### BOLD fMRI:

signal source, data acquisition, and interpretation

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## Discussion series

- Week 1: Biological basis: where's the signal coming from?
- Week 2: Physical basis: what is the signal, how is it measured?
- Week 3: Imaging basics: image formation, noise, and artifacts.
- Week 4: The specific case of BOLD fMRI.
- Week 5: BOLD analysis: what's significant and what's not?
- Week 6: Spikes vs. BOLD: neural activity in visual areas

# BOLD fMRI

- The signal around veins and capillaries
- BOLD and CBF measurements
- Modeling the BOLD response
- Optimizing imaging for BOLD (flip angle, TE)
- Distortion (continued from last week)
- Motion correction prospective routines
- Sample BOLD experiment

#### Arteries, Capillaries, and Veins



dynamic averaging during diffusion of water molecules - Relatively high [dHb] (7

- Relatively high [dHb] (70 - 80%)

- Dynamic averaging from diffusion reduces effect of field inhomogeneities even further

#### Field Dependence



- Higher field means bigger effect from [dHb]induced inhomogeneities

- Less time for dynamic averaging

#### What does Spin Echo do?



Tissue protons experiencing dynamic dephasing see residual BOLD signal.

# Perfusion imaging (FAIR technique)

1) Non-selective inversion pulse before acquisition (free bonus BOLD data in specially marked packages!)





3) Difference image produces perfusion map  $\Delta S \propto 2M_{0A}$ 





### Modeling the BOLD response

Relaxation rates are proportional to blood volume and [dHb]:  $R_2^* \sim V \ [dHb]^{\beta}$ , with  $\beta > 1$  because of diffusion, and the fact that increasing blood volume displaces tissue water

 $dS/S = S_{max}(1 - vc^{\beta})$ 

where v is relative blood volume, and c is relative [dHb]. Using

c = m/f (where m is relative CMRO<sub>2</sub>, and f is relative CBF) v = f<sup>\alpha</sup> (\alpha ~ 0.4, from animal studies)

(this represents just a partial understanding of Box 16 in the Buxton book ... the goal would be to use this to model the temporal dynamics of the response to neural activity ... the hemodynamic response)



# Early Dip

#### Buxton sums it up well: controversial and important





#### High-resolution mapping of isoorientation columns by fMRI

Dae-Shik Kim, Timothy Q. Duong and Seong-Gi Kim

nature neuroscience · volume 3 no 2 · february 2000

Yacoub, E. and Hu, X. (1999). Detection of the early negative response in fMRI at 1.5 Tesla. Mag Reson Med. 41: 1088-1092.



Yacoub et al. (1999). "Further evaluation ..." Mag Reson Med. 41: 436.

Image Number

100

# Optimizing acquisition for BOLD

- Echo time
  - BOLD effect is strongest when  $TE \sim T2^{\boldsymbol{*}}$
- Ernst angle
  - For repetition times ~ T1, steady state signal is greatest when flip angle is less than 90 degrees
- Resolution/SNR trade-off
- Total scan time

# Distortion in EPI images

- Basic problem: a voxel's location is inferred from it's resonant frequency
- Each accumulated 360 phase shift moves the signal one voxel
- Example: chemical shift of fat at 7T
  - Fat resonates at 3.5 ppm
  - At 7T, this is 3.5 x  $10^{-6}$  x 300MHz, or ~1000Hz
  - A 64 x 64 EPI image has a read-out time of 500us
  - So time for phase evolution along phase encode direction is 64 x 500us = 32ms
  - $-32ms \times 1000Hz = 32$  pixel shift



Subcutaneous fat, shifted 32 / 64 pixels in EPI image



#### Prospective motion correction

Navigator echo samples k-space before each image:

 $S(k,\theta) = S_{ref}(k, \theta - \alpha)e^{ik(x\cos + y\sin)}$ 

rotation shows up as rotation

translation shows up as a phase shift



SNAV. Welch et al, MRM 47:32-21 (2002)



Ward et al. (2000). Prospective multi-axial motion correction for fMRI. *Mag Reson Med.* **43**: 459.

ONAV successfully corrects stimulusrelated head motion

 SNAV is more computationally demanding