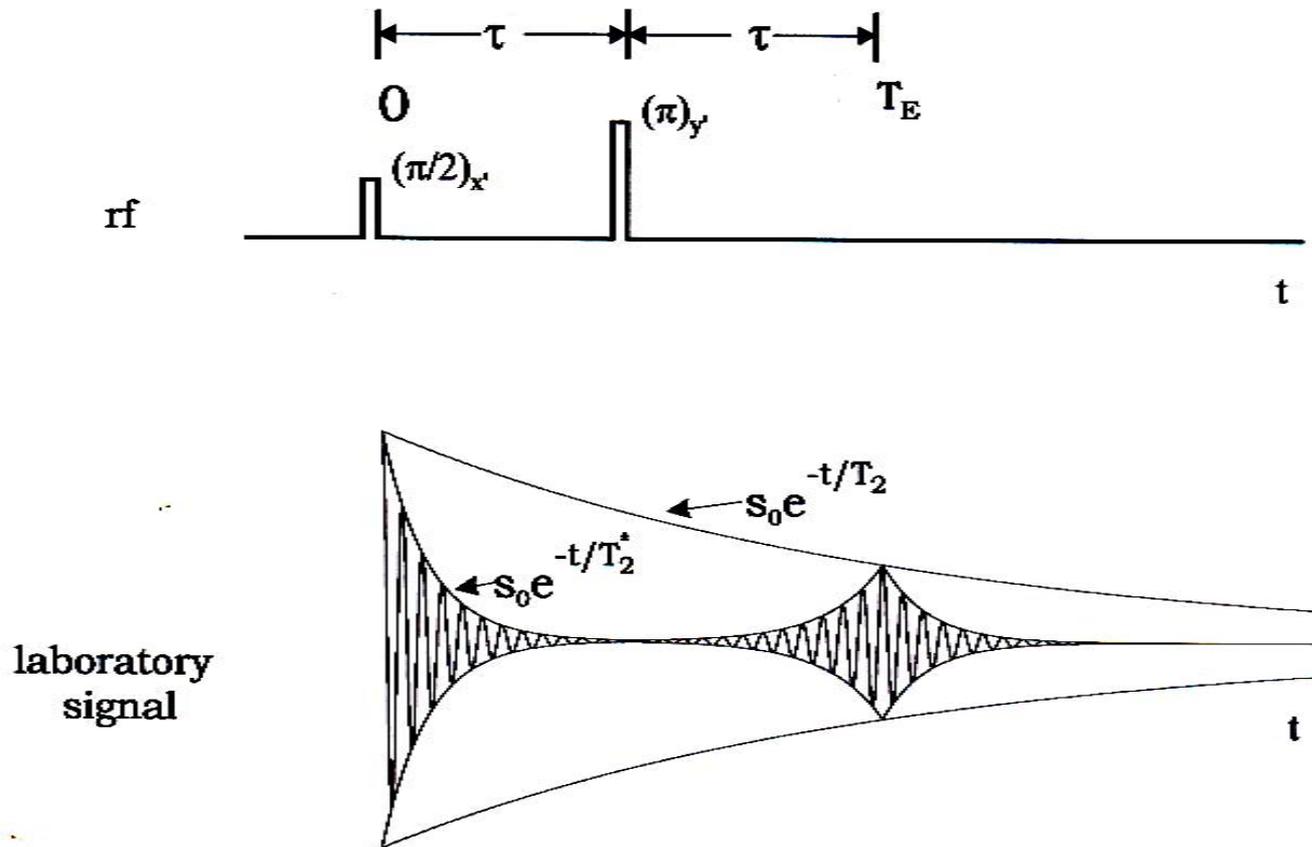
$$\theta_{ML} = \operatorname{argmin} \frac{1}{2} (y-H\theta)^T C^{-1} (y-H\theta)$$

- This also has a closed form solution

$$\theta_{ML} = (H^T C^{-1} H)^{-1} H^T C^{-1} y$$

- If n is not Gaussian at all, ML estimators become complicated and non-linear
- Fortunately, in MR noise is usually Gaussian

Example - estimating T_2 in repeated spin echo data



Example – estimating T_2 in repeated spin echo data

$$s(t) = e^{-t/T_2} \int dr \rho(r)$$

- Need only 2 data points to estimate T_2 :

$$T_{2\text{est}} = [T_{E2} - T_{E1}] / \ln[s(T_{E1})/s(T_{E2})]$$

- However, not good due to noise, timing issues
- In practice we have many data samples from various echoes

Example – estimating T_2

$$y \rightarrow \begin{pmatrix} \ln(s(t_1)) \\ \ln(s(t_2)) \\ \vdots \\ \ln(s(t_n)) \end{pmatrix} = \begin{pmatrix} 1 & -t_1 \\ 1 & -t_2 \\ \vdots & \vdots \\ 1 & -t_n \end{pmatrix} \begin{pmatrix} a \\ r \end{pmatrix}$$

H (points to the matrix)
 θ (points to the vector)

Least Squares estimate:

$$\theta_{LS} = (H^T H)^{-1} H^T y$$

$$T_2 = 1/r_{LS}$$

Can we do better by ML estimate?

- if noise is correlated across time
- if noise variance changes over time

Estimation example - Denoising

- Suppose we have a noisy MR image y , and wish to obtain the noiseless image x , where

$$y = x + n$$

- Can we use Estimation theory to find x ?
- Try: $H = I$, $\theta = x$ in the linear model
- Both LS and ML estimators simply give $x = y$!
- → we need a more powerful model
- Suppose the image x can be approximated by a polynomial, i.e. a mixture of 1st p powers of r :

$$x = \sum_{i=0}^p a_i r^i$$

Example – denoising

$$y \rightarrow \begin{pmatrix} y_1 \\ y_2 \\ \vdots \\ y_n \end{pmatrix} = \begin{pmatrix} 1 & r_1^1 & \dots & r_1^p \\ 1 & r_2^1 & \dots & r_2^p \\ \vdots & \vdots & & \vdots \\ 1 & r_n^1 & \dots & r_n^p \end{pmatrix} \begin{pmatrix} a_0 \\ a_1 \\ \vdots \\ a_p \end{pmatrix} + \begin{pmatrix} n_1 \\ n_2 \\ \vdots \\ n_n \end{pmatrix}$$

H (points to the matrix) and θ (points to the vector a)

Least Squares estimate:

$$\theta_{LS} = (H^T H)^{-1} H^T y$$

$$x = \sum_{i=0}^p a_i r^i$$

Can we do better by ML estimate? YES

Noise in MR can be spatially correlated

- ML with covariance matrix C is better

Multi-variate FLASH

- Acquire 6-10 accelerated FLASH data sets at different flip angles or TR's
- Generate T_1 maps by fitting to:

$$S = \exp\left(-TE/T_2^*\right) \sin \alpha \frac{1 - \exp(-TR/T_1)}{1 - \cos \alpha \exp(-TR/T_1)}$$

- Not enough info in a single voxel
- Noise causes incorrect estimates
- Error in flip angle varies spatially!

Spatially Coherent T_1 , ρ estimation

- First, stack parameters from all voxels in one big vector \mathbf{x}
- Stack all observed flip angle images in \mathbf{y}
- Then we can write $\mathbf{y} = \mathbf{M}(\mathbf{x}) + \mathbf{n}$
- Recall \mathbf{M} is the (nonlinear) functional obtained from

$$S = \exp(-TE/T_2^*) \sin \alpha \frac{1 - \exp(-TR/T_1)}{1 - \cos \alpha \exp(-TR/T_1)}$$

- Solve for \mathbf{x} by non-linear least square fitting, PLUS spatial prior:

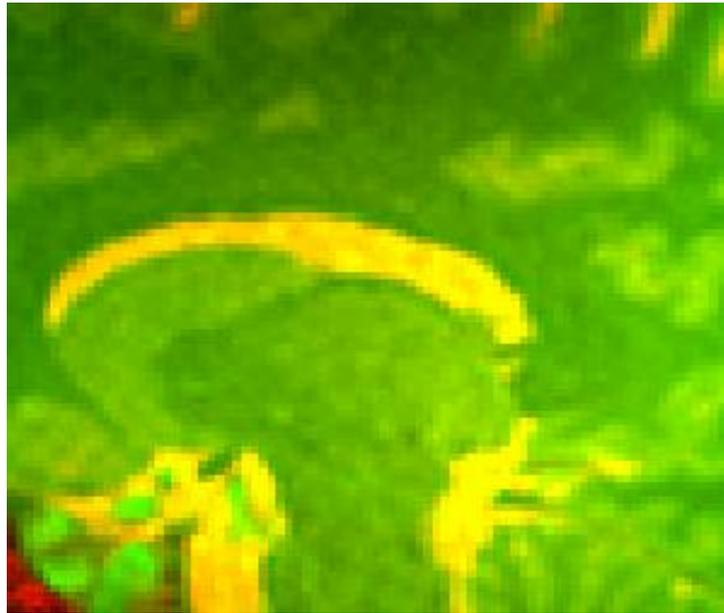
$$\mathbf{x}_{\text{est}} = \arg \min_{\mathbf{x}} \left(\|\mathbf{y} - \mathbf{M}(\mathbf{x})\|^2 + \mu^2 \|D\mathbf{x}\|^2 \right) \leftarrow E(\mathbf{x})$$

Makes $\mathbf{M}(\mathbf{x})$ close to \mathbf{y}

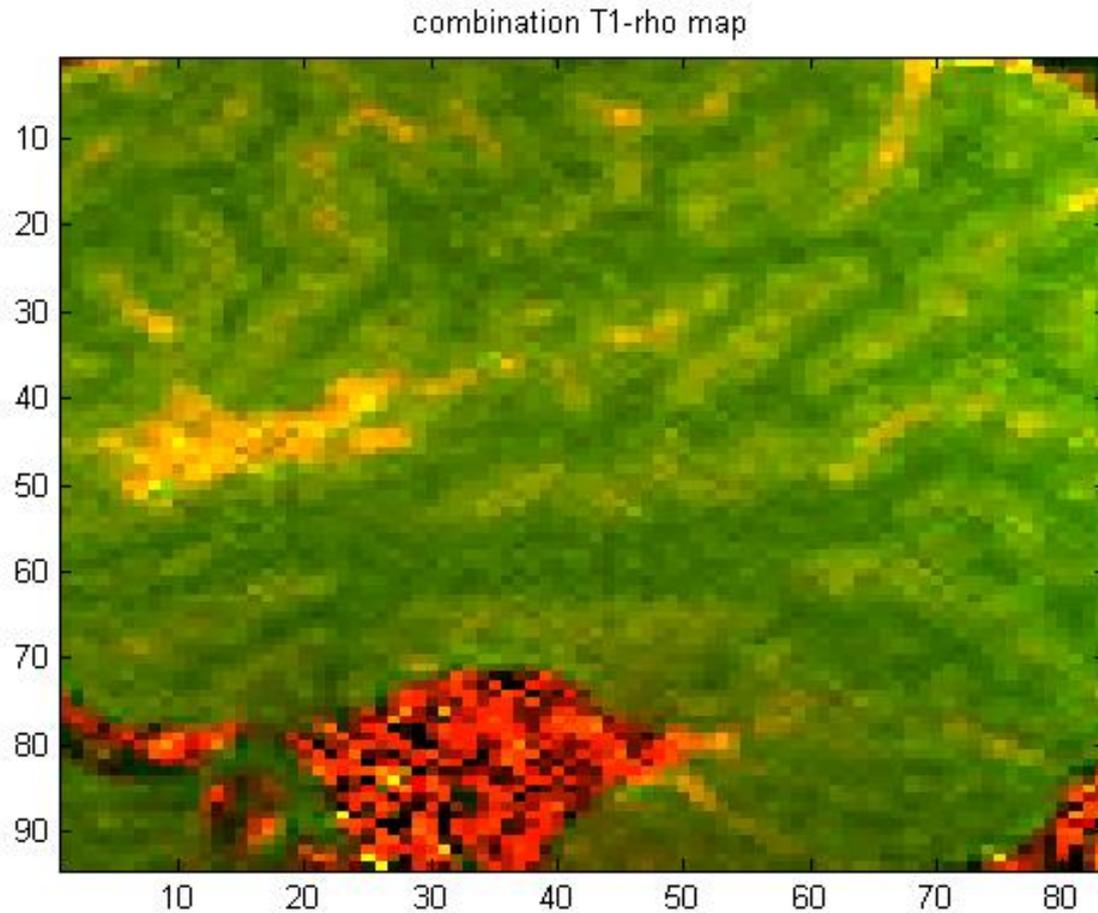
Makes \mathbf{x} smooth

- Minimize via MATLAB's *lsqnonlin* function
- How? Construct $\delta = [\mathbf{y} - \mathbf{M}(\mathbf{x}); \mu D\mathbf{x}]$. Then $E(\mathbf{x}) = \|\delta\|^2$

Multi-Flip Results – combined ρ , T_1 in pseudocolour



Multi-Flip Results – combined ρ , T_1 in pseudocolour



Maximum a Posteriori Estimate

- This is an example of using an image prior
- Priors are generally expressed in the form of a PDF $\Pr(x)$
- Once the likelihood $L(x)$ and prior are known, we have complete statistical knowledge
- LS/ML are suboptimal in presence of prior
- MAP (aka Bayesian) estimates are optimal

Bayes Theorem:

$$\Pr(x|y) = \frac{\Pr(y|x) \cdot \Pr(x)}{\Pr(y)}$$

posterior (points to $\Pr(x|y)$)

likelihood (points to $\Pr(y|x)$)

prior (points to $\Pr(x)$)

Other example of Estimation in MR

- Image denoising: $H = I$
- Image deblurring: $H =$ convolution mtx in img-space
- Super-resolution: $H =$ diagonal mtx in k-space
- Metabolite quantification in MRSI

What Is the Right Imaging Model?

$$y = Hx + n, \quad n \text{ is Gaussian} \quad (1)$$

$$y = Hx + n, \quad n, x \text{ are Gaussian} \quad (2)$$

MAP Sense

- *MAP Sense = Bayesian (MAP) estimate of (2)*

Intro to Bayesian Estimation

- Bayesian methods maximize the posterior probability:

$$Pr(x|y) \propto Pr(y|x) \cdot Pr(x)$$

- $Pr(y|x)$ (likelihood function) = $\exp(- \|y-Hx\|^2)$

- $Pr(x)$ (prior PDF) = $\exp(-G(x))$

- Gaussian prior:

$$Pr(x) = \exp\{- \frac{1}{2} x^T R_x^{-1} x\}$$

- MAP estimate:

$$x_{est} = \arg \min \|y-Hx\|^2 + G(x)$$

- MAP estimate for Gaussian everything is known as Wiener estimate

Regularization = Bayesian Estimation!

- For any regularization scheme, its almost always possible to formulate the corresponding MAP problem
- MAP = superset of regularization



So why deal with regularization??

Lets talk about Prior Models

- Temporal priors: smooth time-trajectory
- Sparse priors: L0, L1, L2 (=Tikhonov)
- Spatial Priors: most powerful for images
- I recommend robust spatial priors using Markov Fields
- Want priors to be general, not too specific
- Ie, weak rather than strong priors

How to do regularization

- First model physical property of image,
- then create a prior which captures it,
- then formulate MAP estimator,
- Then find a good algorithm to solve it!

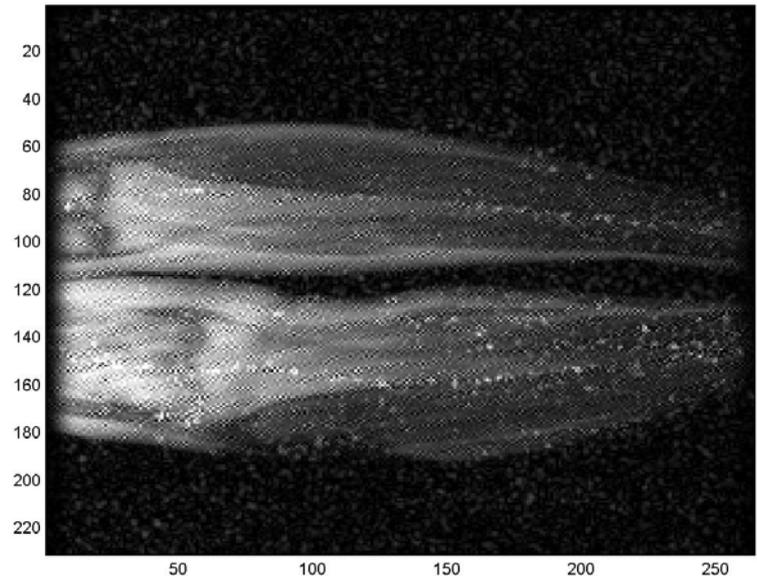
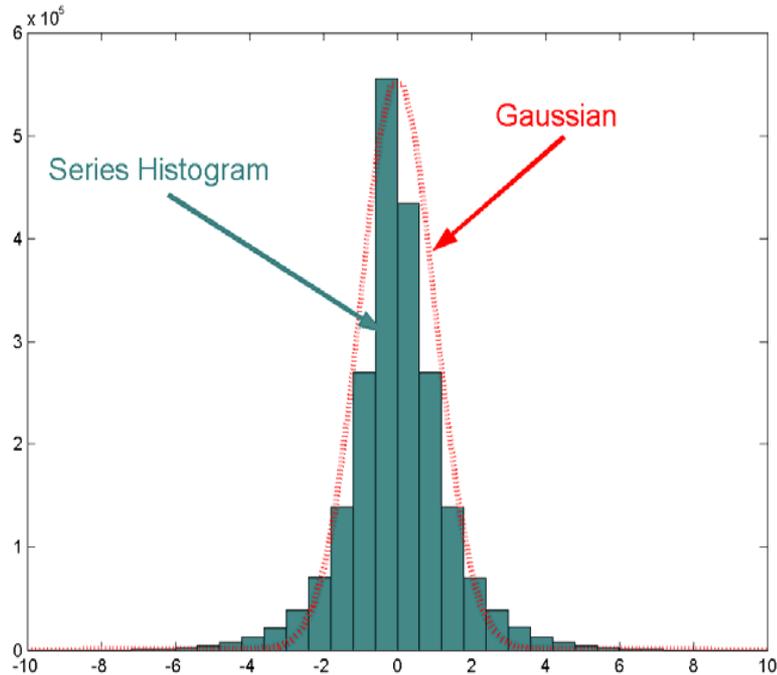
How NOT to do regularization

- DON'T use regularization scheme without bearing on physical property of image!
- Example: L1 or L0 prior in k-space!
- Specifically: deblurring in k-space (handy b/c convolution becomes multiply)
- BUT: hard to impose smoothness priors in k-space → no meaning

Spatial Priors For Images - Example

Frames are tightly distributed around mean

After subtracting mean, images are close to Gaussian



envelope $a(i,j)$

Prior: -mean is μ_x

-local std.dev. varies as $a(i,j)$

Spatial Priors for MR images

- Stochastic MR image model:

$$x(i,j) = \boldsymbol{\mu}_x(i,j) + a(i,j) \cdot (h^{**} p)(i,j) \quad (1)$$

stationary
process

** denotes 2D convolution

$$r(\tau_1, \tau_2) = (h^{**} h)(\tau_1, \tau_2)$$

$\boldsymbol{\mu}_x(i,j)$ is mean image for class

$p(i,j)$ is a unit variance i.i.d. stochastic process

$a(i,j)$ is an envelope function

$h(i,j)$ simulates correlation properties of image x

$$x = ACp + \boldsymbol{\mu} \quad (2)$$

where $A = \text{diag}(a)$, and C is the Toeplitz matrix generated by h

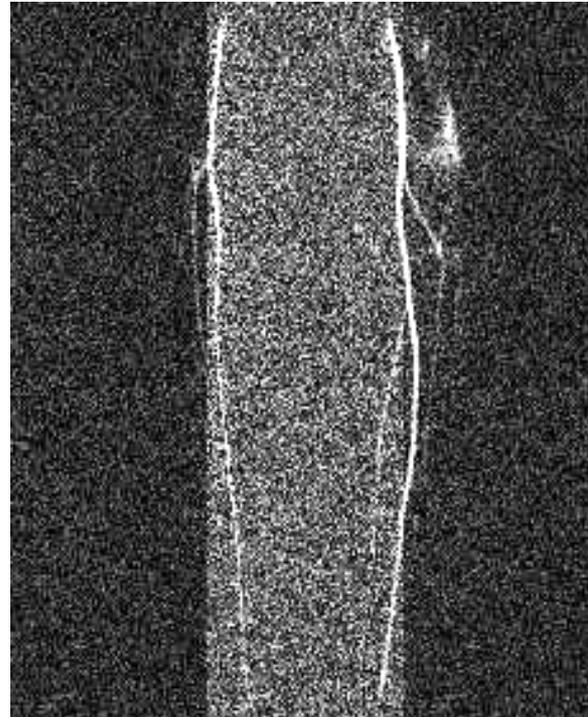
- Can model many important stationary and non-stationary cases

MAP-SENSE Preliminary Results

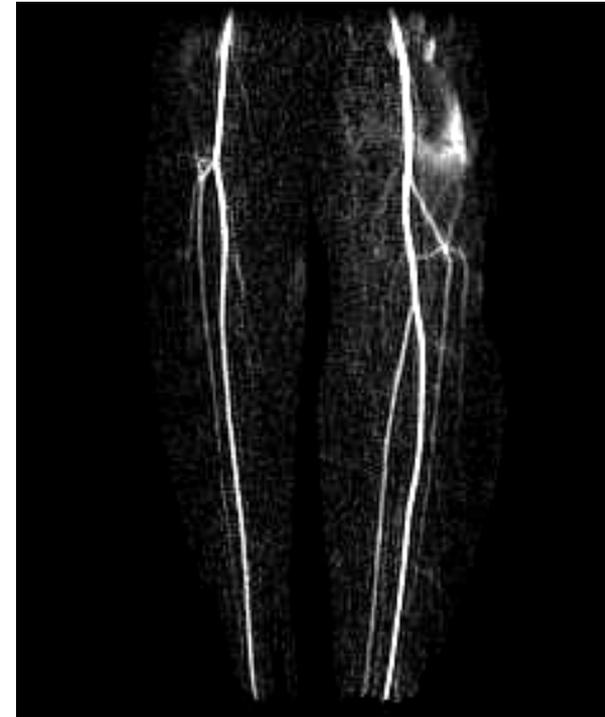
- Scans accelerated 5x
- The angiogram was computed by:
$$\text{avg}(\text{post-contrast}) - \text{avg}(\text{pre-contrast})$$



Unaccelerated

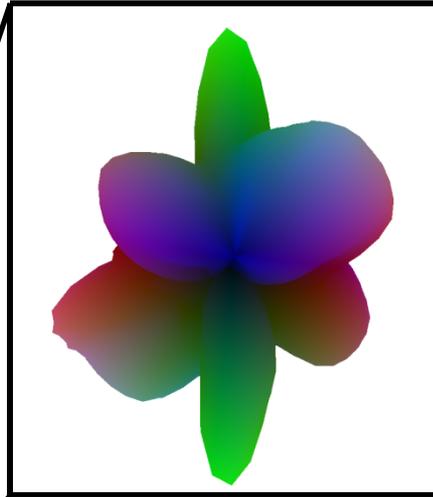
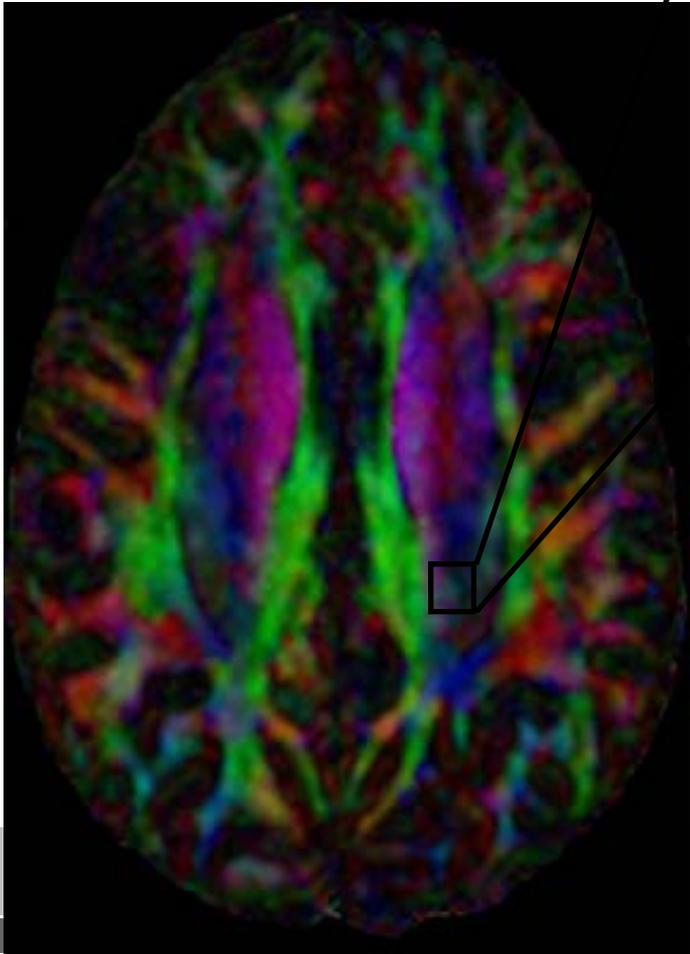


5x faster: SENSE



5x faster: MAP-SENSE

Spatially Constrained High Angular Resolution Diffusion Imaging



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Radiology

Joint work with:
Pratik Mukherjee, MD, PhD
Christopher Hess, MD, PhD
Sri Nagarajan, PhD



University of California
San Francisco

MR Diffusion Imaging

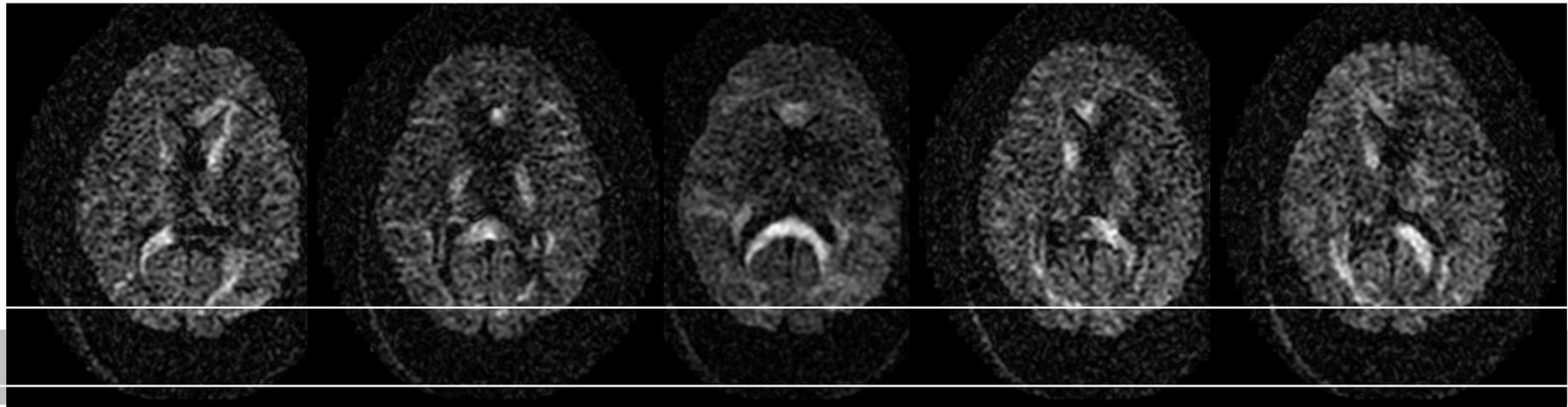
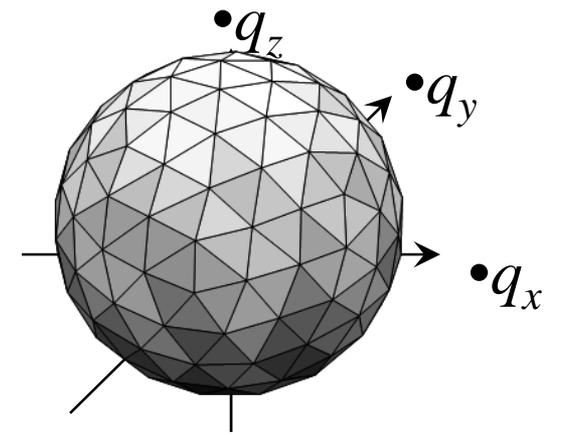
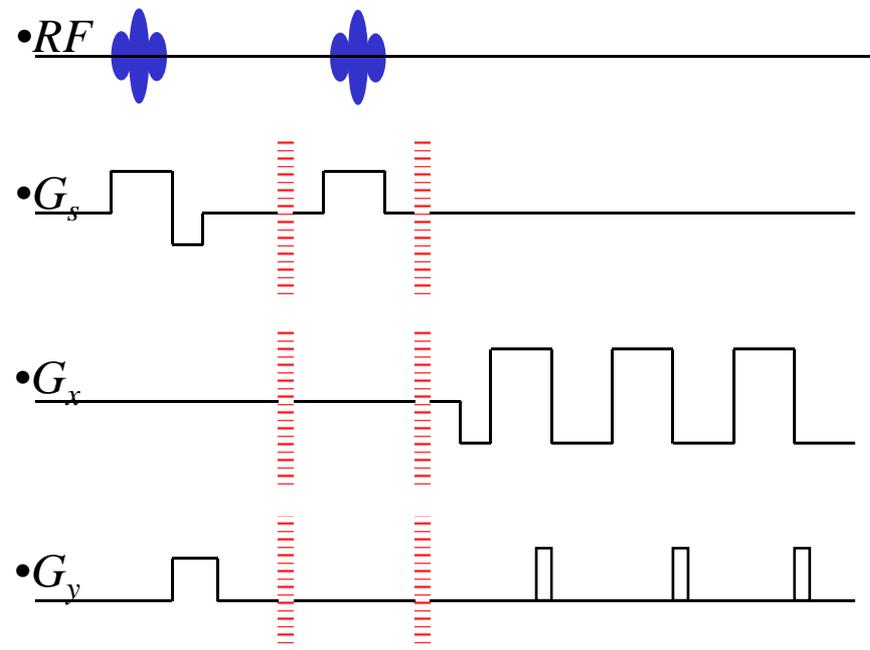
- Diffusion MRI has revolutionized in vivo imaging of brain
- A new contrast mechanism in addition to T1 or T2
- Measures the directionally varying diffusion properties of water in tissue
- Anisotropy of diffusion is an important marker of extant fiber organization
- Enables non-invasive characterization of white matter integrity
- Enables probing of fiber connectivity in the brain, through tractography

Diffusion Tensor Imaging (DTI)

- DTI involves taking 6 directional diffusion imaging measurements
- Then it fits a 3D ellipsoid to these measurements
- Anisotropy of the ellipsoid is correlated with white matter fiber integrity

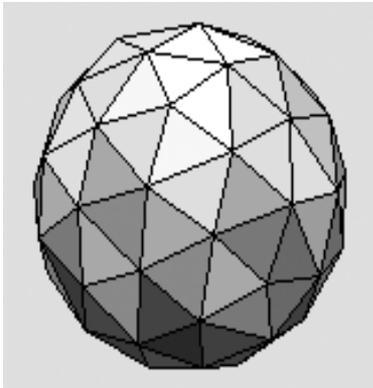
- Cannot resolve crossing fibers
- Fitting an ellipsoid to crossings gives isotropic spheres
 - Erroneously low FA at crossing fibers
 - Messes up tractography, as well as voxel-wise comparisons
- Need much more than 6 directional measurements to resolve crossing fibers

• *Data Acquisition Strategy*

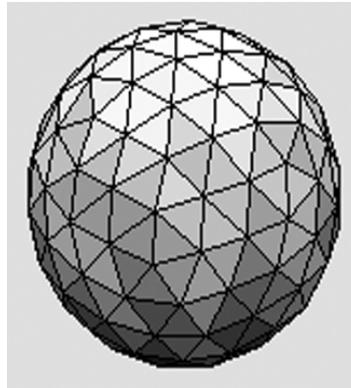


- *High Angular Resolution Diffusion Imaging*
 - *Diffusion-encoding Geometries*

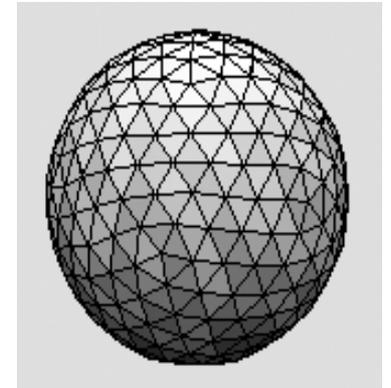
- *55 directions*



- *131 directions*



- *282 directions*



- *Gradient directions are determined using an “electrostatic repulsion” model,*
 - *for the most uniform sampling of 3D space:*
 - *<http://www.research.att.com/~njas/electrons/>*

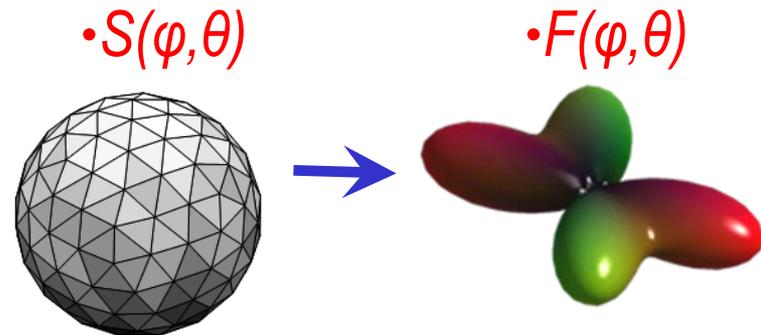
Reconstruction Problem

•Goal:

• *Construct a spherical function that characterizes the angular structure of diffusion anisotropy in each voxel.*

•Solutions:

- *Multi-tensor fitting*
- *Generalized DTI*
- *Persistent angular structure*
- *Spherical encoding*
- *Spherical harmonic “ADC profile”*
- *Circular spectrum mapping*
- *Spherical deconvolution*
- *q-ball imaging*
- *Harmonic q-Ball*



Tuch et al, MRM 2002

Özarslan et al, MRM 2003

Jansons et al, Inv. Prob. 2003

Lin et al, ISMRM 2003

Frank, MRM 2002; Alexander DC et al, MRM 2002

Zhan et al, Neuroimage 2004

Tournier et al, Neuroimage 2004

Tuch et al, Neuron 2003; Tuch MRM 2004

Hess et al, ISMRM 2005; MRM 2006

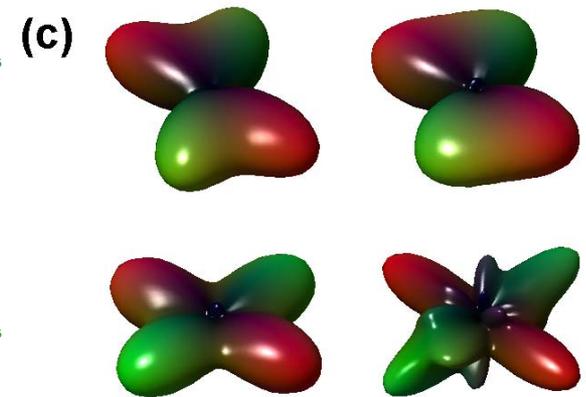
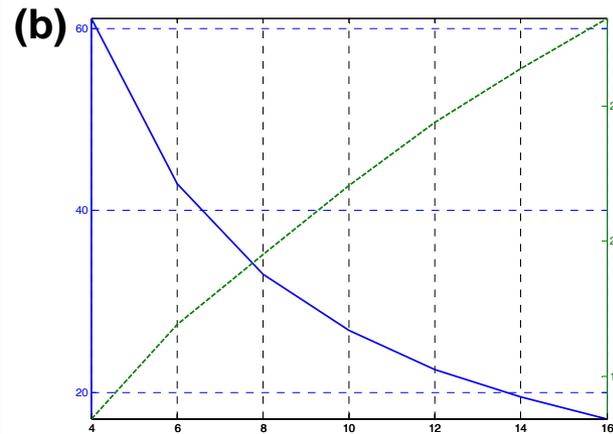
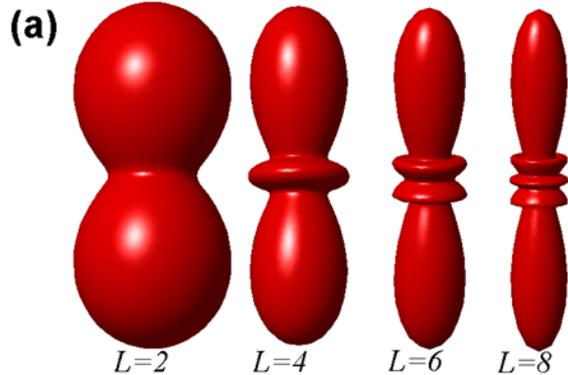
• High Angular Resolution Diffusion Imaging: Spherical Harmonic Q-ball

• Point Spread Functions

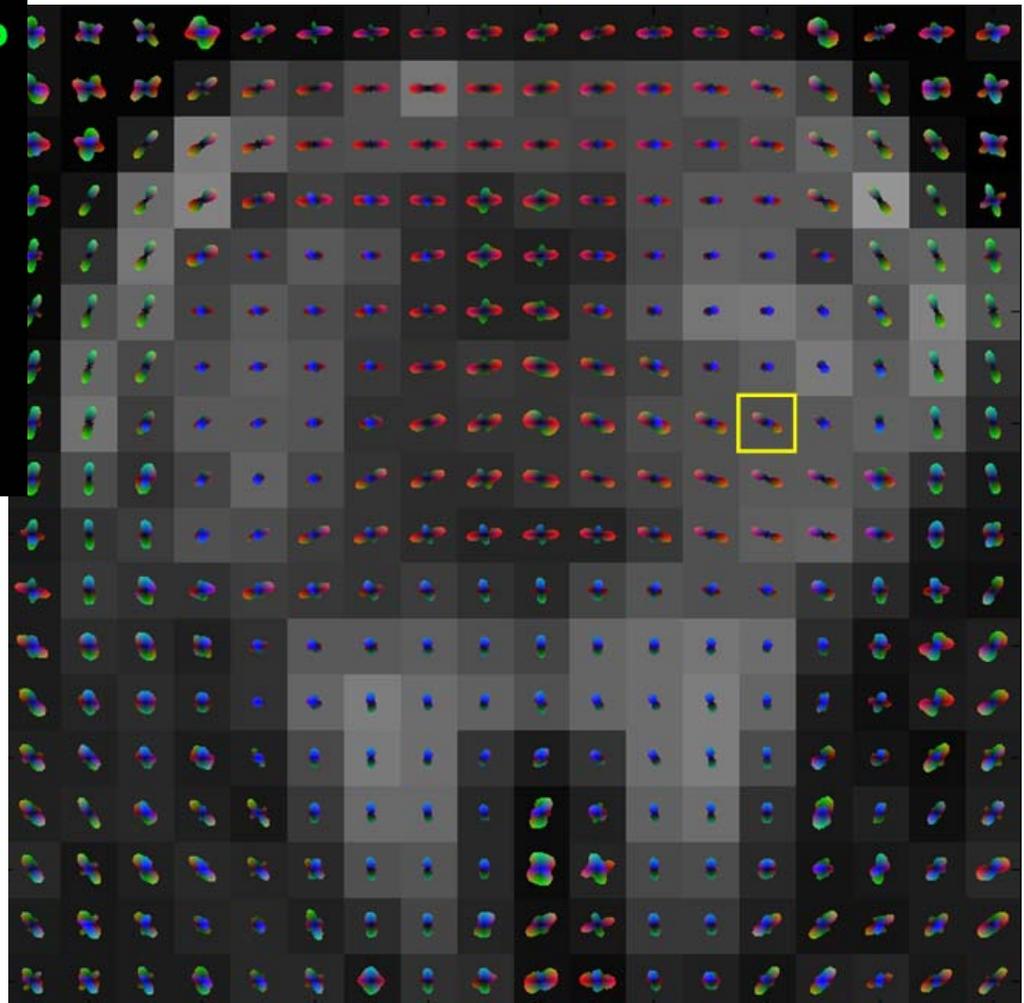
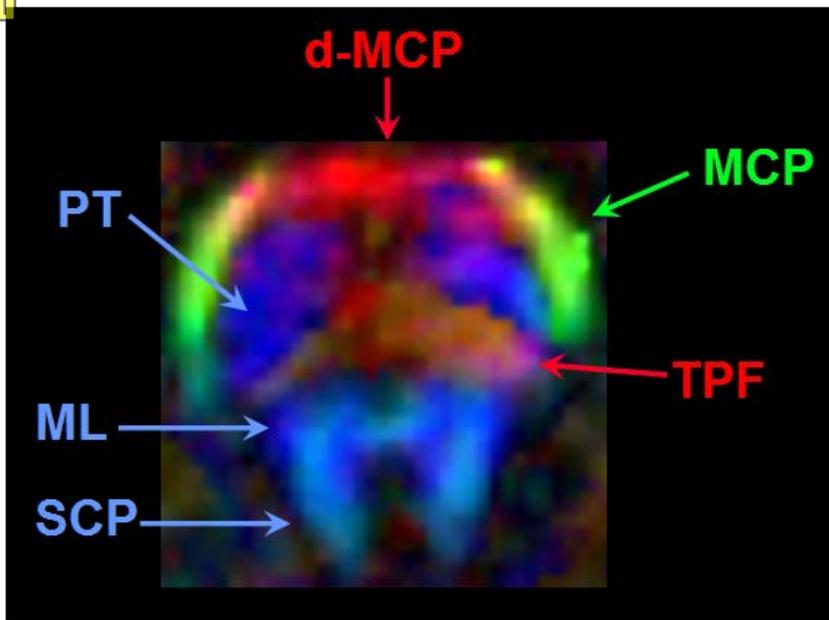
• Model Order vs
• Angular Resolution

• Simulated
• ODFs

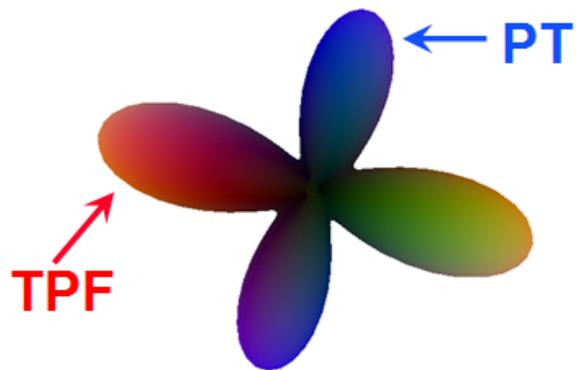
Figure 1



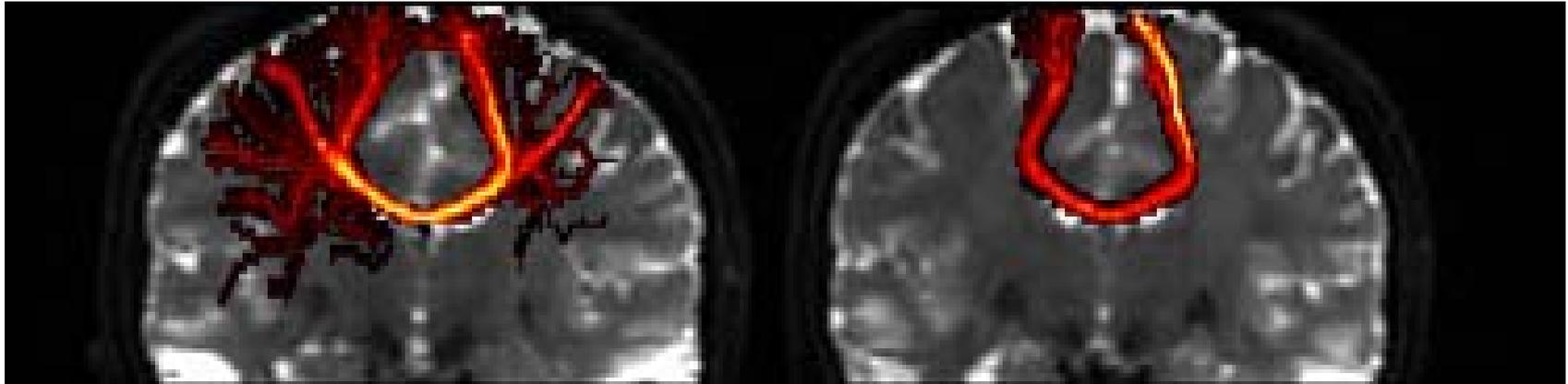
- more computationally efficient
 - more numerically stable
 - more analytically tractable



- Middle cerebellar peduncle (MCP)
- Superior cerebellar peduncle (SCP)
- Pyramidal tract (PT)
- Trans pontocerebellar fibers (TPF)



Clinically Feasible HARDI Tractography



Harmonic q -ball

DTI

- Bootstrapping to generate probability distribution function for orientations
- Probabilistic streamline tracking
- 55 direction HARDI protocol, 1x1x2 mm resolution at $b=3000$ s/mm²

Problems

- ODF reconstruction suffers from noise
- Matrix is ill-conditioned
 - i.e. its inverse “magnifies” small noise values into large ones
- There are not enough diffusion directions
- HARDI with high $b \rightarrow$ very low SNR (< 20)

Current Solutions

- Use efficient representation of ODFs
 - Spherical harmonic basis
 - Radial basis
- Limit the order of the basis to use as few basis functions as possible
 - Currently we use spherical harmonics only up to order 4 or 6.
 - Higher order harmonics contain mostly noise
- (but this limits the angular resolution achievable, thus negating the motivation for HARDI)

Linear System

- Capture the model (whether RBF, SH, ...) into the linear model (matrix) H

$$y = Hx + n$$

Then solve for x by inverting H

Usually inversion of H is ill-posed, so add a regularization term

This process is the same for ALL linear estimation problems!

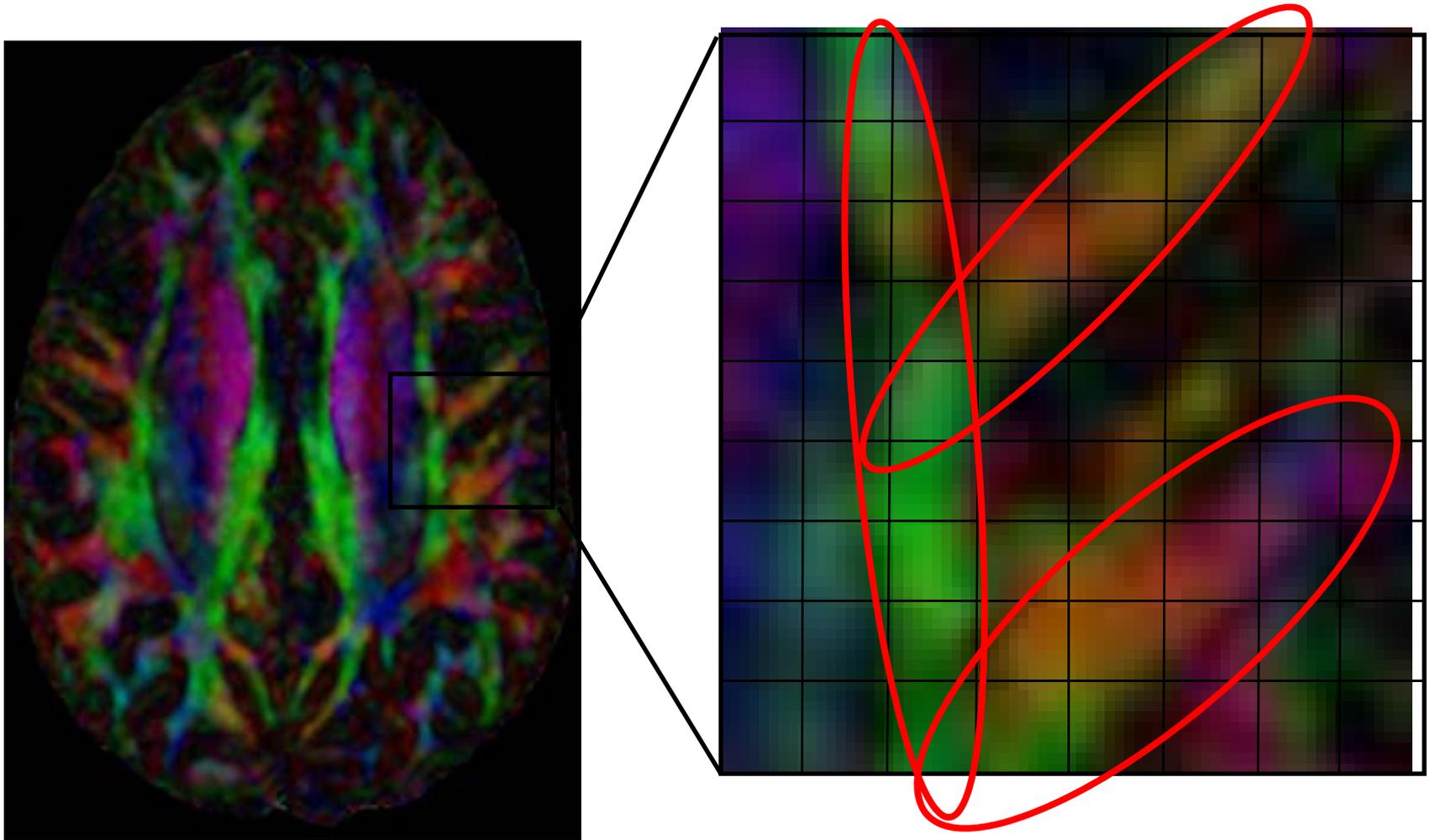
Regularization

- Regularization of matrix inverse
 - Tikhonov
 - Laplace-Beltrami
- Tikhonov Regularization penalizes all harmonic coefficients
- Laplace-Beltrami penalizes higher harmonic coefficients more
- Both methods serve to limit the effective angular resolution of reconstructed ODFs

A New Approach: add spatial constraints

- Fibers are not arbitrarily arranged in space
 - Organized structure – follow coherent fiber tracts
 - ODFs also have this organized structure
 - ODF at one voxel is therefore related to ODF at its neighbours
-
1. How to characterize this neighbourhood relationship?
 2. How to exploit these spatial constraints to improve ODF reconstruction?

Adding Spatial Constraints



- Neighbours are “like” each other, likely to have similar ODFs
- But need to allow for discontinuous boundaries

Iterative Algorithm

Begin with $\boldsymbol{\eta} = \boldsymbol{\eta}_0$. Then for $k = 1$ to K , repeat :

$$\hat{\boldsymbol{\eta}}^k = \arg \min_{\boldsymbol{\eta}} \left\{ \|\mathbf{e} - \bar{\mathbf{P}}\bar{\mathbf{Z}}_o \boldsymbol{\eta}\|^2 + \lambda^2 \|\boldsymbol{\eta}\|^2 + \mu^2 \|\mathbf{D}\mathbf{W}(\boldsymbol{\eta}^{k-1}) \boldsymbol{\eta}\|^2 \right\}$$

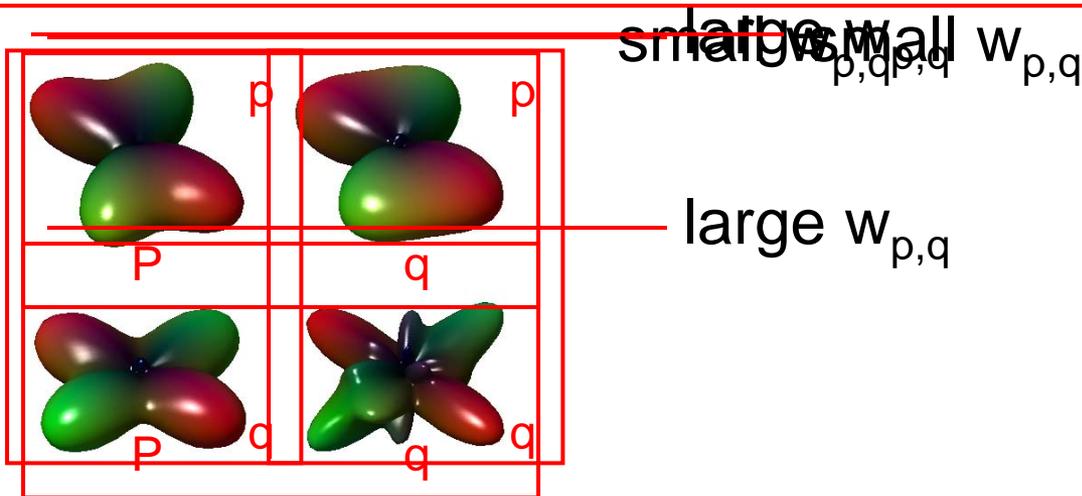
where $\mathbf{W}(\boldsymbol{\eta}) = [w_{p,q}]$, $w_{p,q} = 1 - \frac{\|\boldsymbol{\eta}_p - \boldsymbol{\eta}_q\|}{\|\boldsymbol{\eta}_p\| \|\boldsymbol{\eta}_q\|}$

Iterative Algorithm

Begin with $\boldsymbol{\eta} = \boldsymbol{\eta}_0$. Then for $k = 1$ to K , repeat :

$$\hat{\boldsymbol{\eta}}^k = \arg \min_{\boldsymbol{\eta}} \left\{ \|\mathbf{e} - \bar{\mathbf{P}}\bar{\mathbf{Z}}_Q \boldsymbol{\eta}\|^2 + \lambda^2 \|\boldsymbol{\eta}\|^2 + \mu^2 \|\mathbf{DW}(\boldsymbol{\eta}^{k-1}) \boldsymbol{\eta}\|^2 \right\}$$

where $\mathbf{W}(\boldsymbol{\eta}) = [w_{p,q}]$, $w_{p,q} = 1 - \frac{\|\boldsymbol{\eta}_p - \boldsymbol{\eta}_q\|}{\|\boldsymbol{\eta}_p\| \|\boldsymbol{\eta}_q\|}$



Results – simulation



Results – simulation



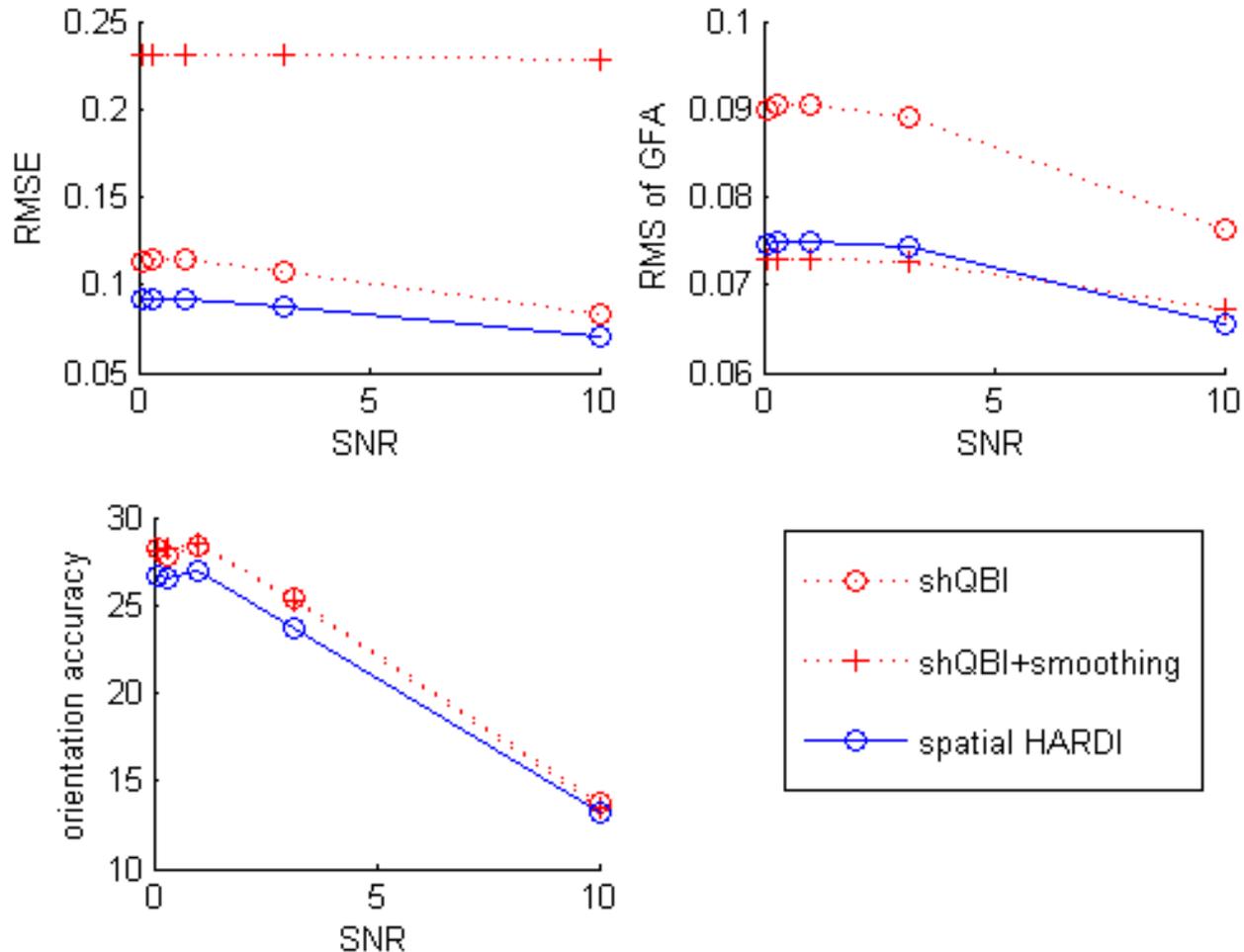
Results – simulation



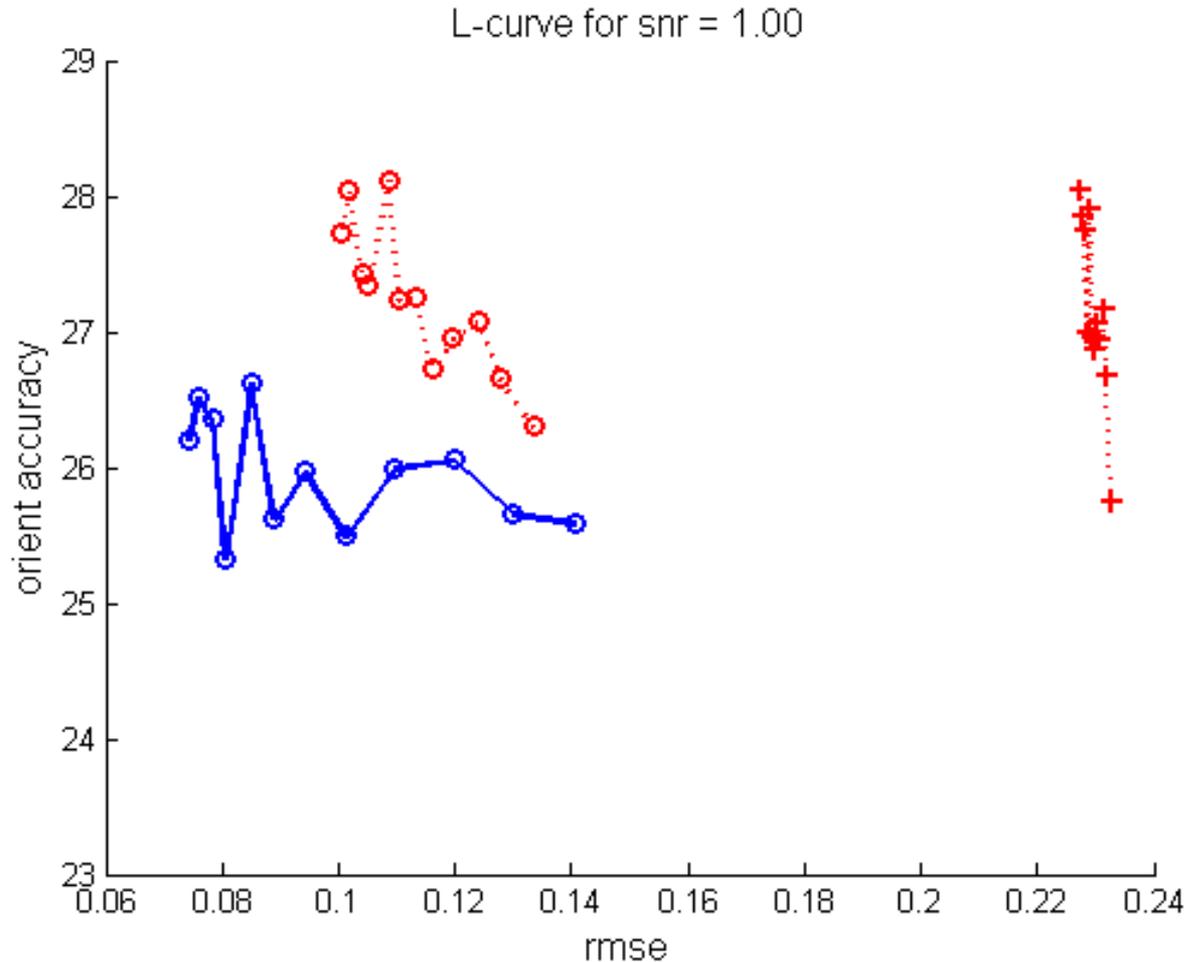
Monte Carlo simulations

- Repeated multiple times for multiple, random 3D tracts within a 15 x 15 x 15 voxel volume
- Repeated for varying :
 - SNR
 - Algorithm parameters (λ , μ)
- Evaluation criteria:
 - RMSE
 - Generalized FA
 - Orientation accuracy

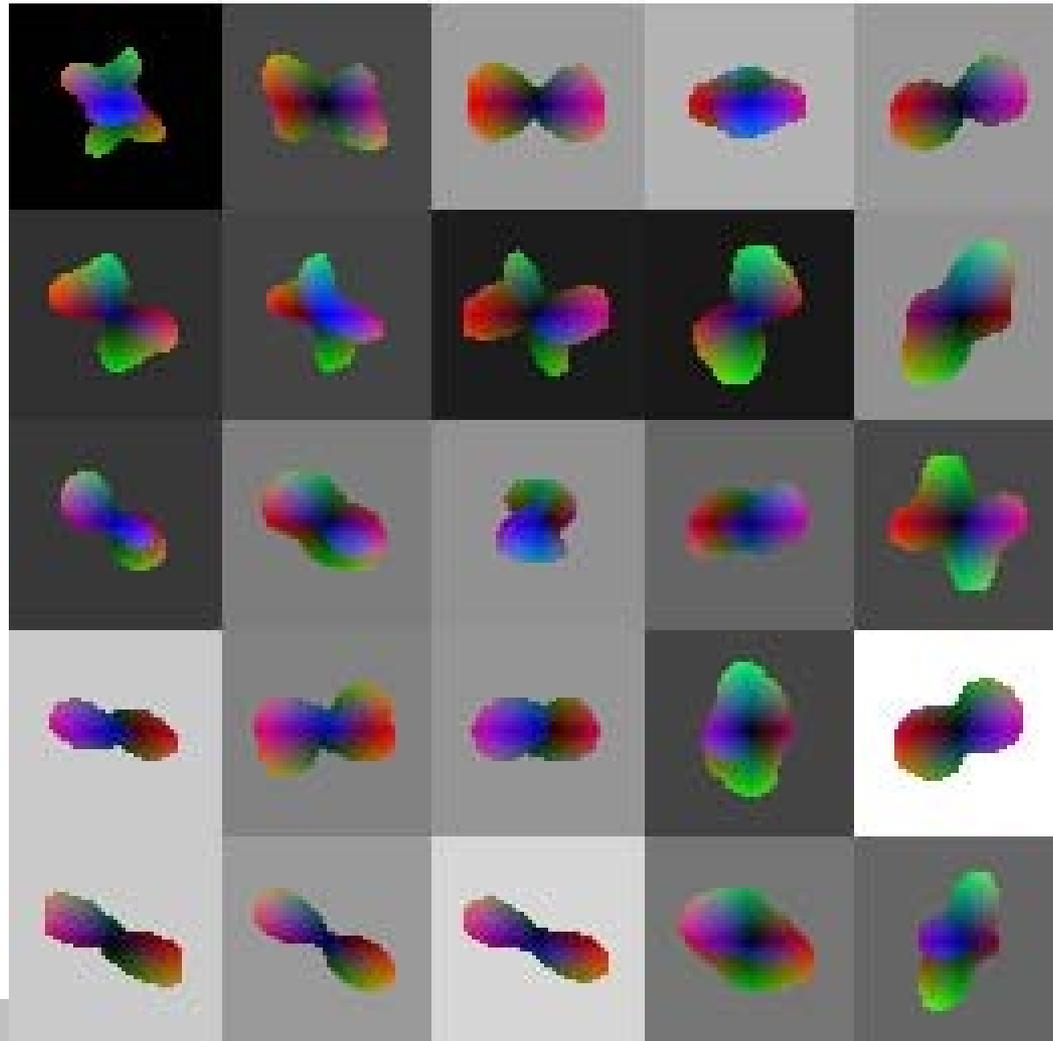
Monte Carlo simulations



Monte Carlo simulations



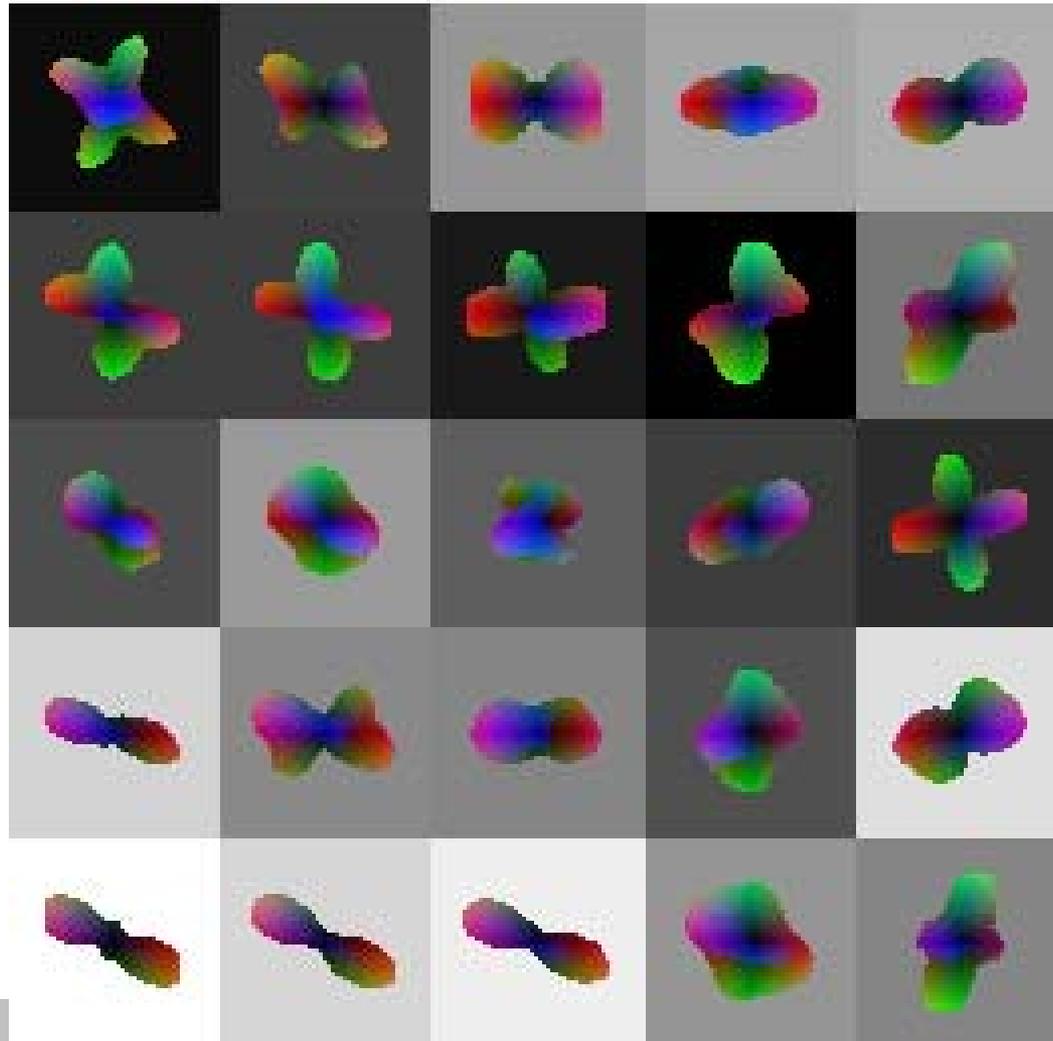
In vivo results



IDEA Lab, Radiology, Cornell

SIT QBI

In vivo results



Part IV : Detection Theory and Examples

- Introduction to optimal detection
- Matched filter detectors
- Detection examples from MR

What is Detection

- Deciding whether, and when, an event occurs
- a.k.a. Decision Theory, Hypothesis testing
- Presence/absence of
 - signal
 - activation (fMRI)
 - foreground/background
 - tissue – WM/GM/CSF (segmentation)
- Measures whether statistically significant change has occurred or not

Detection

- “Spot the Money”



Hypothesis Testing with Matched Filter

- Let the signal be $y(t)$, model be $h(t)$

Hypothesis testing:

$$H_0: y(t) = n(t) \quad (\text{no signal})$$

$$H_1: y(t) = h(t) + n(t) \quad (\text{signal})$$

- The optimal decision is given by the Likelihood ratio test (Nieman-Pearson Theorem)

$$\text{Select } H_1 \text{ if } L(y) = \Pr(y|H_1)/\Pr(y|H_0) > \gamma$$

- It can be shown (Kay 01) to be equivalent to

$$y(t) * h(t) > \gamma'$$

Matched Filter

Matched Filters

- If the profile of a certain signal is known, it can be detected using the Matched Filter
- If the question is not IF but WHERE...
- Maximum of MF output denotes the most likely location of the object $h(t)$

Matched Filters

- Example 1: activation in fMRI
 - Need profile model: hemodynamic response function
- Example 2: Detecting malignant tumours in mammograms
 - need profile model: temporal response to contrast agent
- Example 3: Edge detection
- Example 4: detecting contrast arrival in CE-MRA
- In each case need a model to “match” the signal

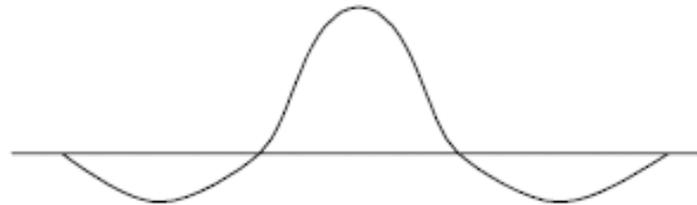
Edge Detection

- Edge information can be used for segmentation
- Detect edges by finding areas of max intensity change
- DoG (Derivative of Gaussian):
$$\nabla^2 (I(x,y) * G(x,y,\sigma))$$
- $G(x,y,\sigma)$ = Gaussian
- ∇^2 = Laplacian operator
- Marr-Hildreth, Canny, Roberts, etc
- Problems: very sensitive to noise, choice of σ

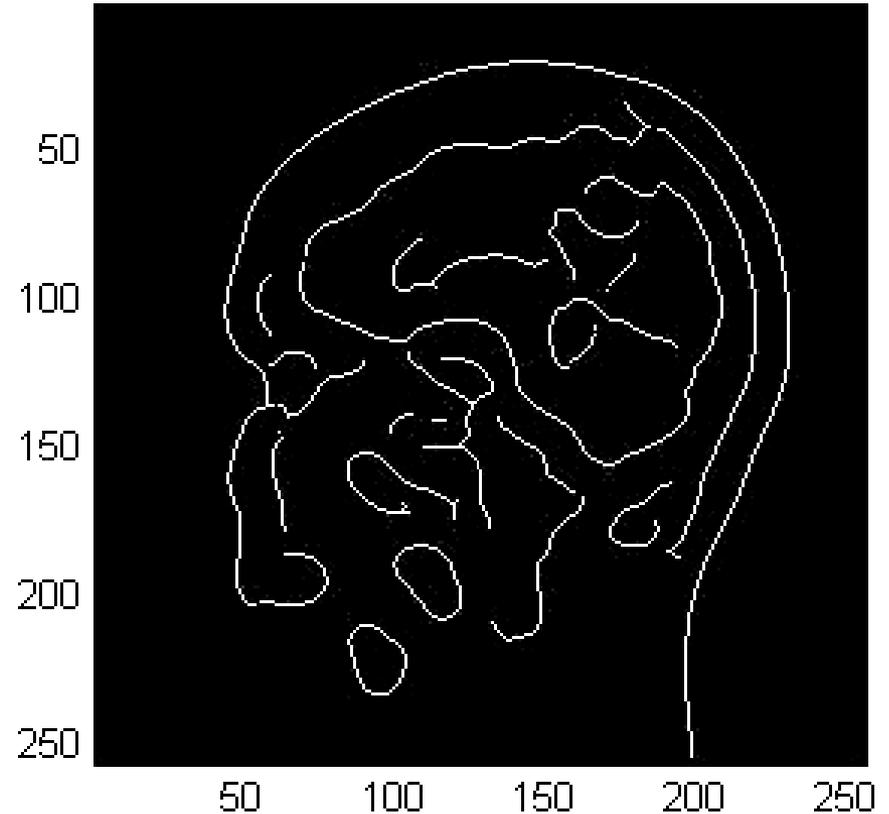
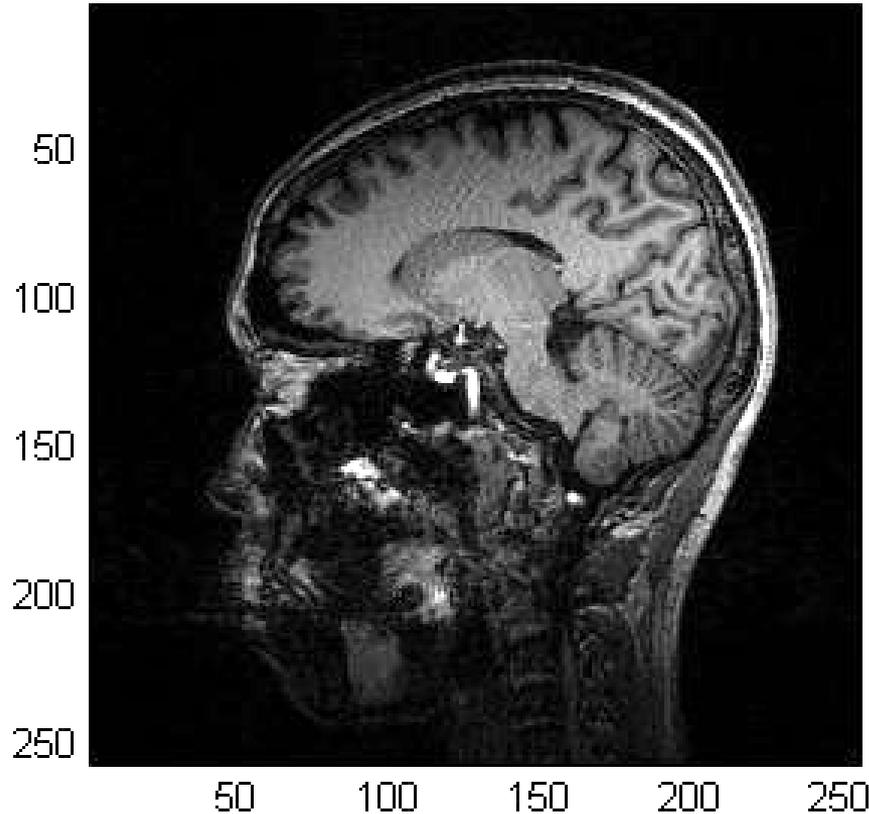
Edge Detection

	1	
1	-4	1
	1	

$$\nabla^2(G_\sigma \star I).$$

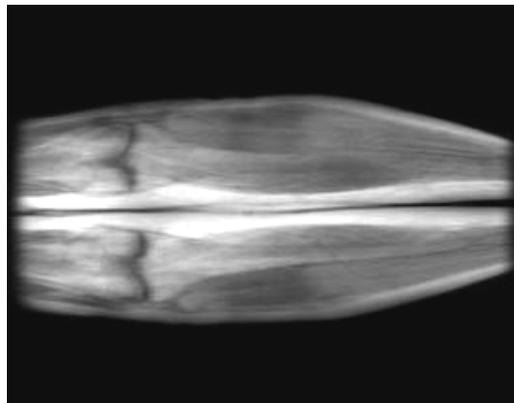


Edge Detection example using MATLAB

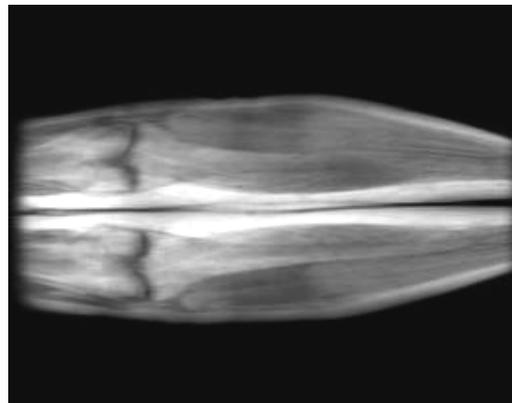


bw = edge(I, 'canny', sigma);

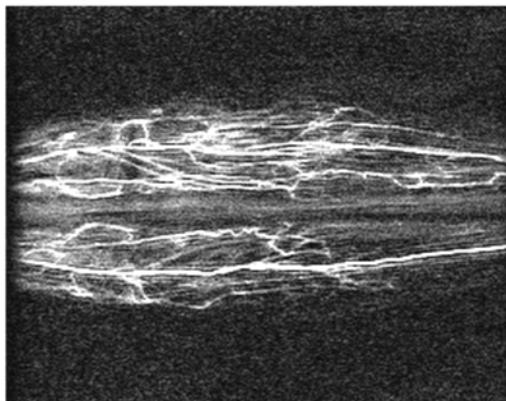
Example: Contrast Arrival in CE-MRA



-



=

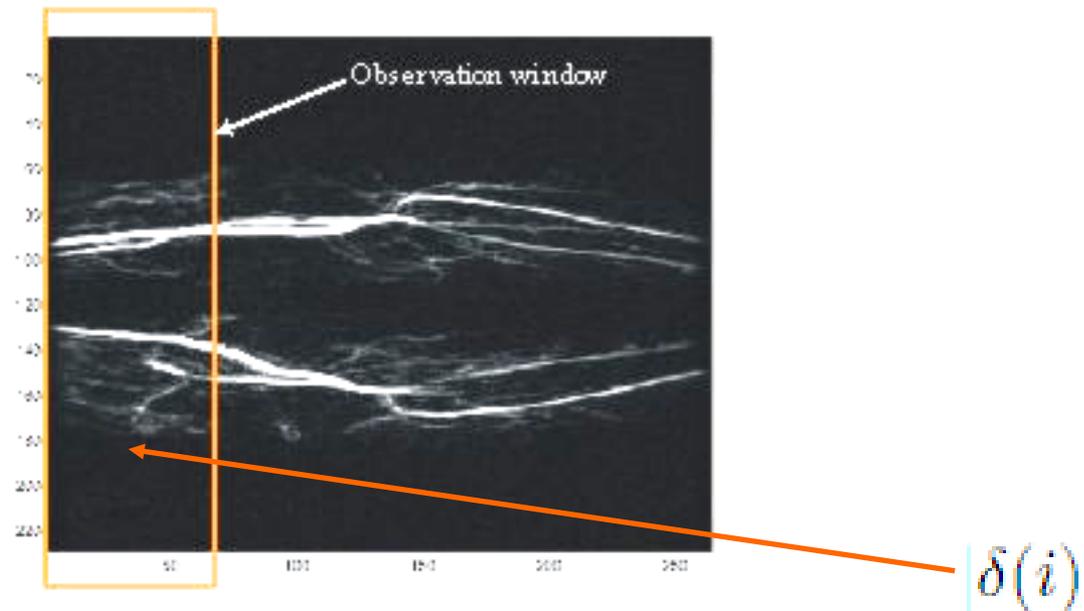


Mask subtraction in MRA gives vasculature

Automatic Detection of Contrast Arrival

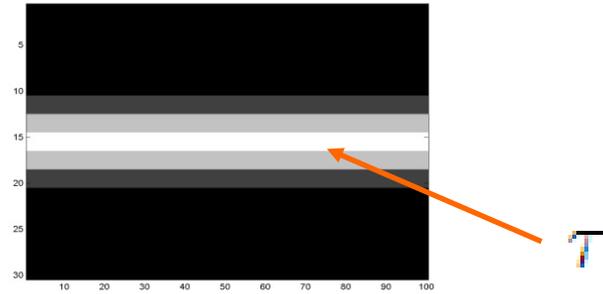
- MRDSA relies on good estimate of contrast arrival
- Completely unsupervised, reliable automatic method
- >90% accuracy, c.f. earlier reported method (~60% accuracy)
 - matched filter - spatial metric
 - keyhole - frequency metric

Vasculature strongly oriented horizontally



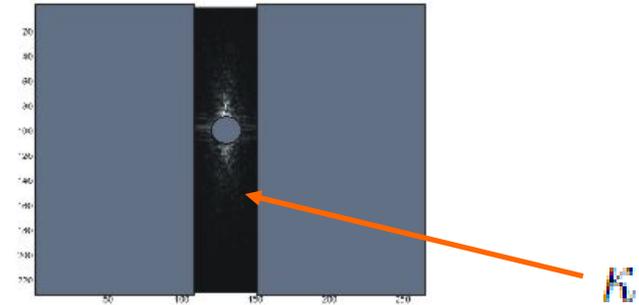


Matched Filter



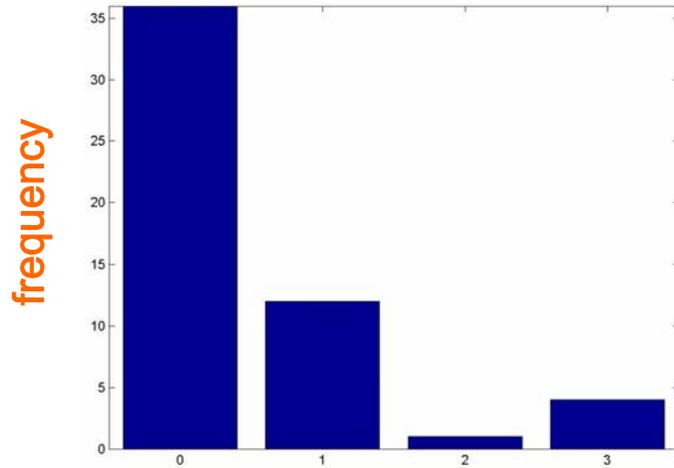
$$\lambda_{MF}(i) = \frac{\|\delta(i) * \tau\|}{\|\tau\| \cdot \|\delta(i)\|},$$

Keyhole



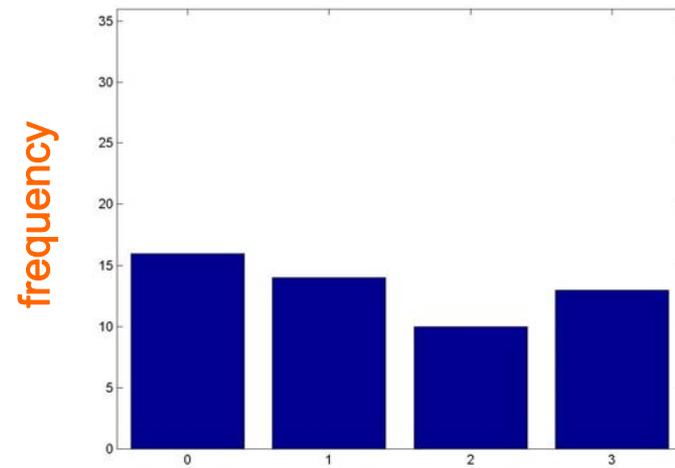
$$\lambda_{KH}(i) = \frac{\|\delta_K(i) \cdot \kappa\|}{\|\delta_K(i)\|}.$$

Results : Our method



accurate -----> inaccurate

Earlier method



accurate -----> inaccurate

Estimation and Detection of MR Signals

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Visit:

<http://www.cs.cornell.edu/~rdz/SENSE.htm>

This concludes today's lecture.

Next week: Classification, Image segmentation, Registration

References

- Simon Kay. Statistical Signal Processing. Part I: Estimation Theory. Prentice Hall 2002
- Simon Kay. Statistical Signal Processing. Part II: Detection Theory. Prentice Hall 2002
- Haacke et al. Fundamentals of MRI.
- Zhi-Pei Liang and Paul Lauterbur. Principles of MRI – A Signal Processing Perspective.

Info on part IV:

- Ashish Raj. Improvements in MRI Using Information Redundancy. PhD thesis, Cornell University, May 2005.
- Website: <http://www.cs.cornell.edu/~rdz/SENSE.htm>