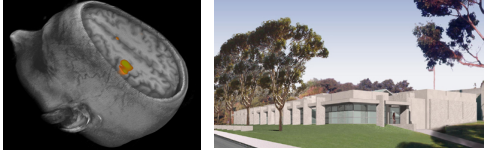


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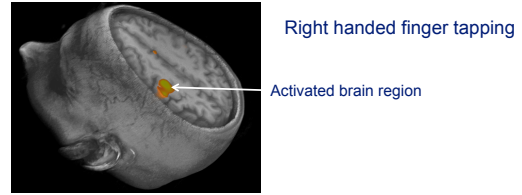
## Functional MRI

Richard B. Buxton  
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W. M. Keck Center for Functional MRI, UCSD

## Mapping Brain Activation



The *blood oxygenation level dependent* (BOLD) effect:

The MR signal is slightly stronger when blood is more oxygenated, and blood oxygenation *increases* when neural activity increases

## Blood Flow and O<sub>2</sub> Metabolism

Blood flow delivers O<sub>2</sub> and glucose and clears CO<sub>2</sub>

Key players:

CMRO <sub>2</sub> = cerebral metabolic rate of O <sub>2</sub>	1.6 micromol/g-min
E = O <sub>2</sub> extraction fraction	40%
CBF = cerebral blood flow	0.5 ml/g-min
[O <sub>2</sub> ] <sub>a</sub> = total arterial O <sub>2</sub>	8 micromol/ml

$$CMRO_2 = E CBF [O_2]_a$$

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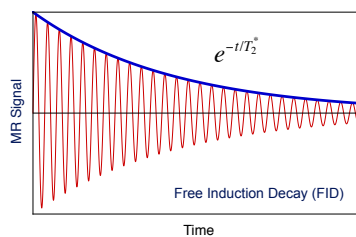
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$$CMRO_2 = E CBF [O_2]_a$$

E decreases with neural activation! [Fox and Raichle, *PNAS* 83:1140 (1986)]

Venous deoxy-hemoglobin = E [O<sub>2</sub>]<sub>a</sub> ↓

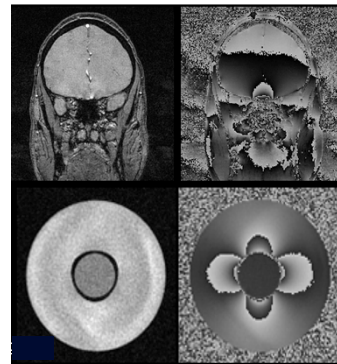
## Key properties of the NMR signal



Resonant Frequency:  $\nu_0 = \gamma B_0$   
(128 MHz at 3T)

Relaxation Time:  $T_2^*$   
(~50 ms at 3T)

## Magnetic Susceptibility Effects

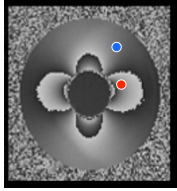


Large scale field gradients:  
Susceptibility differences between air, water and bone

$$\Delta B \approx \Delta \chi B_0$$

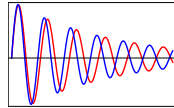
Microscopic field gradients:  
Deoxy-hemoglobin (dHb) alters the susceptibility of blood

### Signal decay due to deoxy-hemoglobin



Magnetic field variations within a voxel lead to protons precessing at slightly different rates

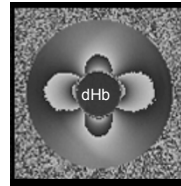
Local signals become out of phase, reducing the net signal



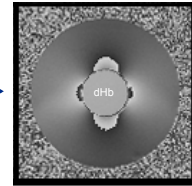
Deoxy-hemoglobin distorts the magnetic field around blood vessels, reducing the MR signal

### The BOLD response to activation

Blood flow increases more than  $O_2$  metabolism, increasing venous oxygenation (less deoxy-hemoglobin)

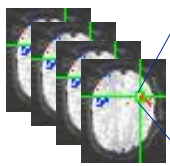


Neural Activation

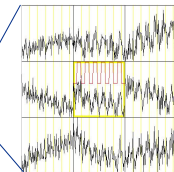


Magnetic field is less distorted, and the MR signal increases

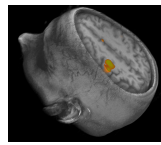
### The basic fMRI experiment



Dynamic imaging during "on" and "off" blocks of the stimulus

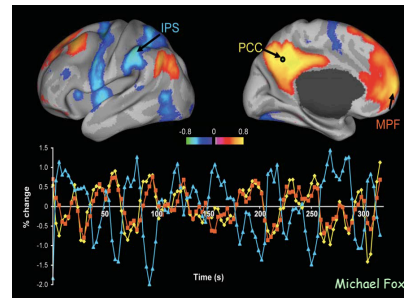


Correlation of each voxel's time course with the stimulus pattern.



Map areas of significant correlation overlaid on the anatomical image

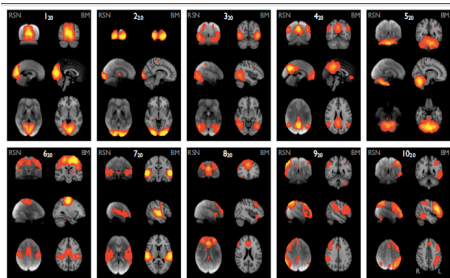
### Brain networks from resting state correlations



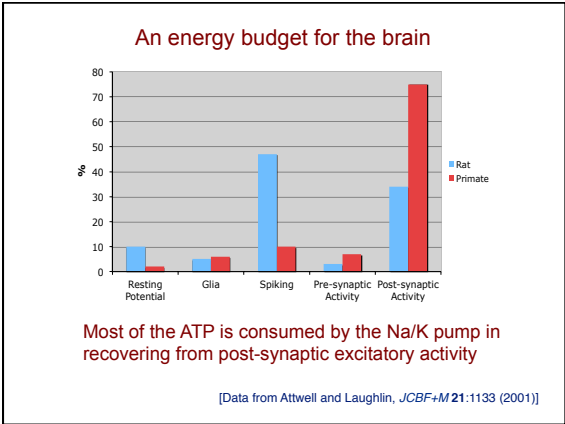
Functional Connections

### Correspondence of activation maps and resting correlation patterns

(Smith et al, PNAS 2009)



### The physiological basis of the BOLD effect



### The importance of continuous delivery of metabolites

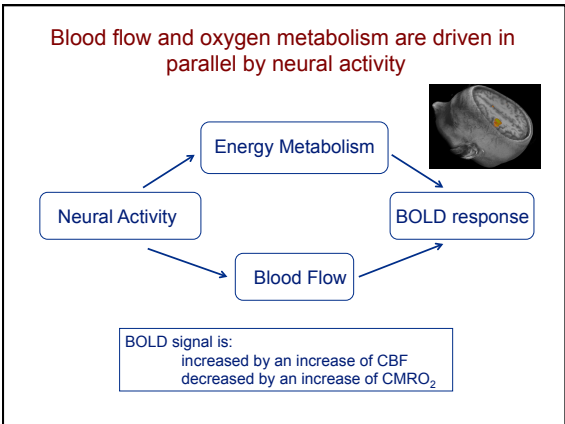
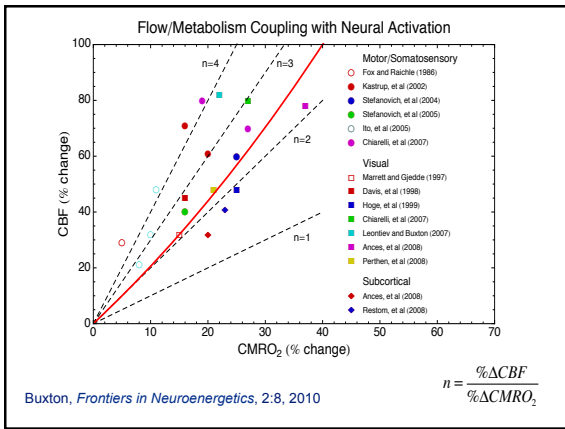
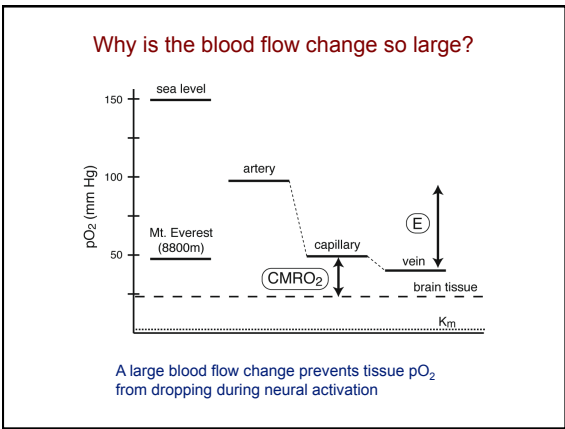
**Brain energy metabolism**

Glycolysis:  $\text{Glc} \rightarrow 2 \text{ATP}$

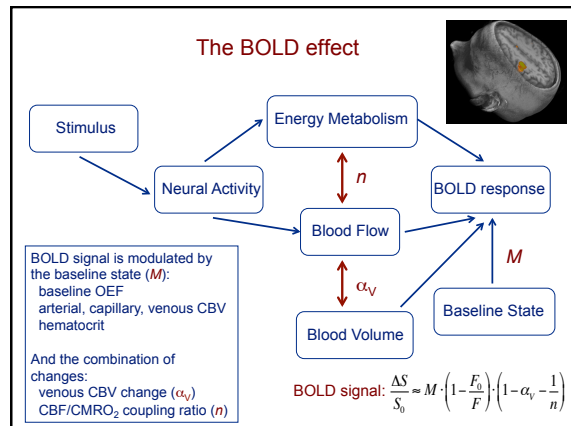
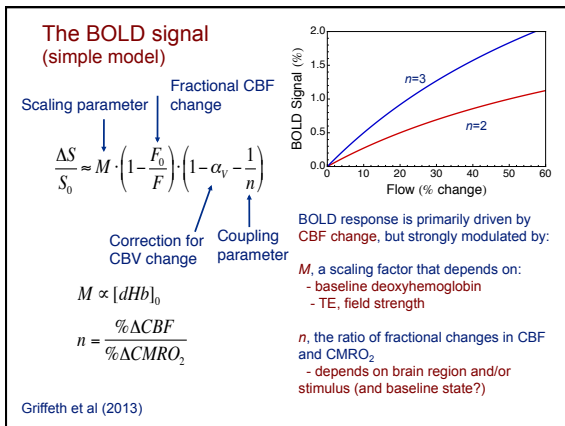
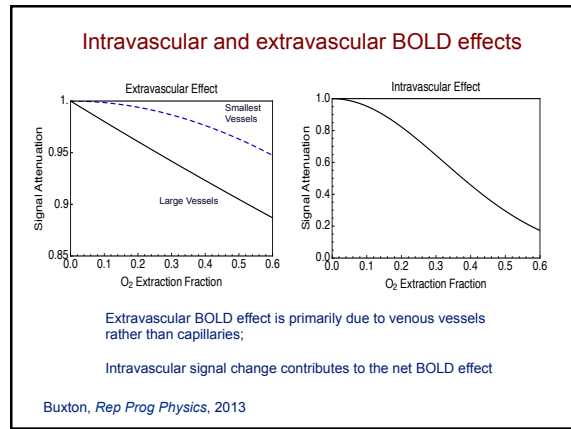
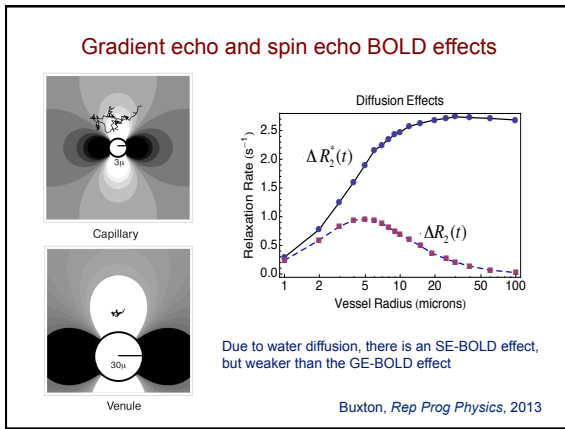
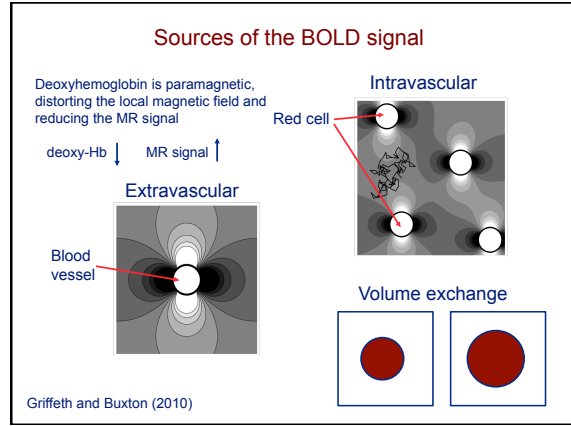
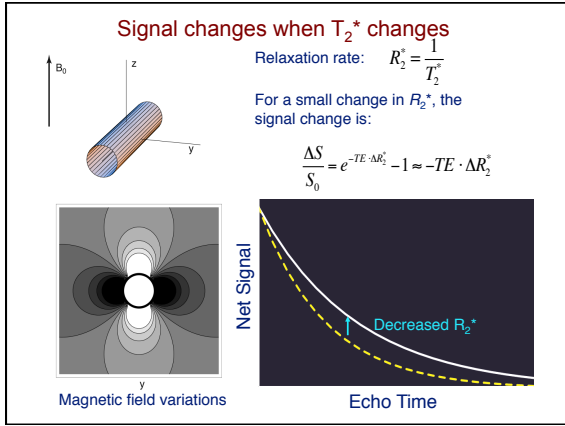
Oxidative metabolism:  $\text{Glc} + \sim 6 \text{O}_2 \rightarrow \sim 34 \text{ATP}$

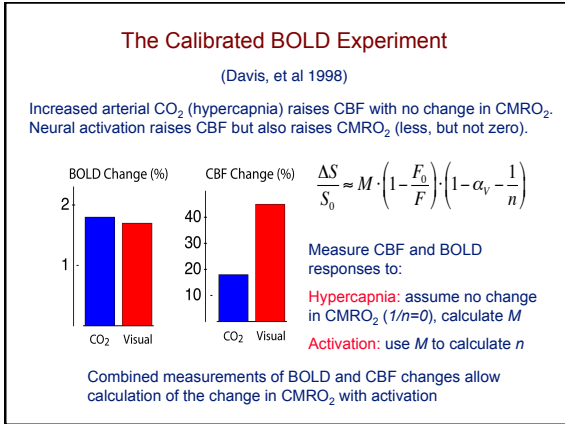
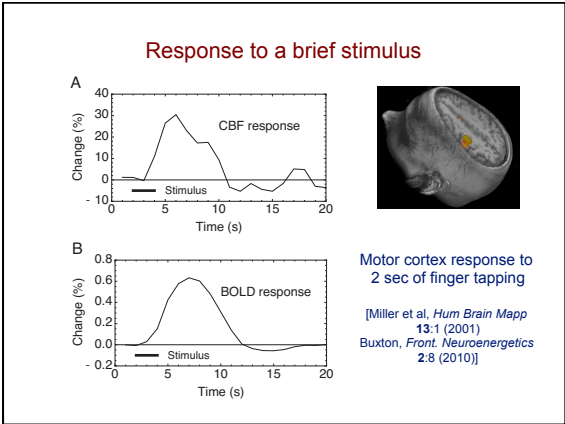
Metabolite	Metabolic rate	Tissue concentration	Lifetime
Glucose	0.3 mM/min	2.0 mM	~7 min
$\text{O}_2$	1.6 mM/min	0.03 mM (tissue only)	~1 sec
$\text{O}_2$		0.3 mM (plus blood)	~10 sec

There is very little reserve of  $\text{O}_2$  in the brain, requiring a responsive blood flow



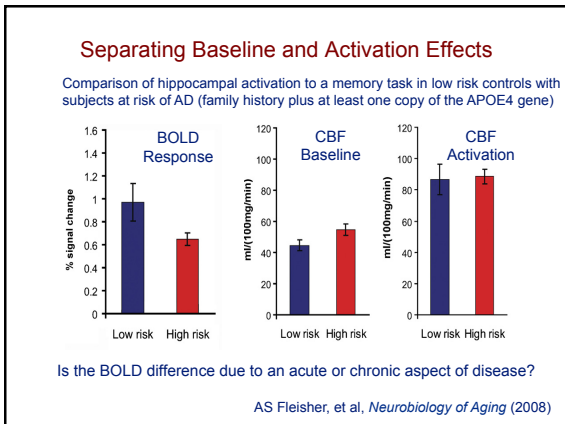
### The physical basis of the BOLD effect





### Interpreting the BOLD Response:

How can we go beyond mapping where brain activity changes?



### Caffeine

Caffeine blocks adenosine receptors

Adenosine has two effects on the brain:  
inhibits neural activity  
increases blood flow

Basic questions about CBF and CMRO<sub>2</sub>:  
How does caffeine alter the baseline state?  
How does caffeine alter the response to a stimulus?

