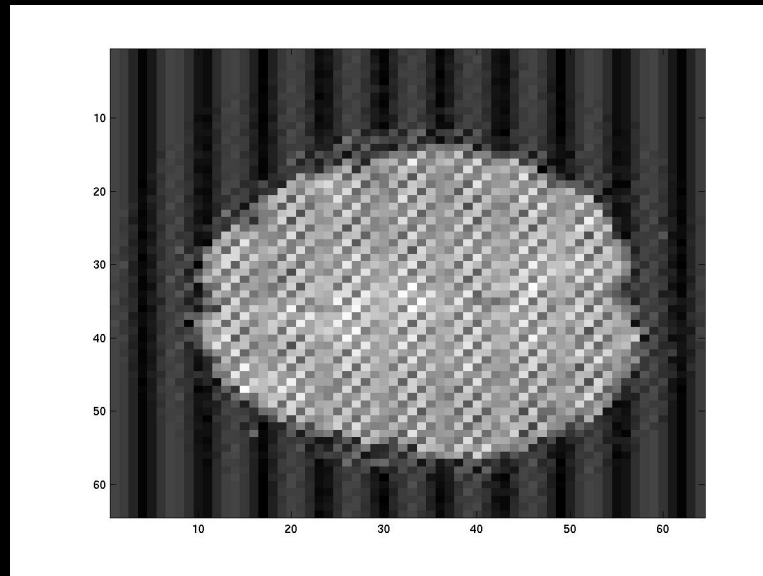


White Pixel Artifact

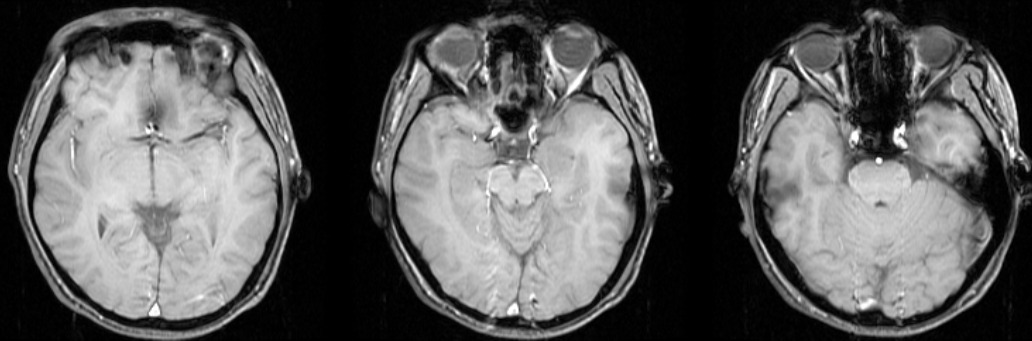
- Caused by a noise spike during acquisition
- Spike in K-space \leftrightarrow sinusoid in image space



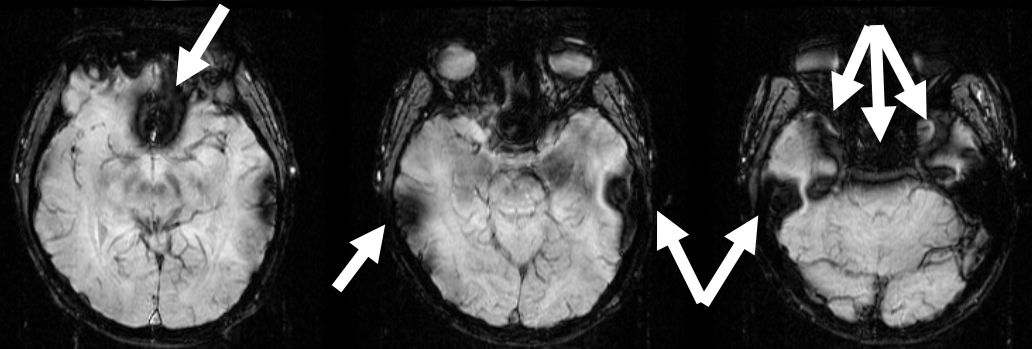
Susceptibility Artifacts

- Off-resonance artifacts caused by adjacent regions with different Susceptibility
- **BOLD** signal requires susceptibility weighting... but this also leads to image artifacts

No
Susceptibility
Contrast

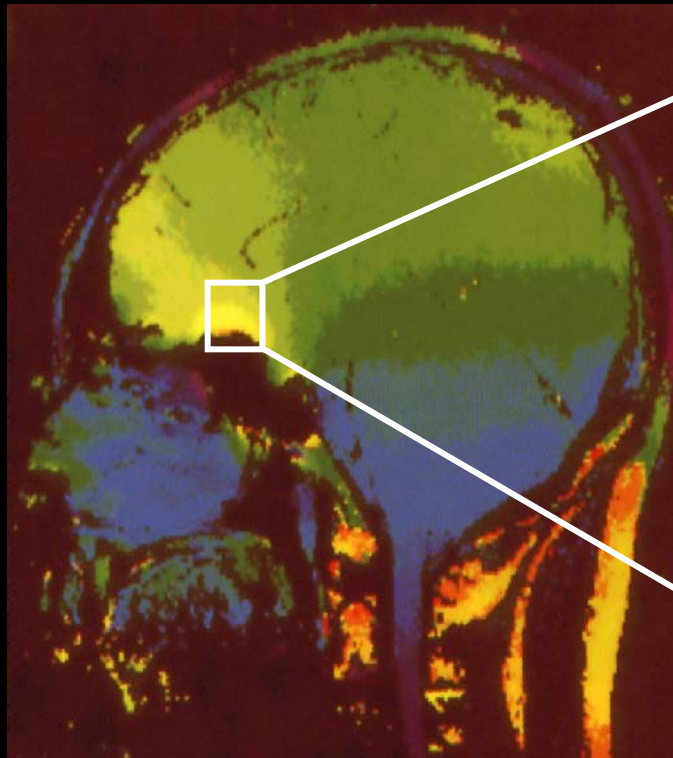


High
Susceptibility
Contrast



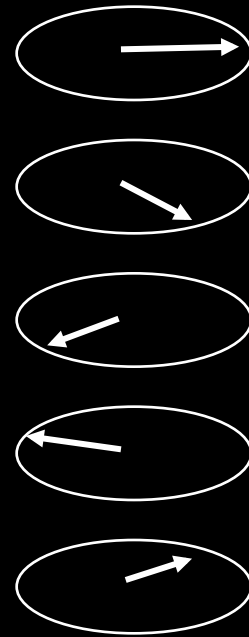
Through-Plane Dephasing

Magnetic Fields
in the Head



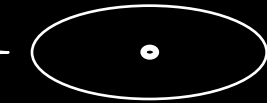
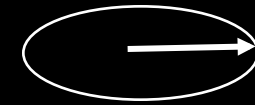
Li et al. Magn. Reson. Med. 36:710
(1996)

Low Field



High Field

Ideal

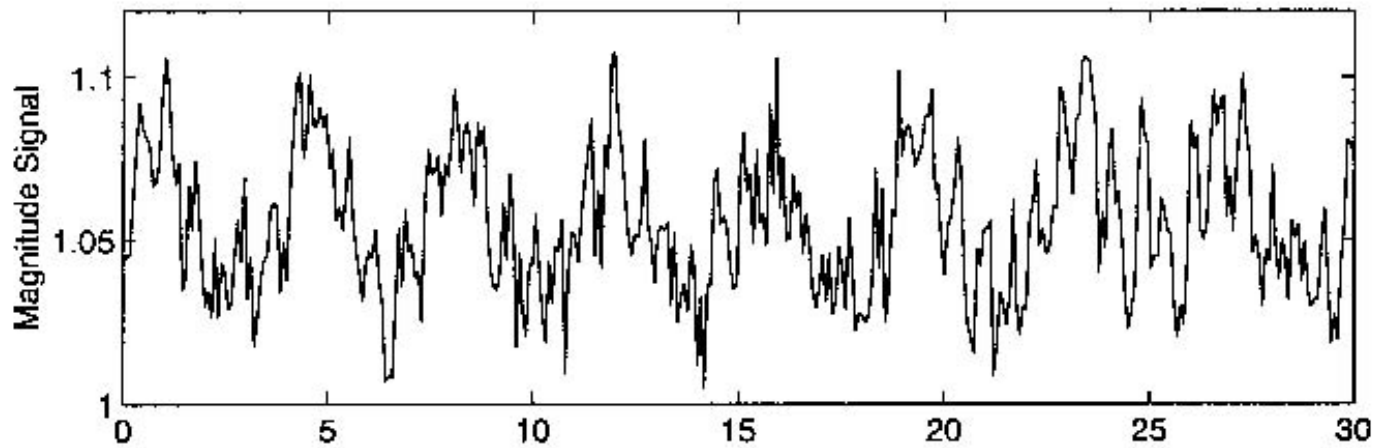


Signal Loss

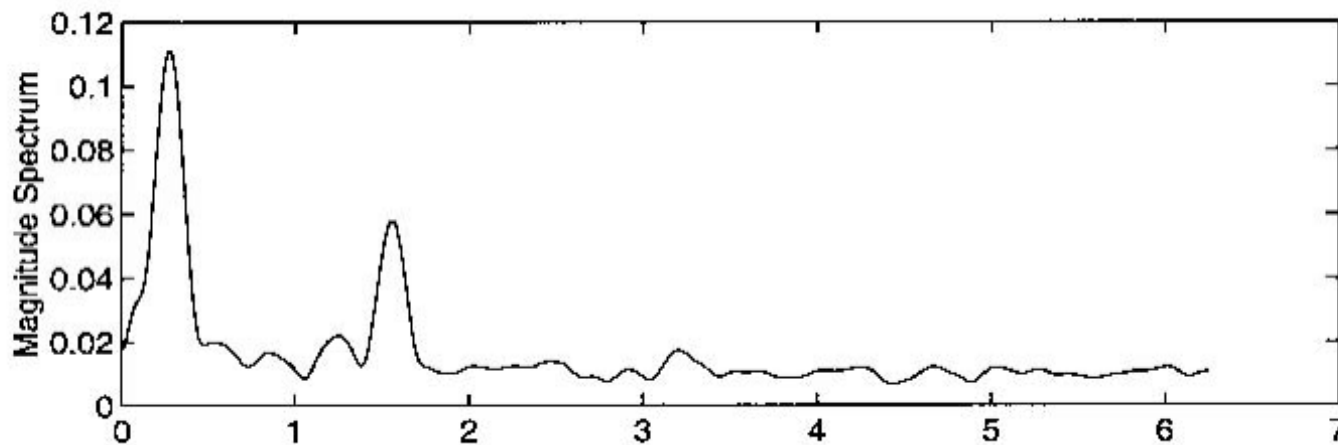
Susceptibility Artifacts

- Local gradients cause:
 - extra dephasing when the gradient occurs through the imaging plane (destructive interference). Result is signal loss.
 - distortion (skewing of the k-space trajectory in different voxels) when the gradient happens in the plane. Signal loss and distortions of the image.
- Solution: this is an active research field, lots of tricks you can do, but they all have an associated cost in time, SNR, computation, hardware..
 - Simplest: design acquisition parameters such that the artifacts are minimized.
 - Z-shimming: apply set of additional gradients
 - Active shims: create additional gradient using materials, coils
 - iterative reconstructions: crunch the numbers

Physiological oscillations



Time
domain



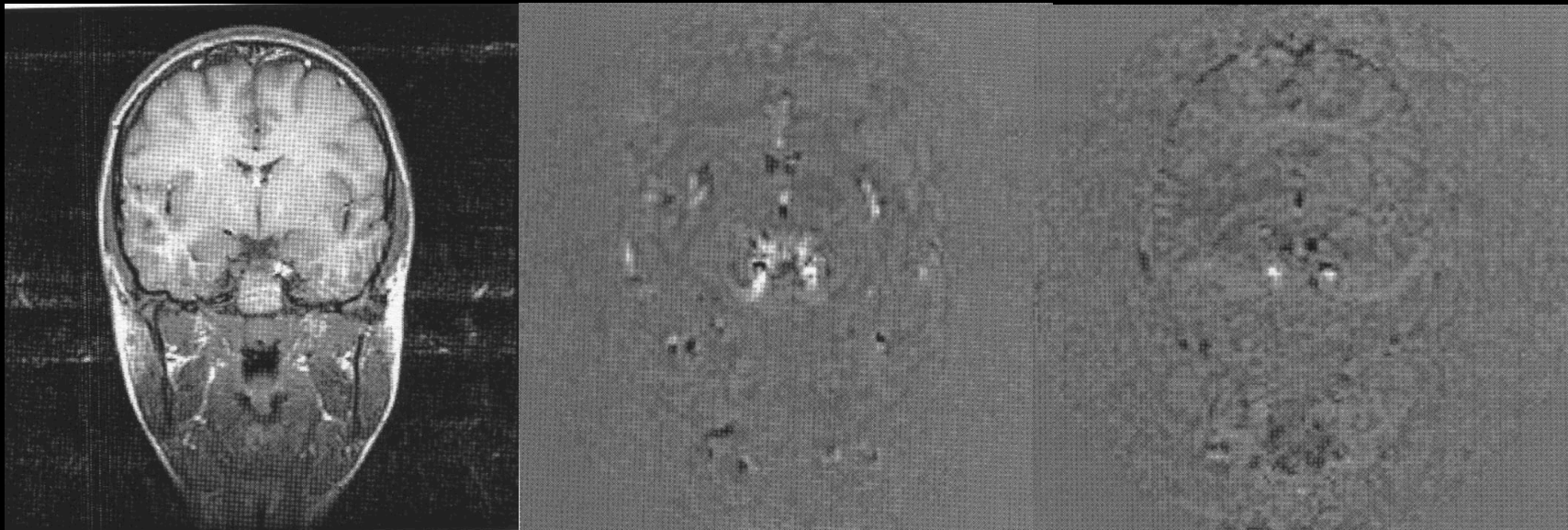
Frequency
domain

Cardiac and Respiratory Variance

anatomy

Residual Variance w/o
Physio correction

Residual Variance w/
Physio correction



Cardiac Noise

- Blood flow is pulsatile -> changes blood volume, and velocity.
- Other blood flow effects on MR signal:
 - Flow enhancement (incoming spins have not received any RF , fully relaxed -> more signal)
 - Flow void (sometimes spins flow so fast through the plane that they don't see the RF pulse, or they flow out before they can be encoded -> less signal)
 - Flow induced displacement (additional phase acquired because of in-plane movement -> distorted/displaced signal, ghosting)

Reduction of cardiac effects during Acquisition

- Use a smaller flip angle - reduces flow enhancements and voids.
- Use flow “spoilers” to remove vascular signals. (pair of symmetric gradient pulses, a.k.a. “crushers”, refocus the signal from stationary spins but not from moving spins.)
- Use fast acquisition (single shot) to reduce ghosting.
- “Cardiac Gating”

Reduction of cardiac artifacts after acquisition

- Digital Filters ...
- Measure cardiac waveform and include in analysis as a confound.
- Note: watch out for aliasing!!
 - heartbeat ~from 0.5-2 Hz, typically ~1 Hz
 - typical Nyquist frequency < 0.5 Hz

Respiration

- Air and Tissue difference in χ :
Distortion of B_0 field

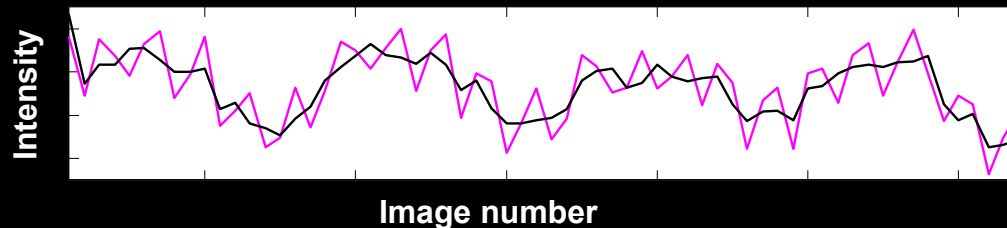


Phase difference between inspiration and expiration for a coronal slice.

- Chest movement changes the shape of the B_0 field. Changes gradients too.
- Resonant frequency changes slightly (Recall that $\omega_0 = \gamma B_0$)
- Blood Pressure changes slightly with respiration (pulsation of arteries and hence blood volume)

Corrections for Respiration

- Fast image acquisition (single shot)
- “Notch” or “band-stop” Filters
- Record Respiratory waveform and use as a confound.
(Note- sometimes it’s correlated with task of interest)



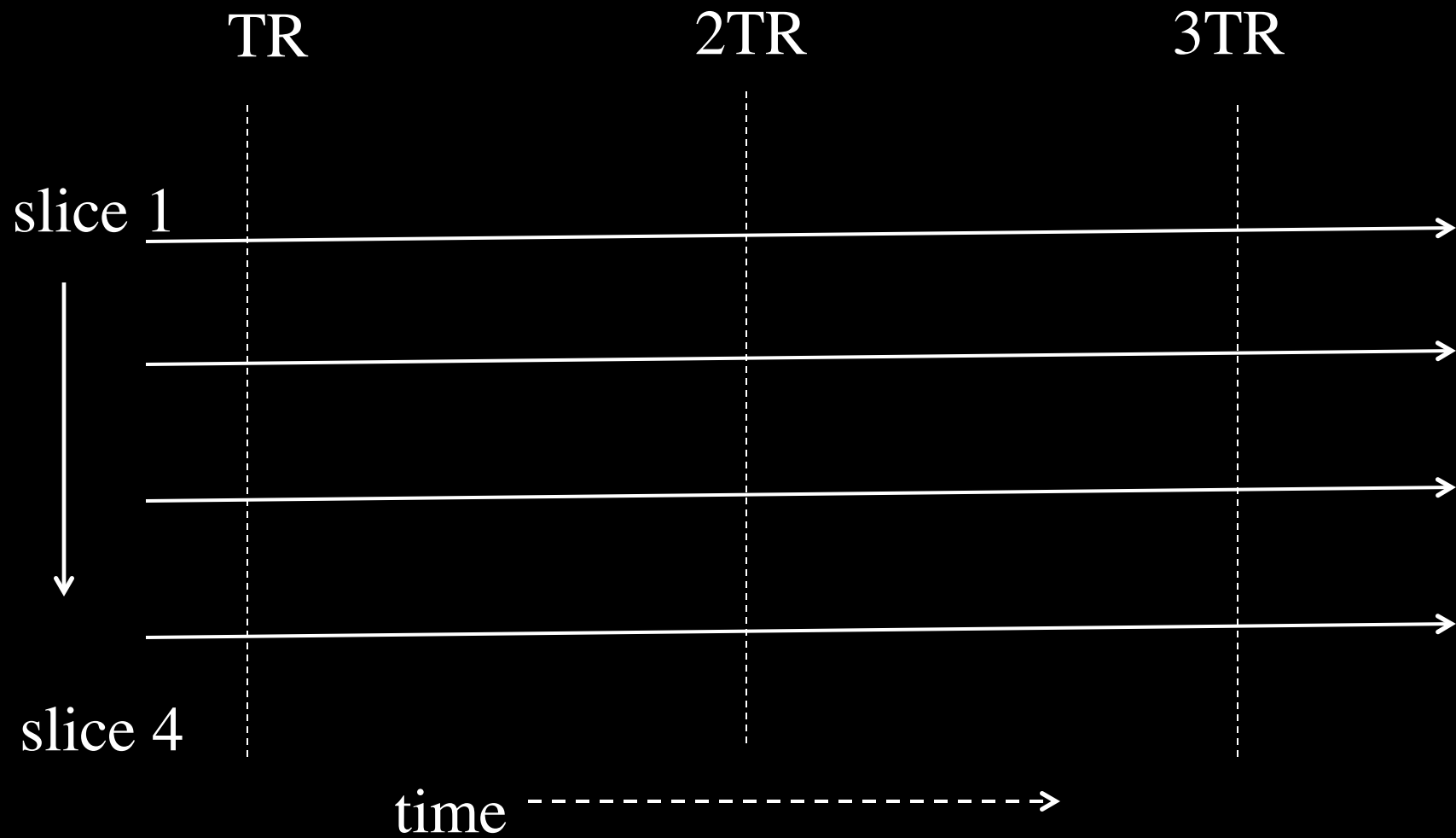
Timecourse before (purple) and after (black) regression correction.

- Aliasing is not as much of a problem as in cardiac fluctuations, but might still interfere with design
 - Respiration \sim from 0.1 to 0.5 Hz, typically 0.3 Hz
 - BOLD \sim from 0.01 to 0.05 (broad)
 - typical fMRI Nyquist frequency $<$ 0.5 Hz

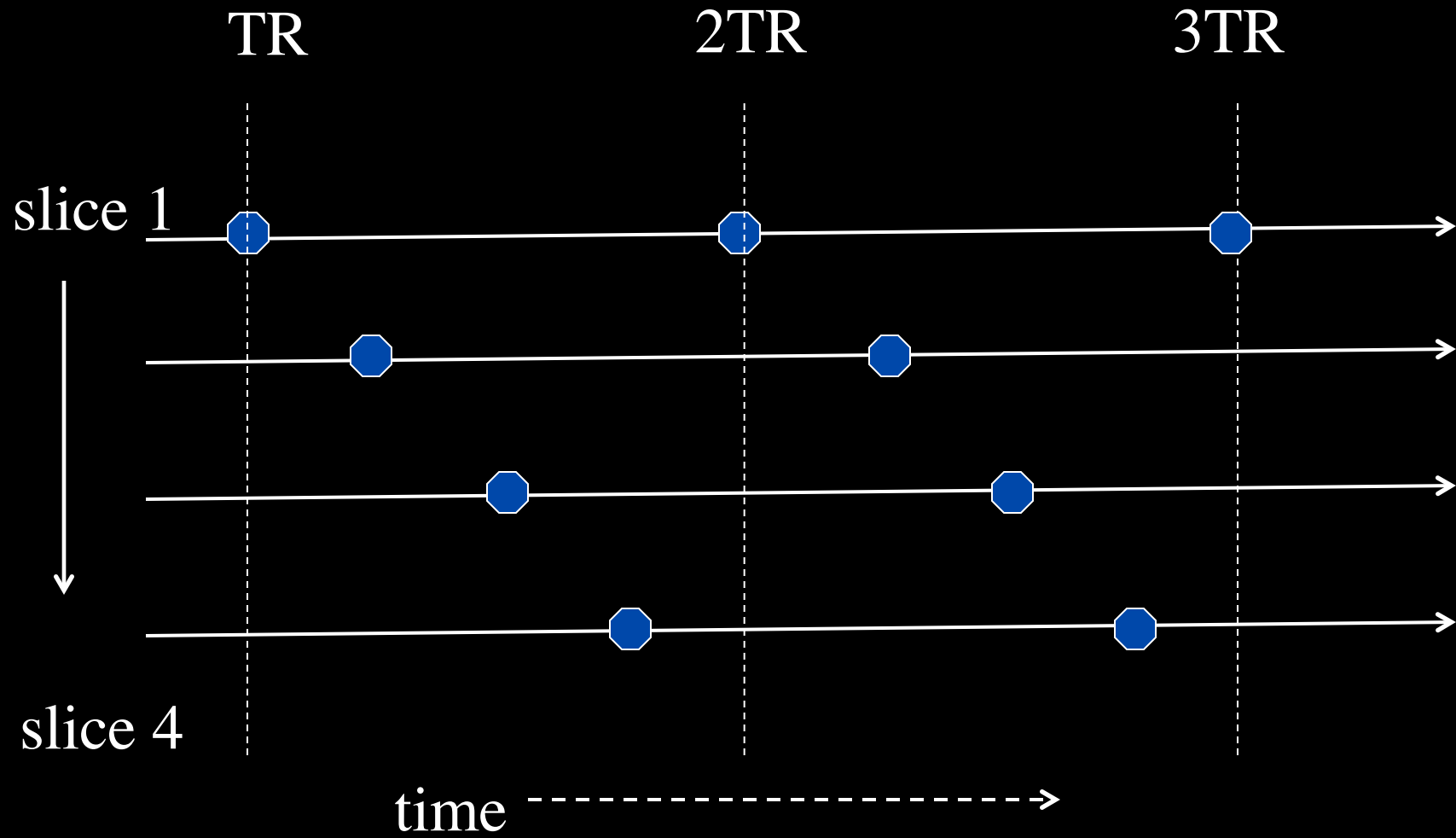
Timing Errors

- MR images are typically collected one slice at a time (exception: 3D imaging)
- The slices can be collected sequentially or interleaved.
- Delay between slice excitations is typically
$$TR / (\text{num. slices})$$
- Therefore, the time series are time-shifted differently in each slice

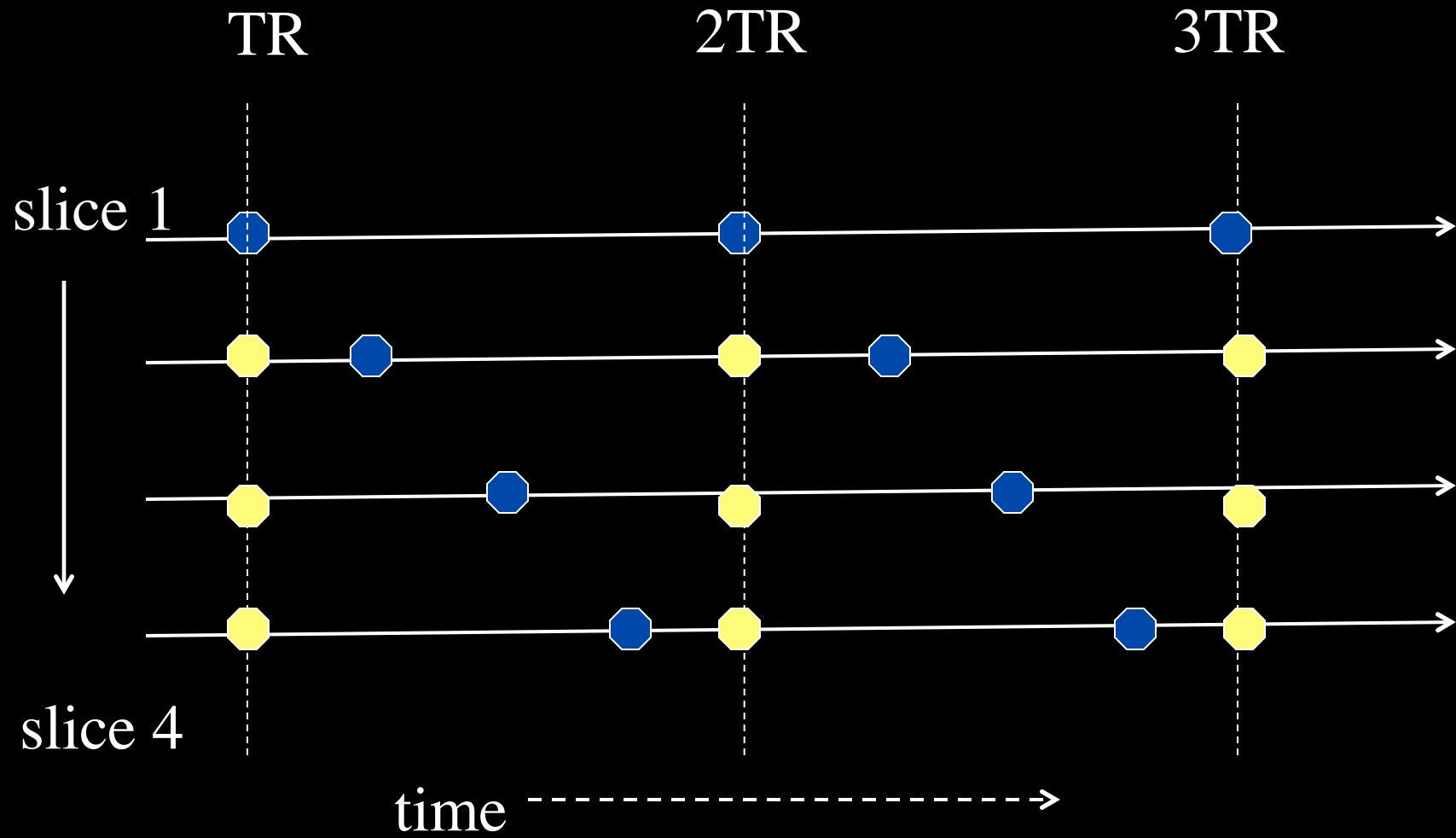
FMRI data “layout”



Acquisition



Acquisition



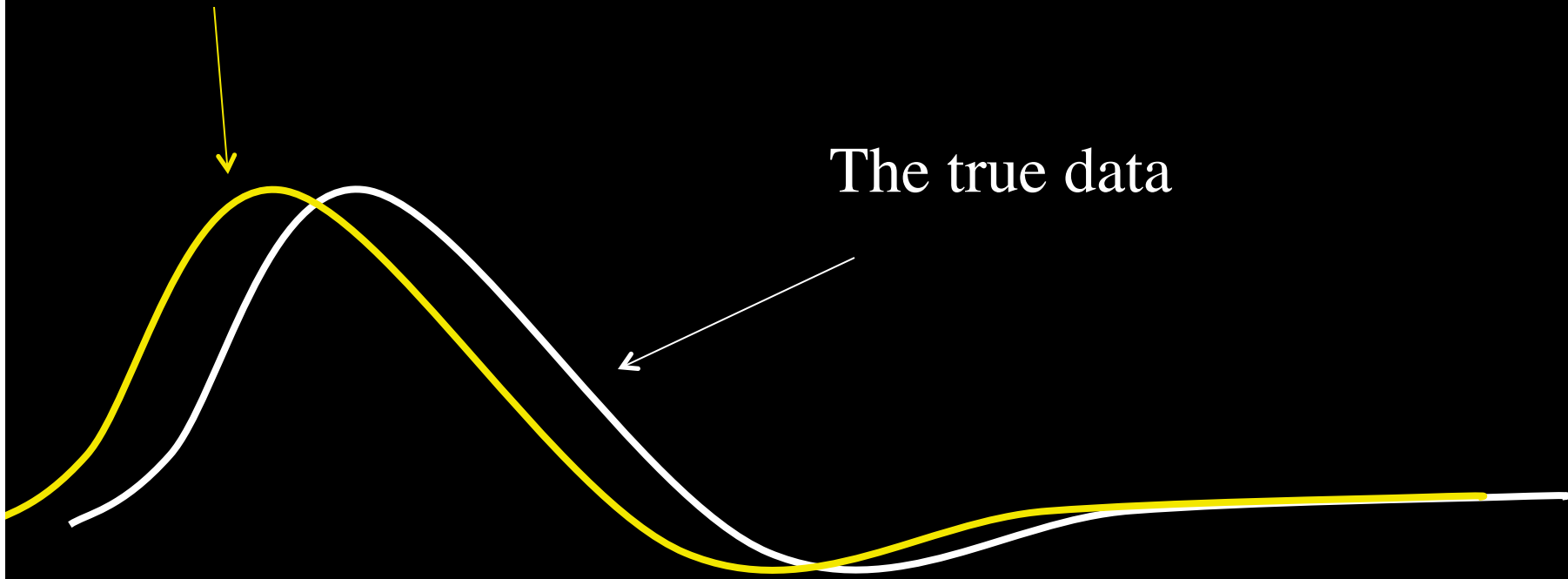
Sampling Error in Time



Sampling Error in Time

How the data looks

The true data



so shift it back!

Movement

In 2 Dimensions:

- shift from (x_1, y_1) to (x_2, y_2) :

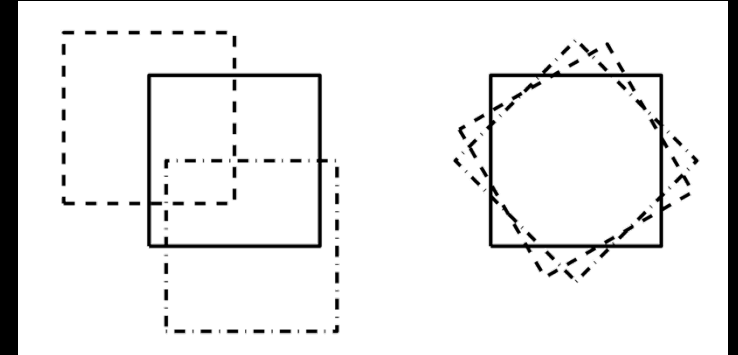
$$x_2 = x_1 + \Delta x$$

$$y_2 = y_1 + \Delta y$$

- Rotation from (x_1, y_1) to (x_2, y_2) :

$$x_2 = x_1 \cos(\theta) + y_1 \sin(\theta)$$

$$y_2 = -x_1 \sin(\theta) + y_1 \cos(\theta)$$



2-D Transformation matrix

- Both Together (note that the order matters)

$$x_2 = x_1 \cos(\theta) + y_1 \sin(\theta) + \Delta x$$

$$y_2 = -x_1 \sin(\theta) + y_1 \cos(\theta) + \Delta y$$

or In Matrix Form ...

$$\begin{pmatrix} x_2 \\ y_2 \\ 1 \end{pmatrix} = \begin{pmatrix} \cos(\theta) & \sin(\theta) & \Delta x \\ -\sin(\theta) & \cos(\theta) & \Delta y \\ 0 & 0 & 1 \end{pmatrix} \begin{pmatrix} x_1 \\ y_1 \\ 1 \end{pmatrix}$$

2-D Transformation matrix

$$(x_2, y_2) = A(x_1, y_1)$$

this extends to N-dimensions too

3-D Rotation matrices

$$\begin{pmatrix} \cos(\theta) & \sin(\theta) & 0 & 0 \\ -\sin(\theta) & \cos(\theta) & 0 & 0 \\ 0 & 0 & 1 & 0 \\ 0 & 0 & 0 & 1 \end{pmatrix} \quad \begin{pmatrix} \cos(\theta) & 0 & \sin(\theta) & 0 \\ 0 & 1 & 0 & 0 \\ -\sin(\theta) & 0 & \cos(\theta) & 0 \\ 0 & 0 & 0 & 1 \end{pmatrix} \quad \begin{pmatrix} 1 & 0 & 0 & 0 \\ \cos(\theta) & 1 & \sin(\theta) & 0 \\ -\sin(\theta) & 0 & \cos(\theta) & 0 \\ 0 & 0 & 0 & 1 \end{pmatrix}$$

xy plane
rotation

xz plane
rotation

yz plane
rotation

Estimation of Movement

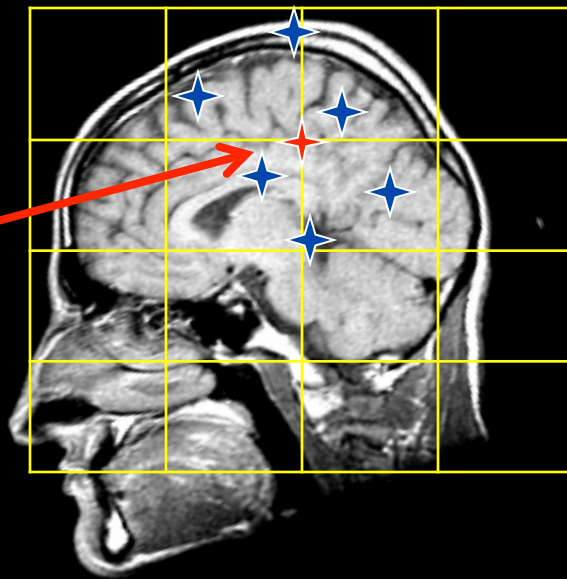
1. Choose a set of translations, rotations
2. Combine the six transformations matrices (linear operators) into one “rigid body” transformation

$$r_2 = A r_1$$

3. Resample the images at the new locations
4. Are the two images more alike?
5. Repeat and search for the best matrix A

Movement

Interpolate this
point from its
neighbors



Resampling the image

- Think of realignment as transforming the sampling grid, rather than the image.
- Interpolation:
 - Choose weighting function (kernel):
 - Nearest neighbor
 - bi-linear, tri-linear interpolation
 - sinc interpolation

Comparing images: cost function

- How do you know two images match?
 1. Least squares difference

$$\Sigma (I_1 - I_2)^2$$

2. Normalized correlation, correlation ratio

$$\frac{\Sigma(I_1 - I_2)}{(\text{Var}(I_1) \text{Var}(I_2))^{1/2}} \quad \frac{\text{Var}(E[I_1 I_2])}{\text{Var}(I_2)}$$

3. Mutual information

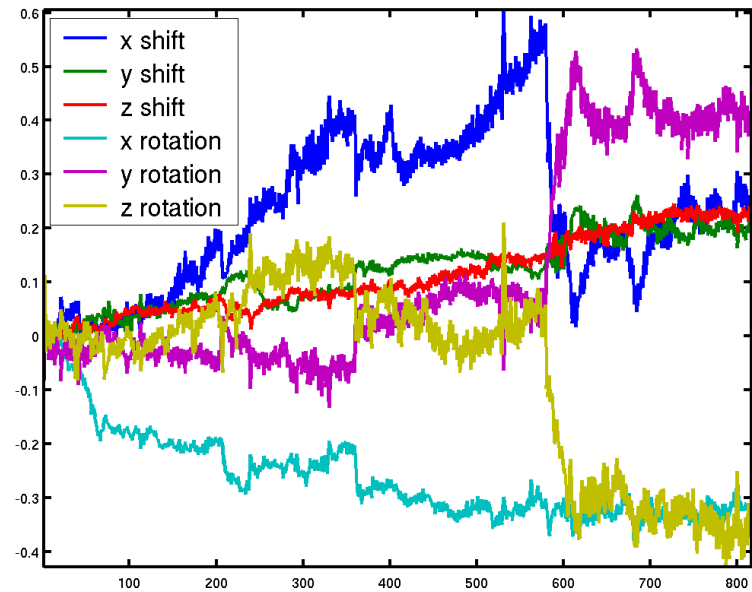
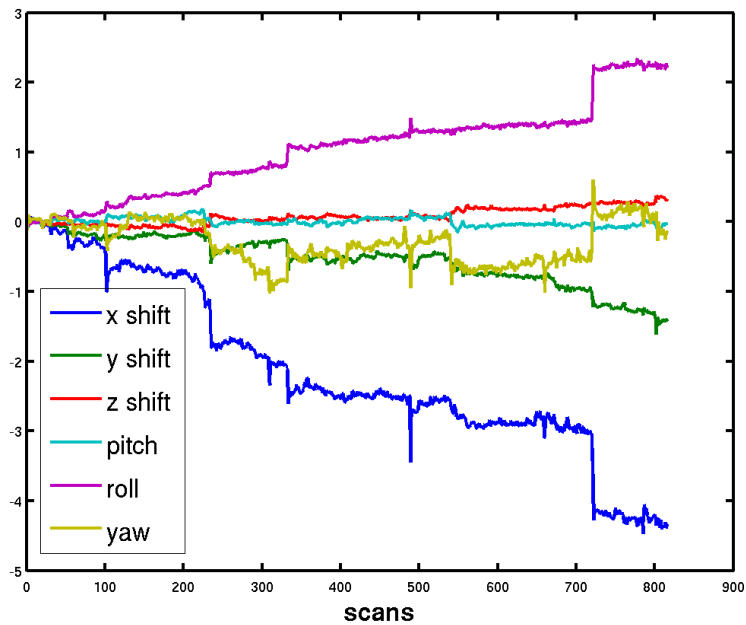
$$M(I_1, I_2) = \sum_{i,j} p(I_1, I_2) \log_2 \left(\frac{p(I_1, I_2)}{p(I_1)p(I_2)} \right)$$

4. others ...M. Jenkinson and S.M. Smith. Medical Image Analysis, 5(2):143-156, June 2001

Search Strategies

- least squares ($Y=\beta X$) ...
- Steepest descent: vary parameters and compute the gradient in the cost function (error). Keep going as long as it gets better.
- There are variations on this theme:
 - simplex
 - Newton's method / gradient descent
 - Adaptive methods
 - others...

Sample Movement Parameters



Movement Noise

- In addition to mixing voxels, you introduce a fluctuation in signal intensity during realignment
- This is a complicated function of the movement:
 - affects the k-space data acquisition
 - mixes partial volumes,
 - interpolation methods also have an effect on intensity.

Movement Noise corrections

- Minimize movement while acquiring data whenever possible !!
- Include movement parameters as confounding regressors.
 - Complicated function, but the signal fluctuation is well correlated with the movement parameters.
 - Including movement regressors will strongly reduce variance.
 - If movement is correlated with task = BIG TROUBLE!

Putting it all together : pre-processing stream

Functional Time Series

B0 map correction

Physio correction

Reconstruction

Slice Timing correction

Motion- realignment

SPM

Anatomical Images

Reconstruction

B1 homogeneity correction

brain extraction

registration

normalization

Statistical Map
in Standard
Space

