

# BENEFRI Workshop 2019

Methods in Experimental Neurosciences:  
From Animal Models to Humans

*u<sup>b</sup>*

b  
UNIVERSITÄT  
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## fMRI in Neuroscience

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University of Bern

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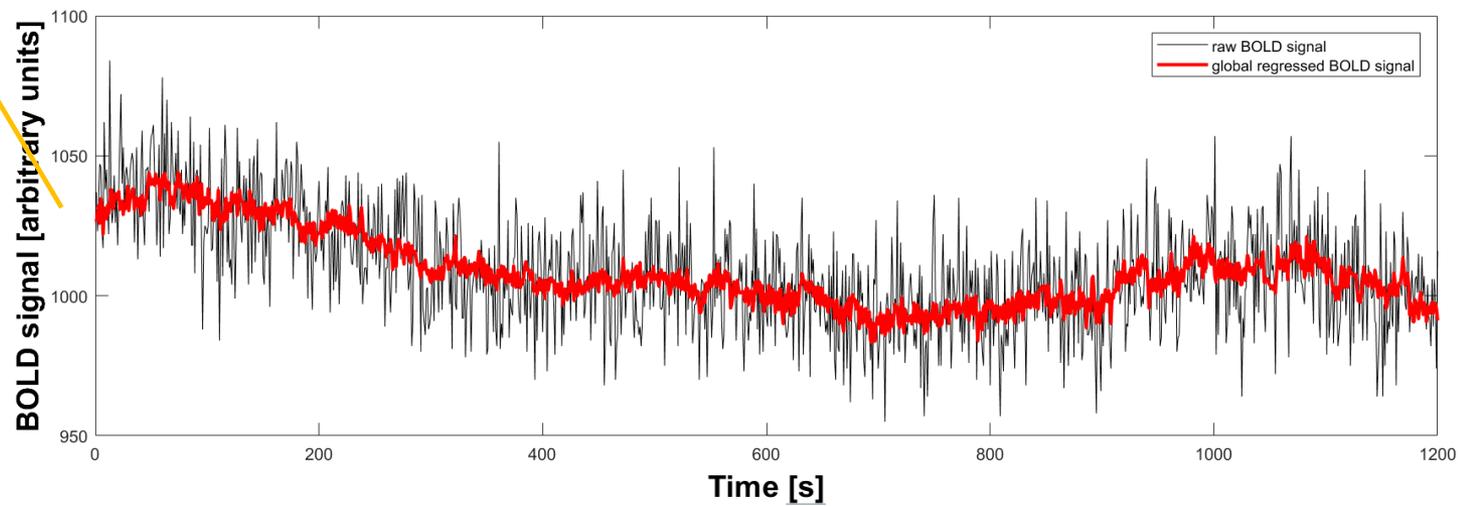
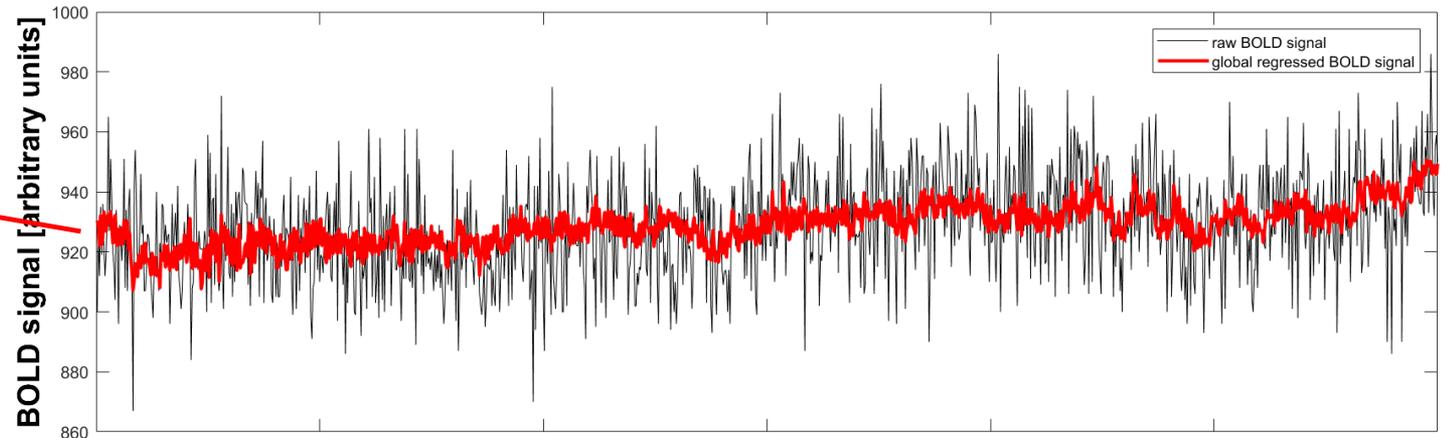
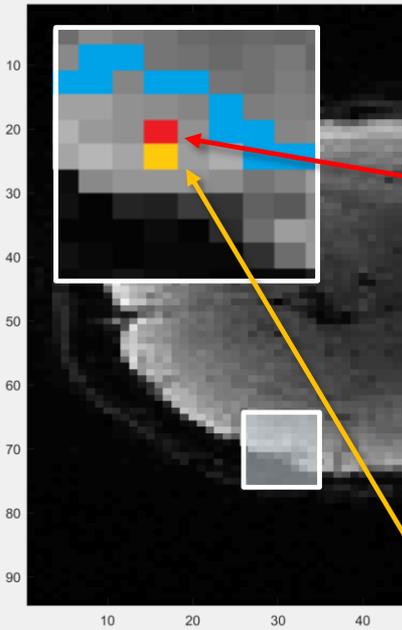


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DIENSTE BERN

# roadmap

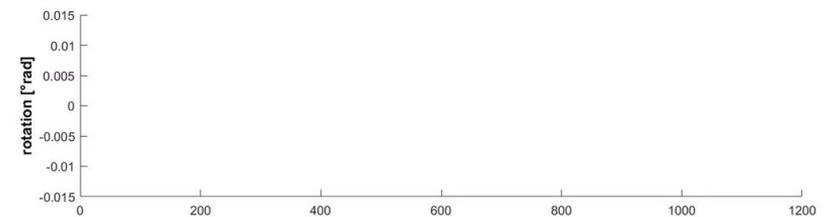
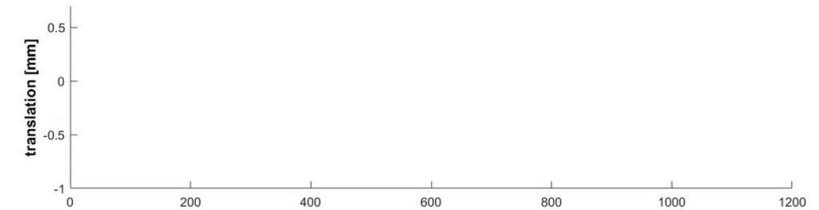
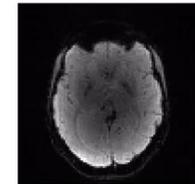
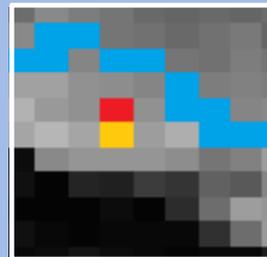
- 9:15-10:00**     **Basic concepts of **functional** Neuroimaging**  
fMRI Signal, task-dependent fMRI, resting state fMRI, Functional Network Analysis, processing pipeline, statistical testing, Random Effects, General Linear Model and MRI physics.
- 10:15-11:00**     **Basic concepts of **structural** Neuroimaging**  
Voxel Based Morphometry, Cortical Thickness, Cortex based inter-subject alignment, Diffusion Tensor Imaging, Tract-Based Spatial Statistics.
- 11:15-12:00**     **Advanced Neuroimaging Methods in Neurosciences**  
Non-BOLD fMRI, Cerebral Blood flow (CBF), calibrated fMRI, Multimodal Imaging.

# Variance in the neighbourhood

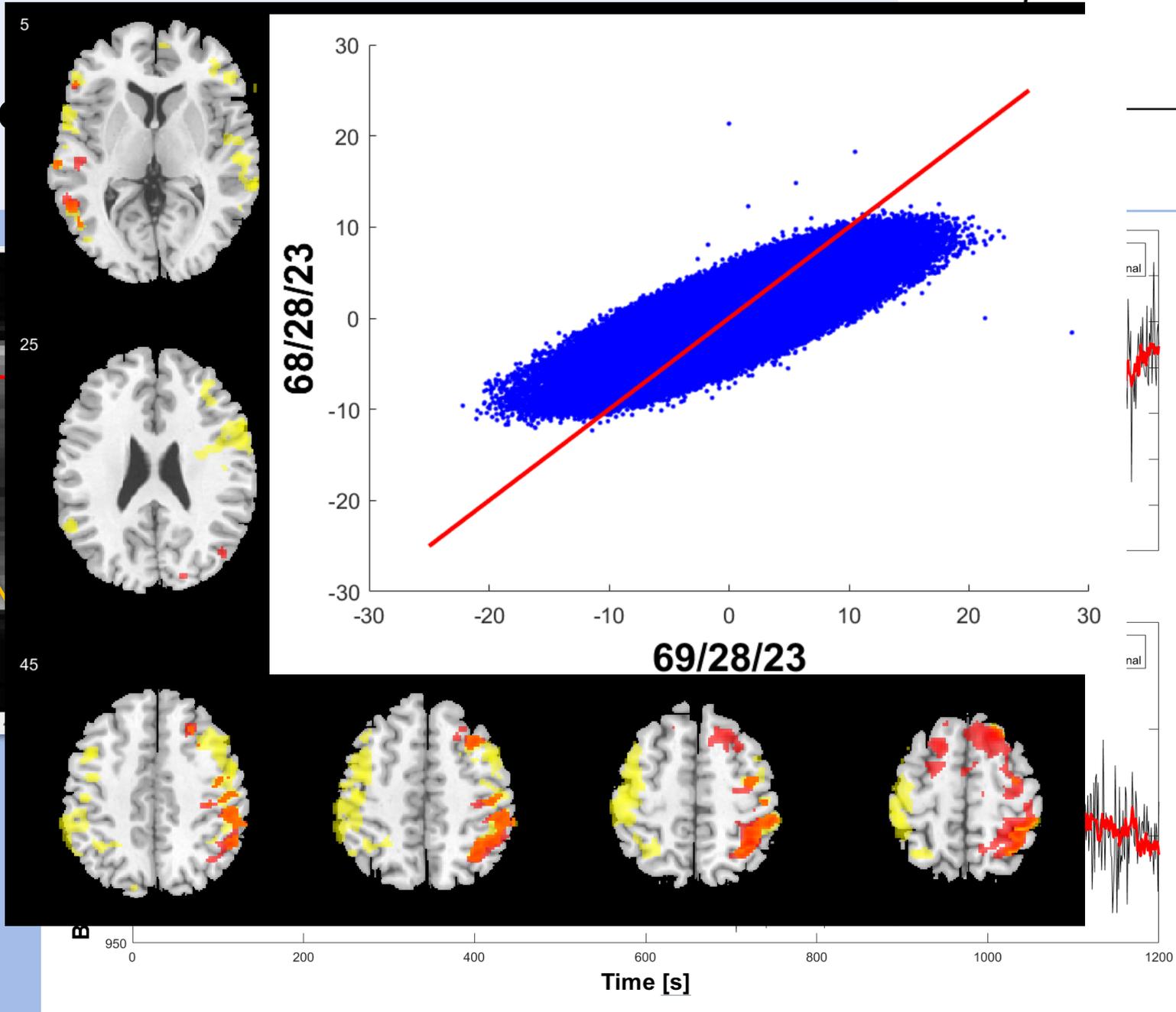


# Closer lock at BOLD signal

Variance in:		68/28/23	69/28/23
gm		301.18	415.69
BOLD (fit)		238.26	250.76
gm - BOLD(fit)		50.41	177.43
wm	298.35		
csf	273.24		
motion	0.06		



# Variance



# roadmap

- 9:15-10:00**     **Basic concepts of **functional** Neuroimaging**  
fMRI Signal, task-dependent fMRI, resting state fMRI, Functional Network Analysis, processing pipeline, statistical testing, Random Effects, General Linear Model and MRI physics.
- 10:15-11:00**     **Basic concepts of structural Neuroimaging**  
Voxel Based Morphometry, Cortical Thickness, Cortex based inter-subject alignment, Diffusion Tensor Imaging, Tract-Based Spatial Statics.
- 11:15-12:00**     **Advanced Neuroimaging Methods in Neurosciences**  
Non-BOLD fMRI, Cerebral Blood flow (CBF), calibrated fMRI, Multimodal Imaging.

## fMRI “roadmap”

- > Slice Time correction
- > Coregistration (2D fmri → 3D anatomy)
- > Segmentation (3D anatomy)
- > Normalisation (3D anatomy)
- > 1. and 2. level statistics

# White paper on fMRI

<https://www.humanbrainmapping.org/files/2016/COBIDASreport.pdf>

Committee on Best Practices in Data Analysis and Sharing (COBIDAS)

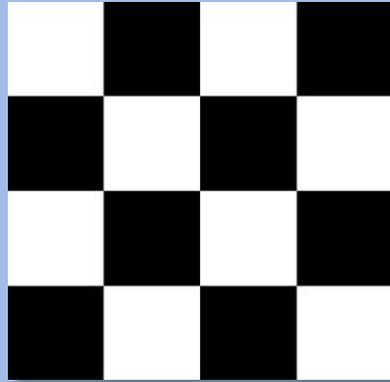
Organization of Human Brain Mapping (OHBM)

Suggestions and recommendations on how to deal with fMRI Data

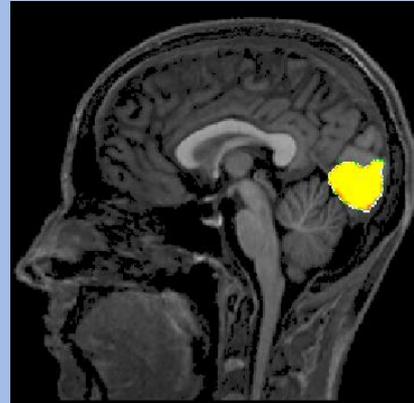
Data acquisition, Design, Data analysis, etc.

# Functional Magnetic Resonance Imaging

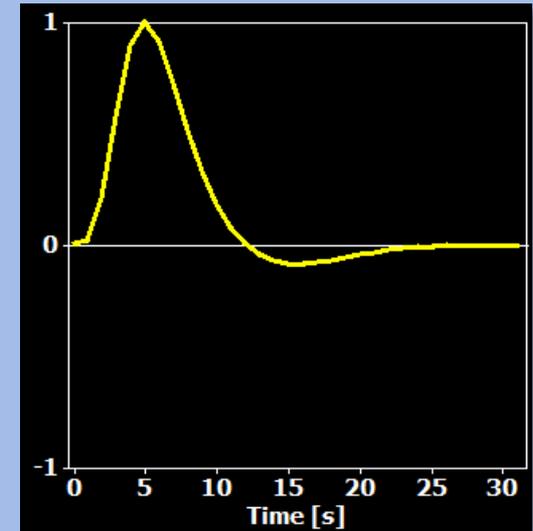
## fMRI: measure of neuronal activity?



Stimulus



Reaction



Measure

**BOLD = Blood Oxygen  
Level Dependent**

Concentration between  
Oxyhemoglobin (diamagnetic)  
and Deoxyhemoglobin (paramagnetic)  
in the veins

If neuronal activity high



**BOLD contrast high**

# Image Orientation

*Proc. Natl. Acad. Sci. USA*  
Vol. 87, pp. 9868–9872, December 1990  
Biophysics

## **Brain magnetic resonance imaging with contrast dependent on blood oxygenation**

(cerebral blood flow/brain metabolism/oxygenation)

S. OGAWA, T. M. LEE, A. R. KAY, AND D. W. TANK

Biophysics Research Department, AT&T Bell Laboratories, Murray Hill, NJ 07974

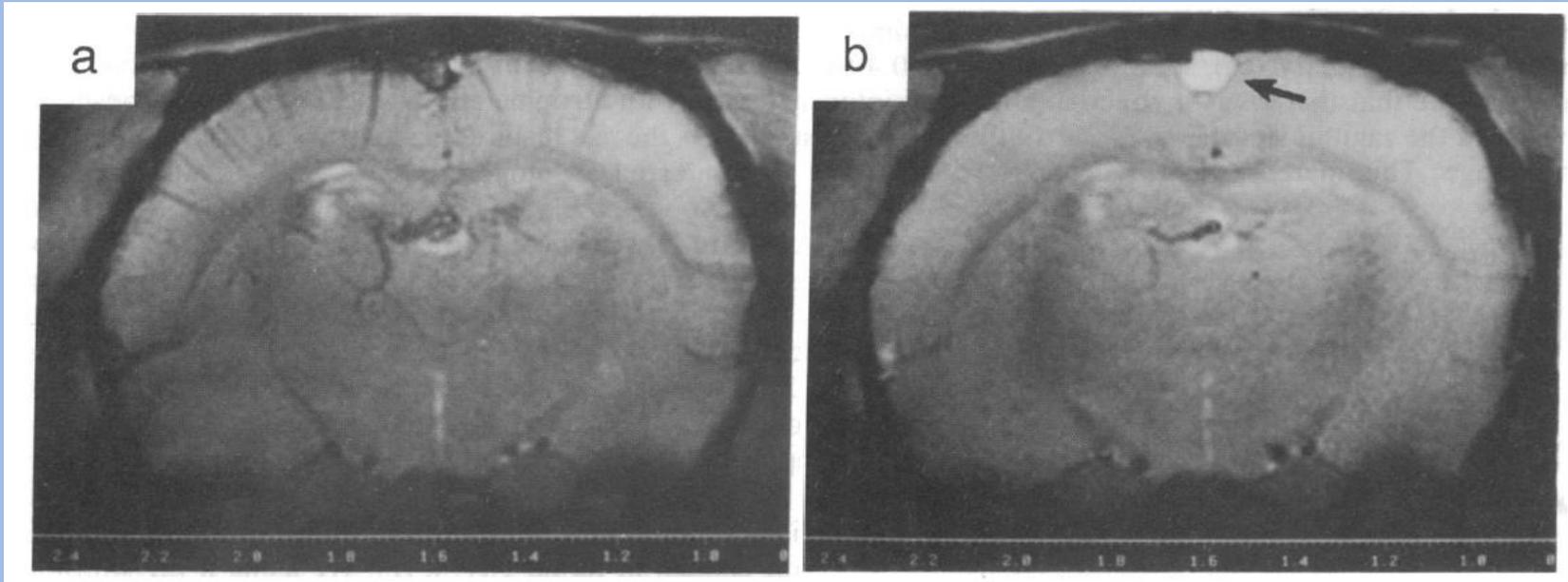
*Proc. Natl. Acad. Sci. USA*  
Vol. 89, pp. 5951–5955, July 1992  
Neurobiology

## **Intrinsic signal changes accompanying sensory stimulation: Functional brain mapping with magnetic resonance imaging**

(cerebral blood flow/blood oxygenation/visual cortex/positron emission tomography/magnetic susceptibility)

SEJI OGAWA<sup>†</sup>, DAVID W. TANK<sup>†</sup>, RAVI MENON<sup>‡</sup>, JUTTA M. ELLERMANN<sup>‡</sup>, SEONG-GI KIM<sup>‡</sup>,  
HELLMUT MERKLE<sup>‡</sup>, AND KAMIL UGURBIL<sup>‡</sup>

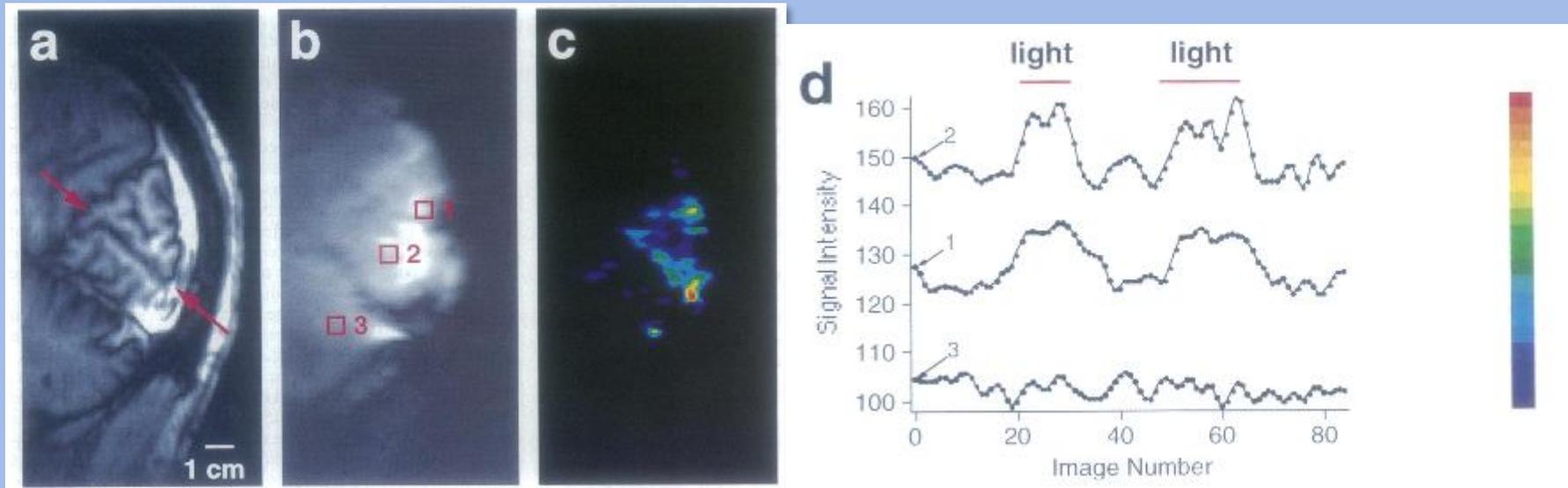
# What does Neuroimaging means?



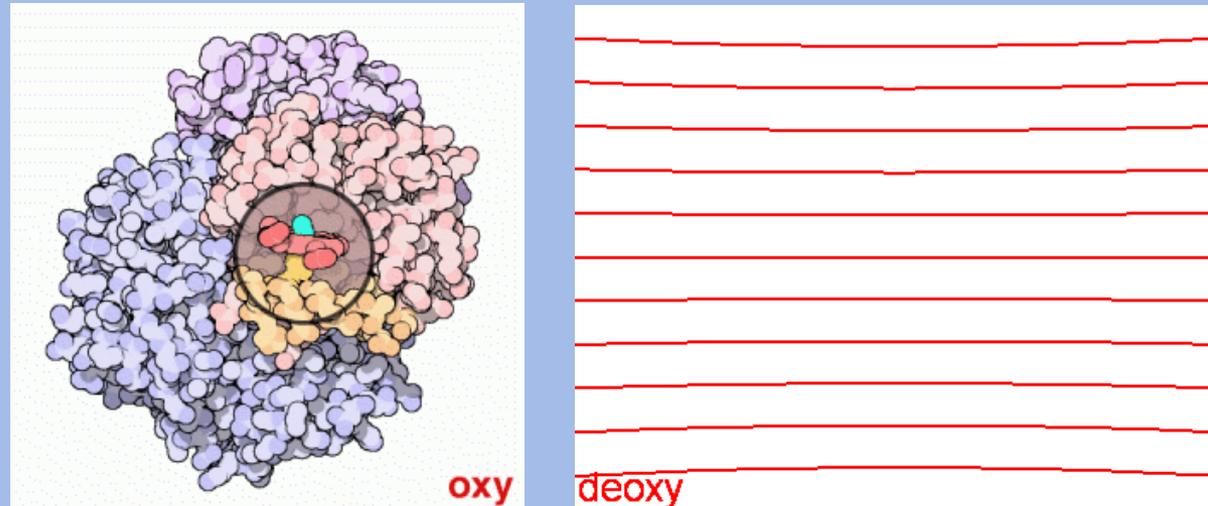
**100% O<sub>2</sub>**

**90% O<sub>2</sub> / 10% CO<sub>2</sub>**

# Contrast signal in different regions



# Active agent responsible for fMRI



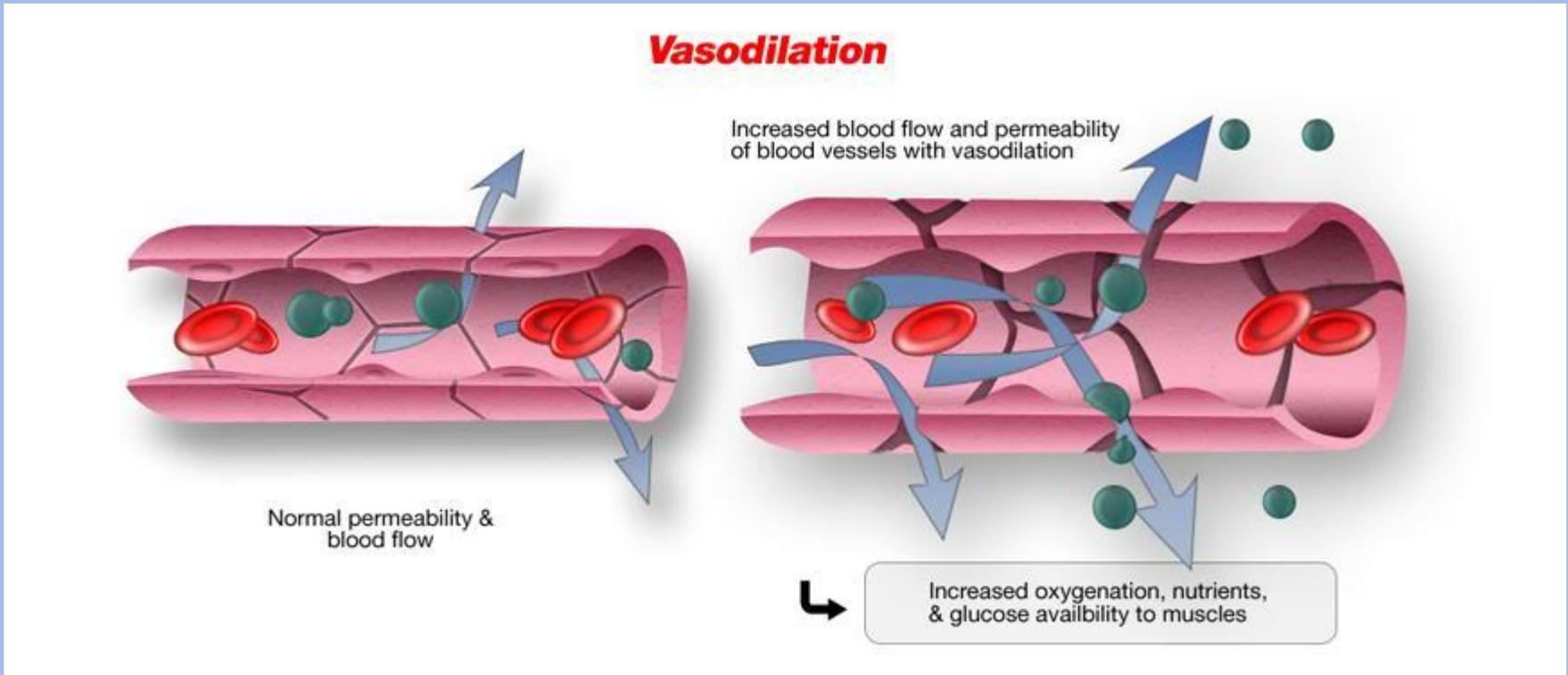
Deoxyhemoglobin: **paramagnetic** ( $\chi > 0$ )  
Hb (4 unpaired  $e^- \rightarrow S=2$ )

$T_2^* \downarrow$

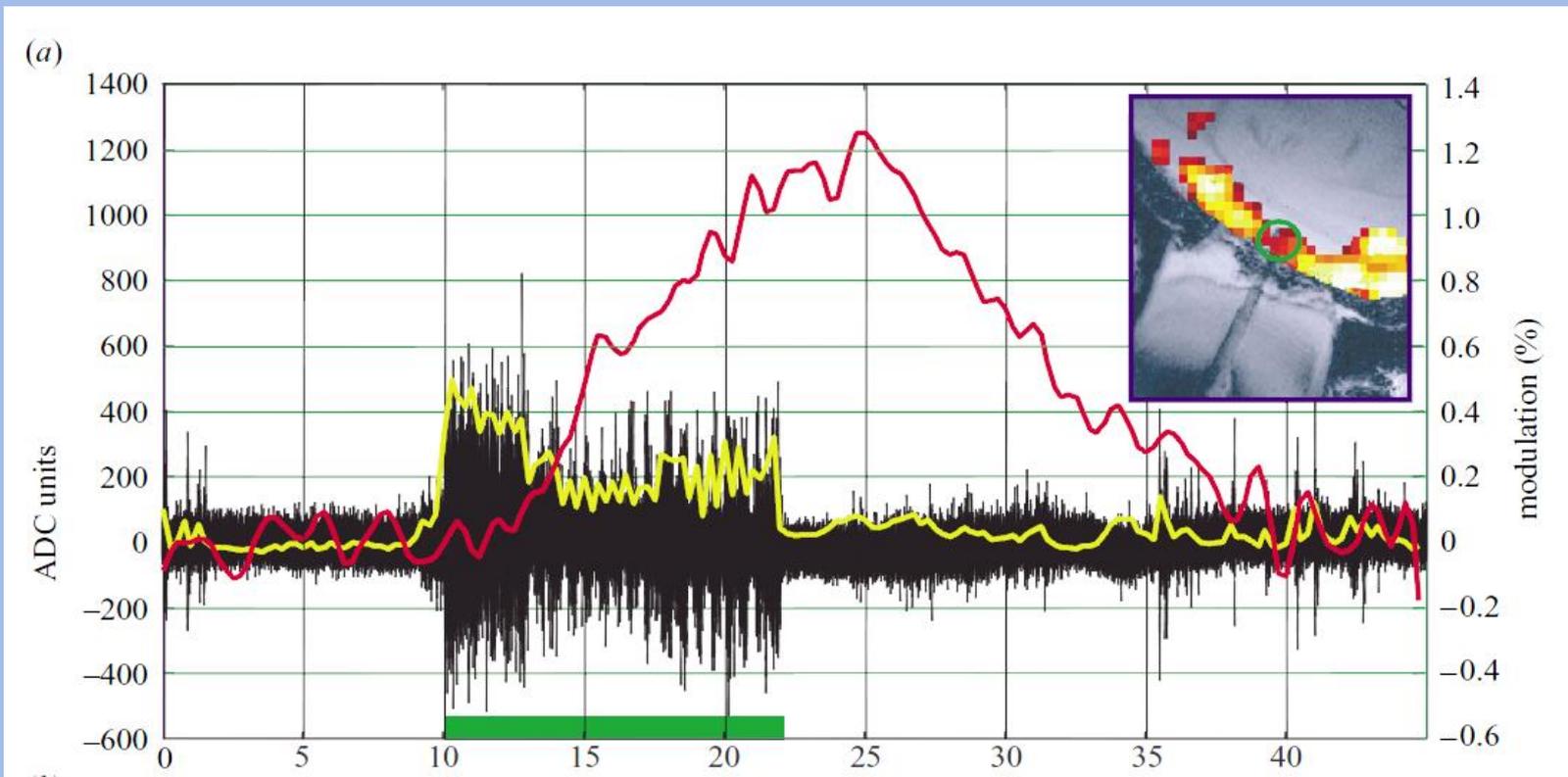
Oxyhemoglobin: **diamagnetic** ( $\chi < 0$ )  
HbO<sub>2</sub>  $S=0$

$T_2^* \uparrow$

# fMRI: vasodilatation



# fMRI: Neuronal Activity

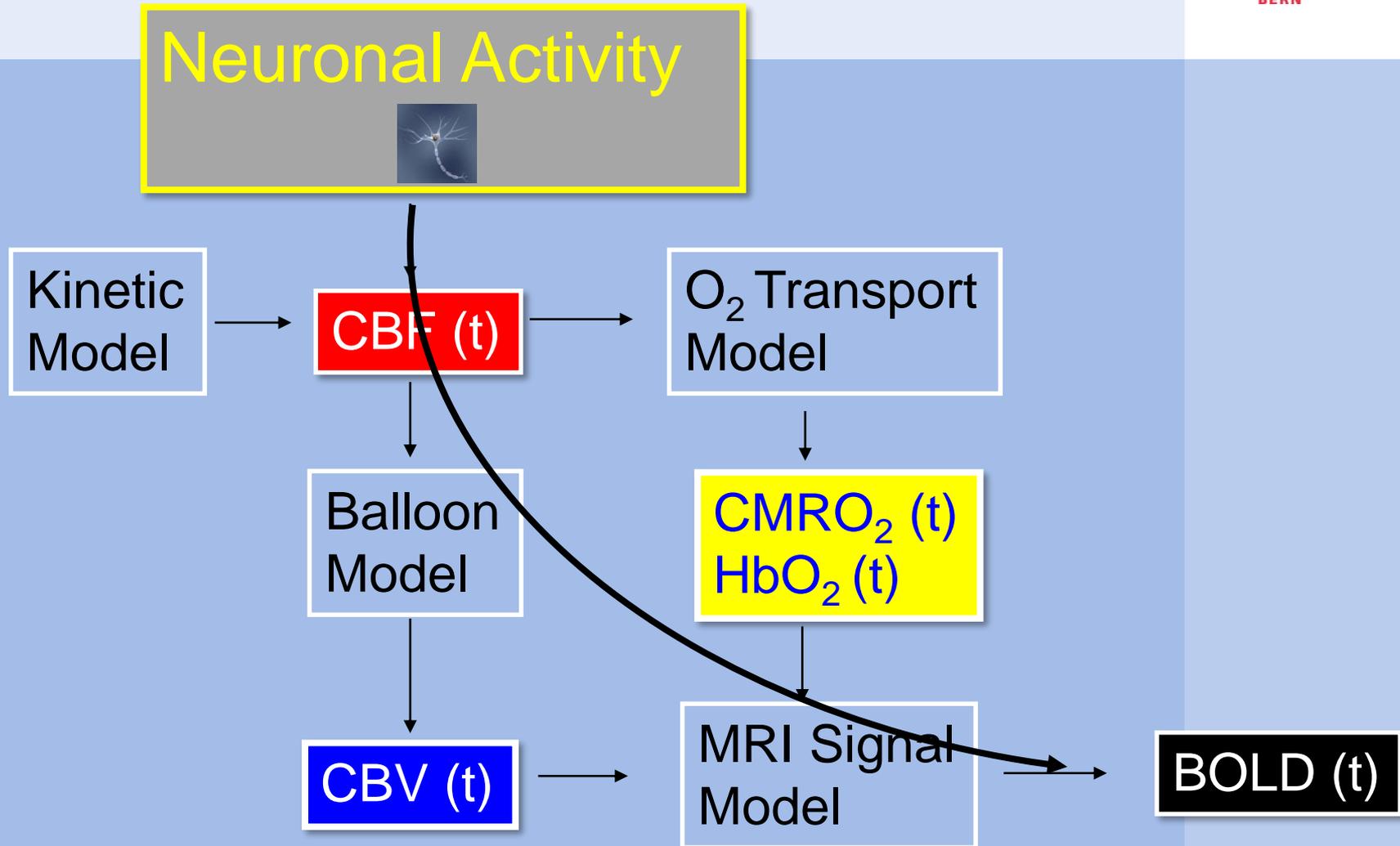


Spike Activity (black / yellow)

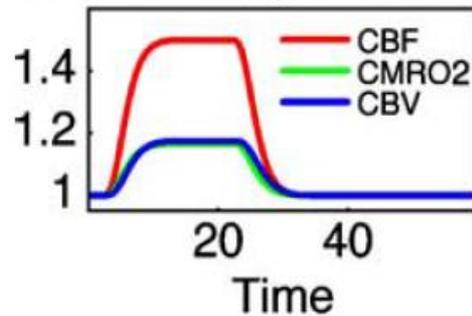
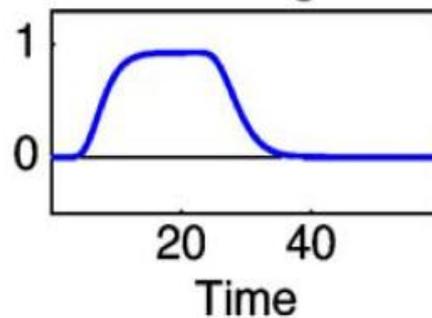
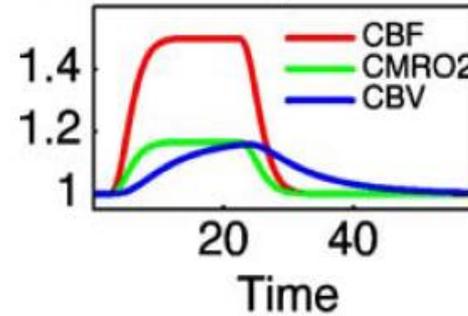
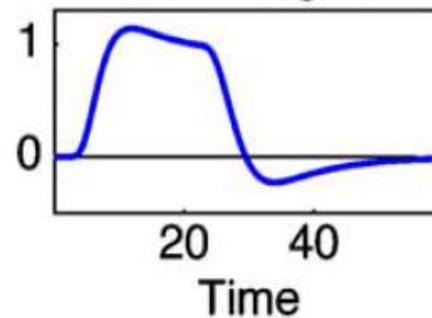
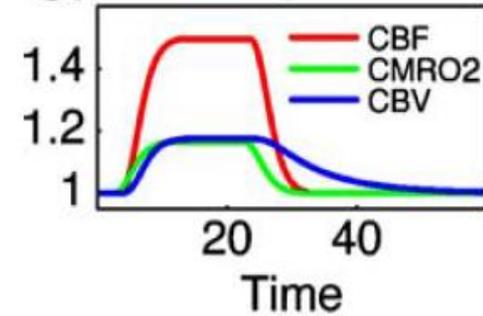
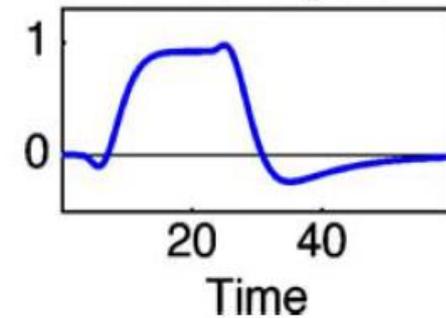
BOLD Activity (red)

(Simultan Intracortical / BOLD Measure on wake animal)

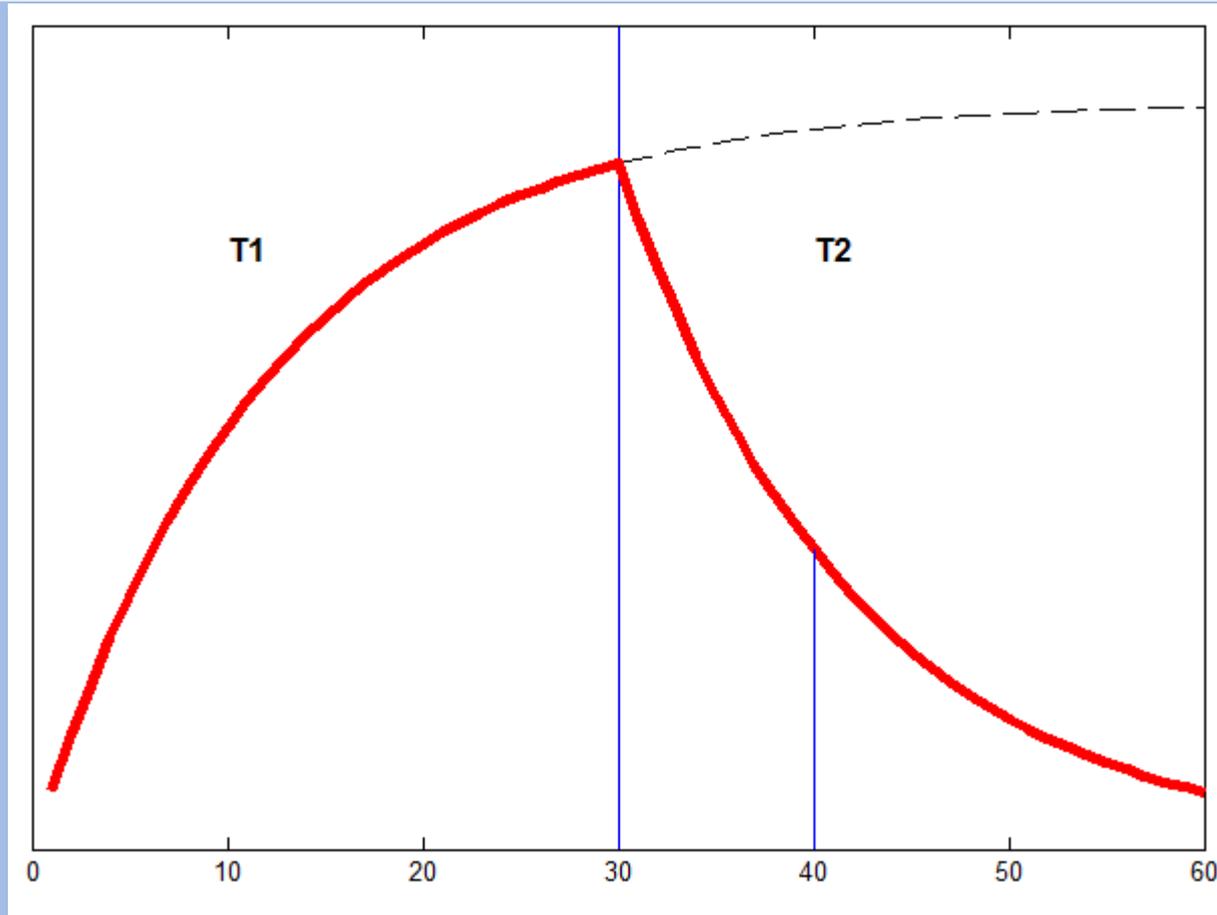
# Understanding BOLD



# Balloon Model for understanding BOLD

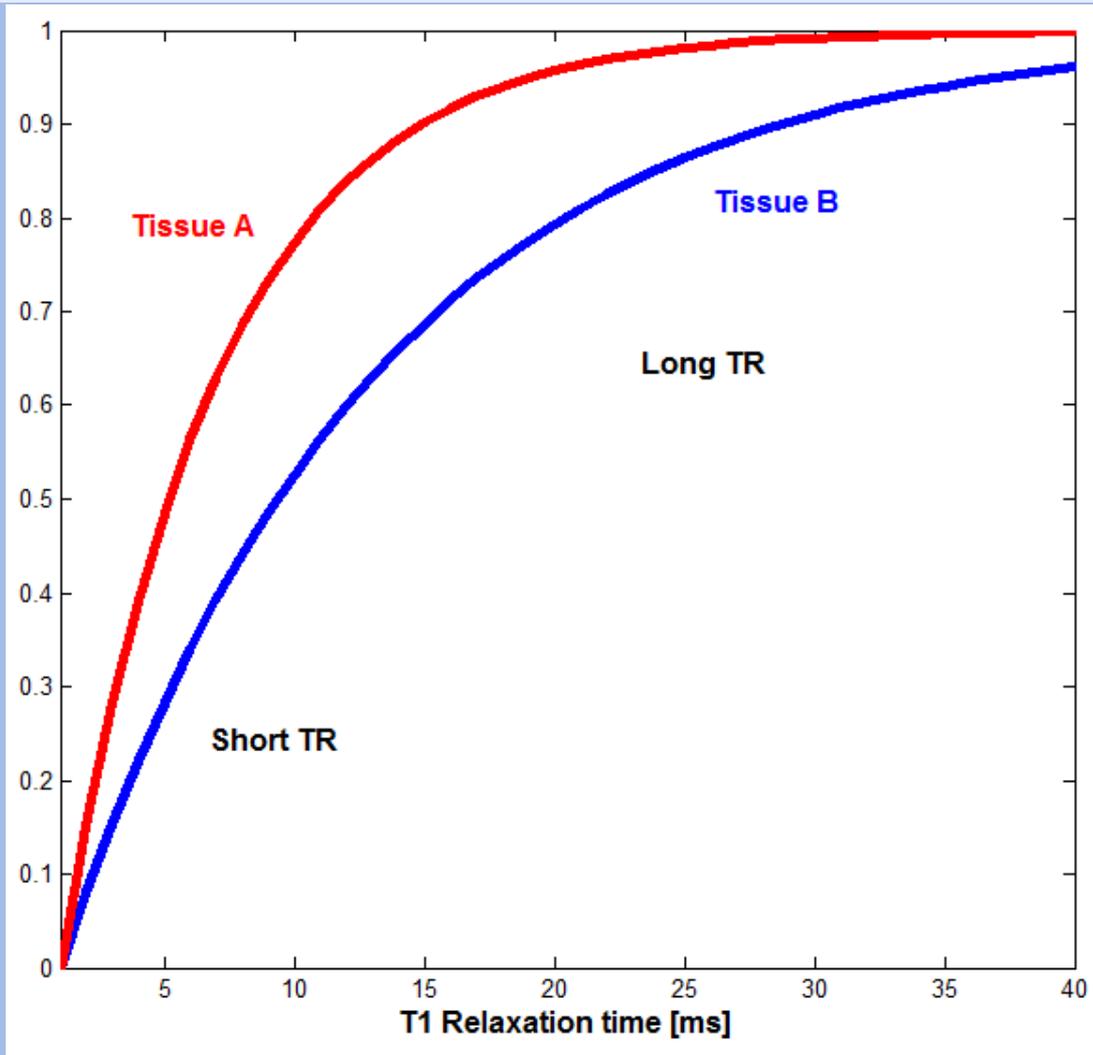
**A. Hemodynamics****BOLD Signal****B. Hemodynamics****BOLD Signal****C. Hemodynamics****BOLD Signal**

# TR, TE (repetition- and echo time)

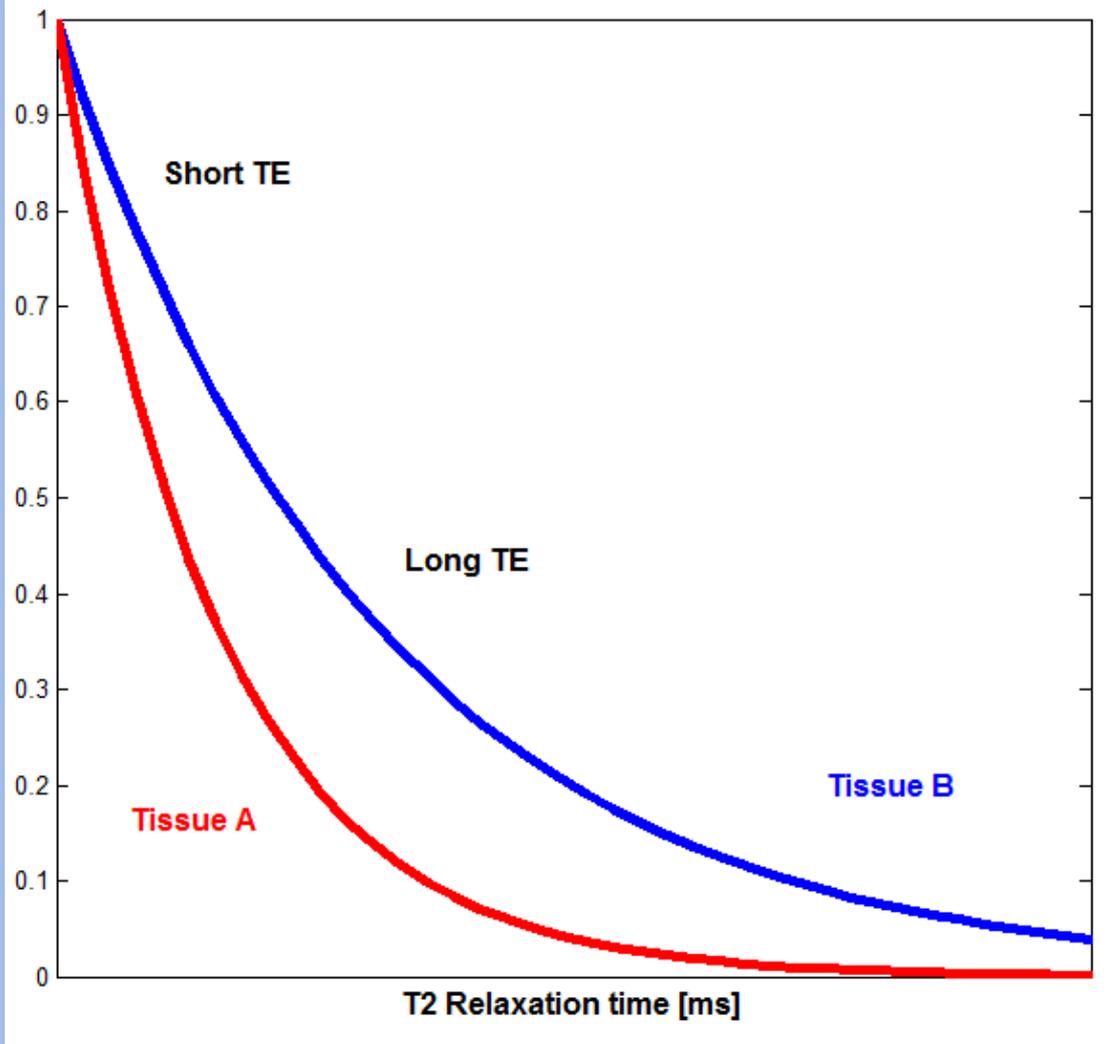


TR TE

# TR, $T_1$ relaxation



# TE, T<sub>2</sub> relaxation

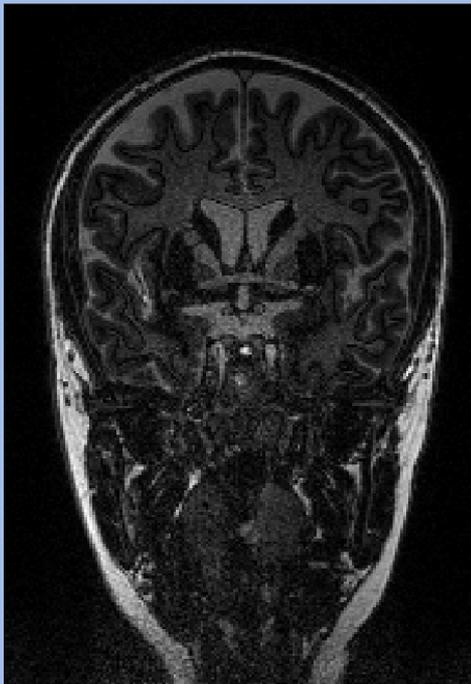


# T<sub>1</sub> and T<sub>2</sub> relaxation times in tissue

Tissue	T <sub>1</sub> [ms]	T <sub>2</sub> [ms]
Gray matter	600	80
White matter	950	100
Blood @ 3T	1450	275
Cerebro Spinal Fluid (CSF)	4500	2200
Fat	250	60

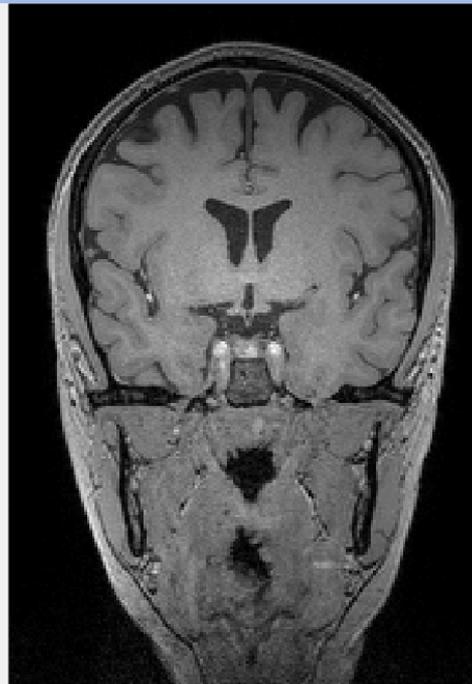
# Typical T1 sequence: mp2rage (TA: 8 min 22 sec)

1<sup>st</sup> inversion



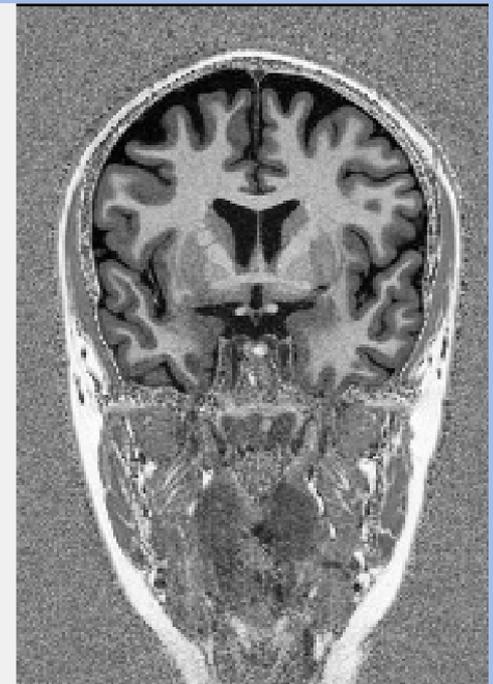
$T_{i1}=700$  ms  
Flip.angl.= $4^\circ$

2<sup>nd</sup> inversion

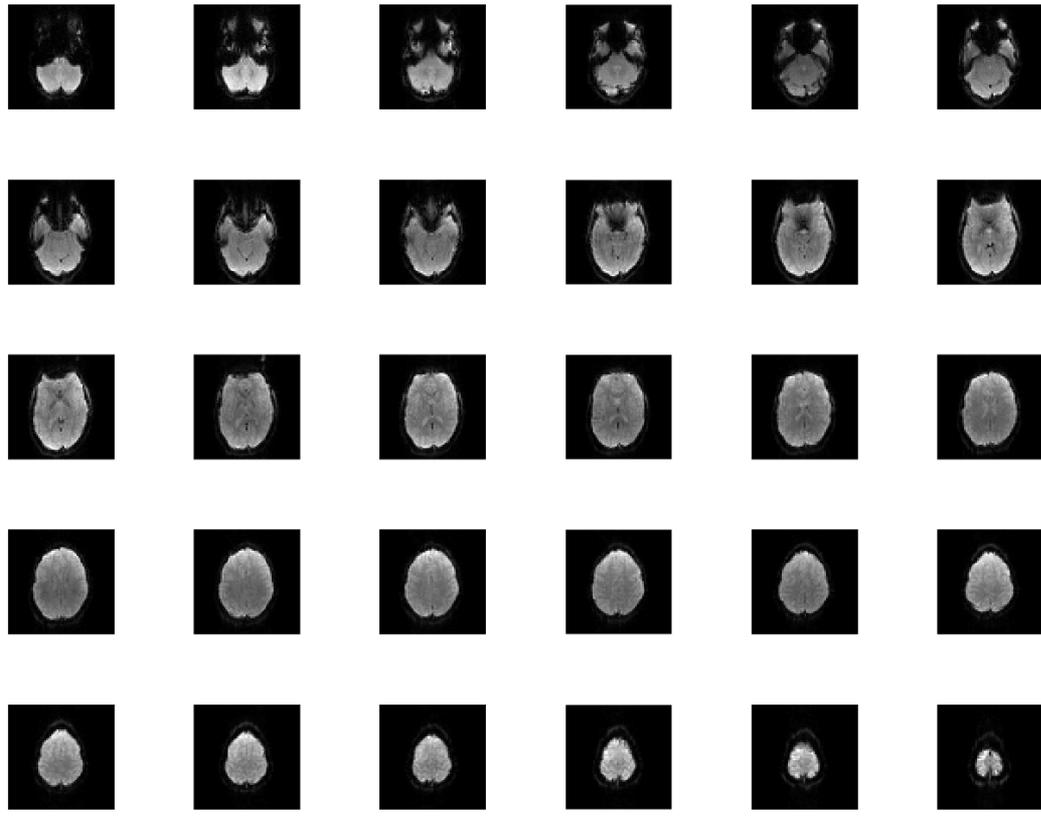


$T_{i2}=2500$  ms  
Flip.angl.= $5^\circ$

combined



# Typical T2 sequence: multi-band

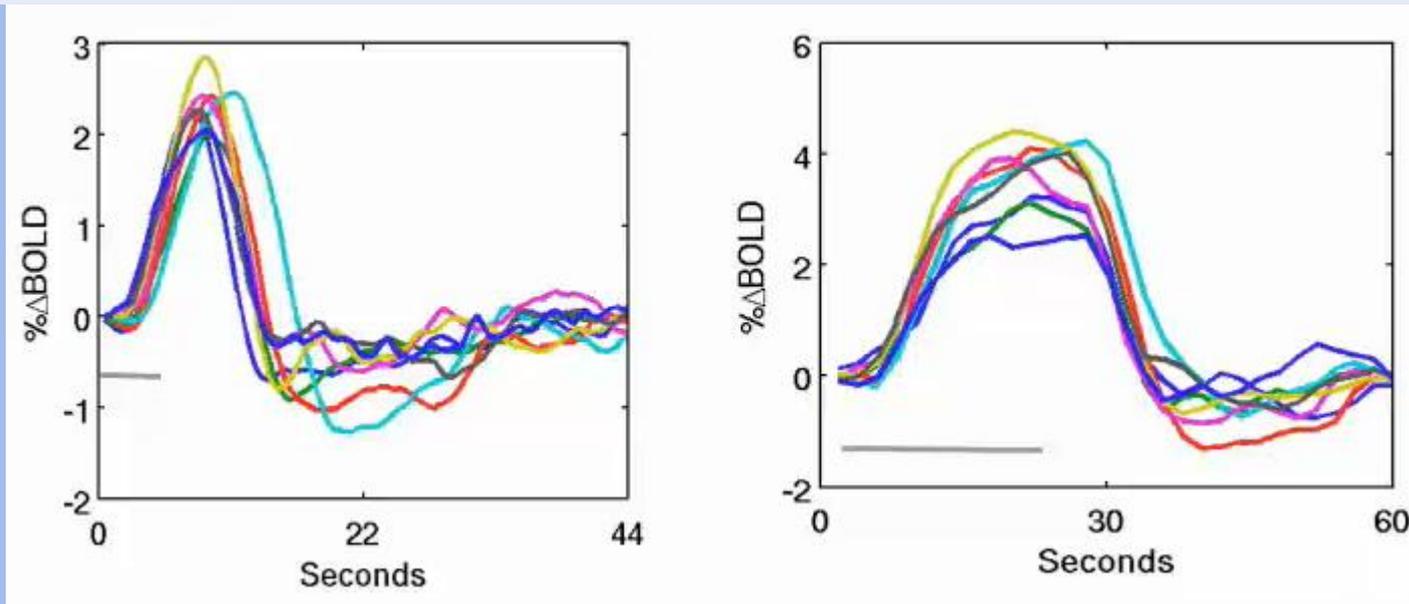


TR=300 ms

TE=30ms

64 x 64; 3.6 mm<sup>3</sup>

# Variability of BOLD signal



Healthy subjects/ same acquisition time/same age/male/same.....

- Vascular origin of variability ?
- Origin of variability due to different Neuronal Activation ?
- etc.

# fMRI sources of variance

-  Sequence  
Susceptibility
-  Drug/Coffee/Nicotine/ etc.  
Circadian rhythm/Time
-  Respiration  
Cardiac pulsatility in brain  
3D Motion
-  Age  
Healthy/Patient
-  (list not complete)



Task-related variability  
Trial-to-trial variability

# Understanding fMRI signal

$$BOLD_{signal} = \frac{Signal}{Noise} = \frac{\sum_{i=1}^{\infty} S_i}{\sum_{i=1}^{\infty} N_i}$$

$$BOLD_{signal} = \frac{\sigma_{Signal}}{\sigma_{Noise}}$$

**Maximize**

Signal

Minimize

Noise

# Signal to Noise Ratio (SNR)

$$SNR \propto \rho \frac{FOV_x FOV_y}{\sqrt{N_x N_y} bw} \rho_s \sqrt{N_{average}} B_0 f$$

$\rho, \rho_s$ : proton density, slice thickness

FOV: Field of view in x,y

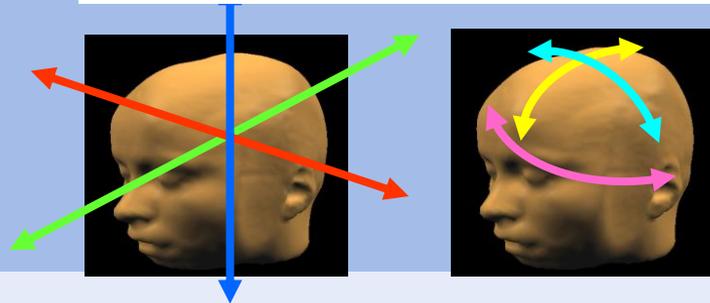
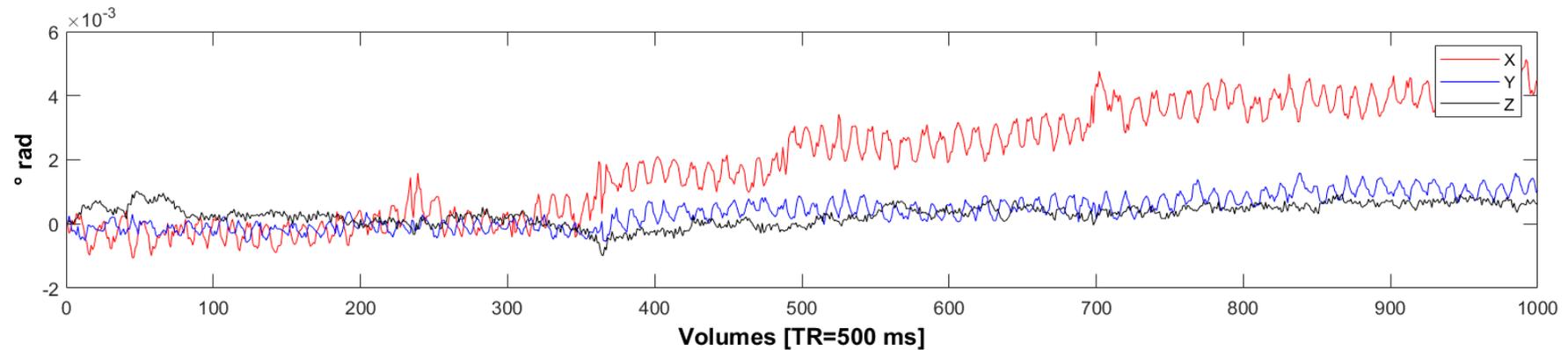
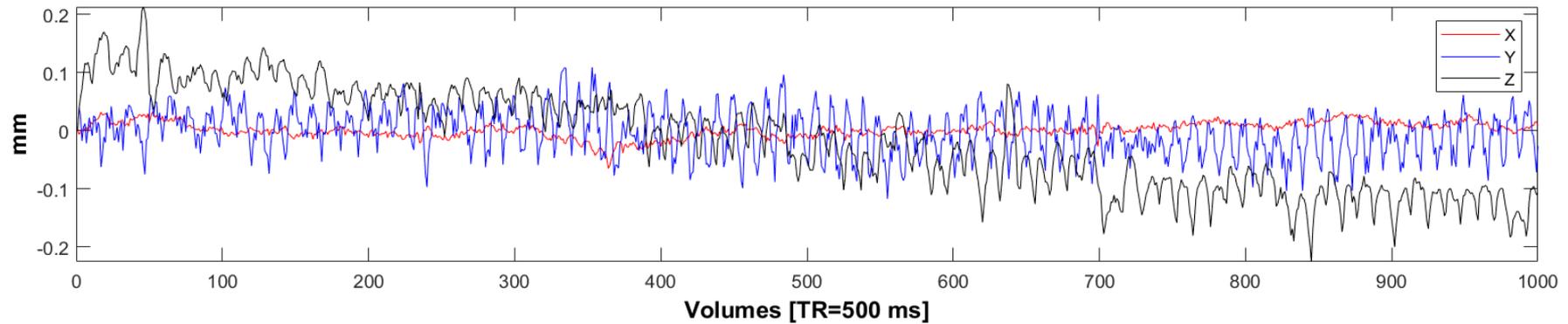
N: Number of points in x,y

bw: sampling bandwidth

$B_0$ : static magnetic field

f: sequence parameter (TR, TE, coil, etc...)

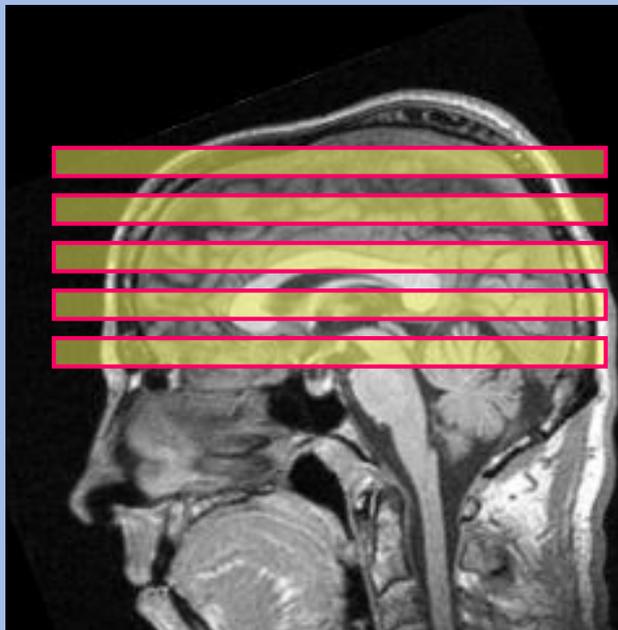
# 3D Head Motion



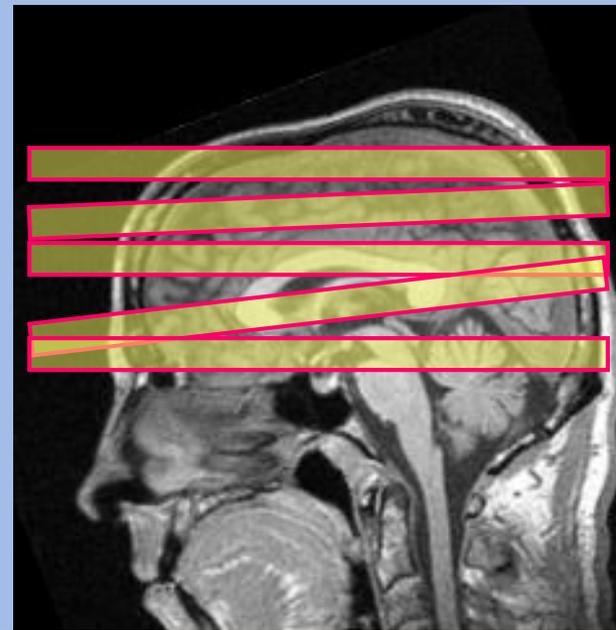
# Difficulties

> Motion is 3-dimensional

expected acquisition:



slices are actually acquired like this...



## Estimating the motion parameters – from data

*Function to minimize*

- > Coefficient of variation of Ratio

$$E \equiv \frac{\sigma_R}{\mu_R} \quad R \equiv \frac{T(\text{image}_i)}{\text{image}_{\text{base}}}$$

*Woods, et al.*

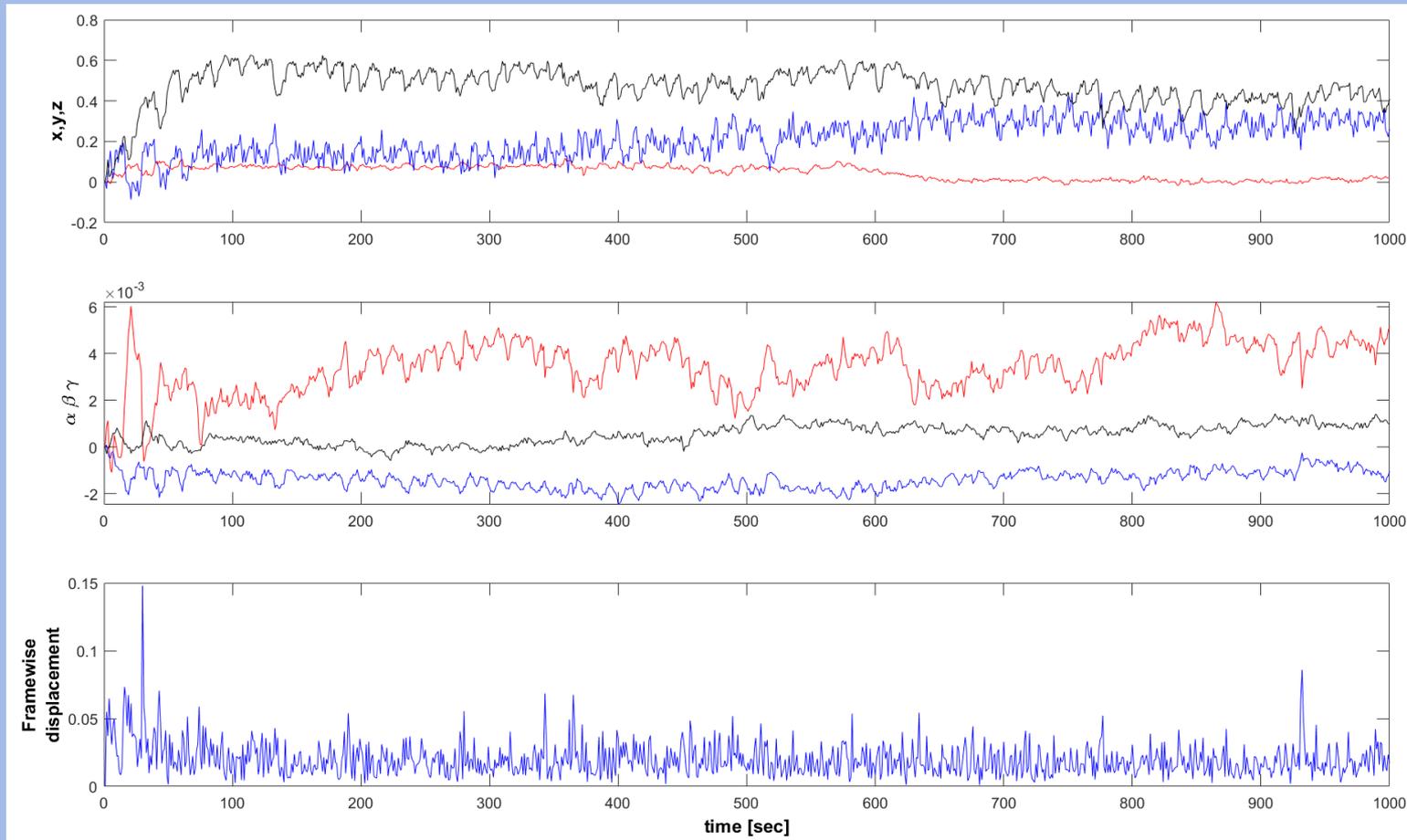
- > Squared difference

$$E \equiv \sum [T(\text{image}_i) - \text{image}_{\text{base}}]^2$$

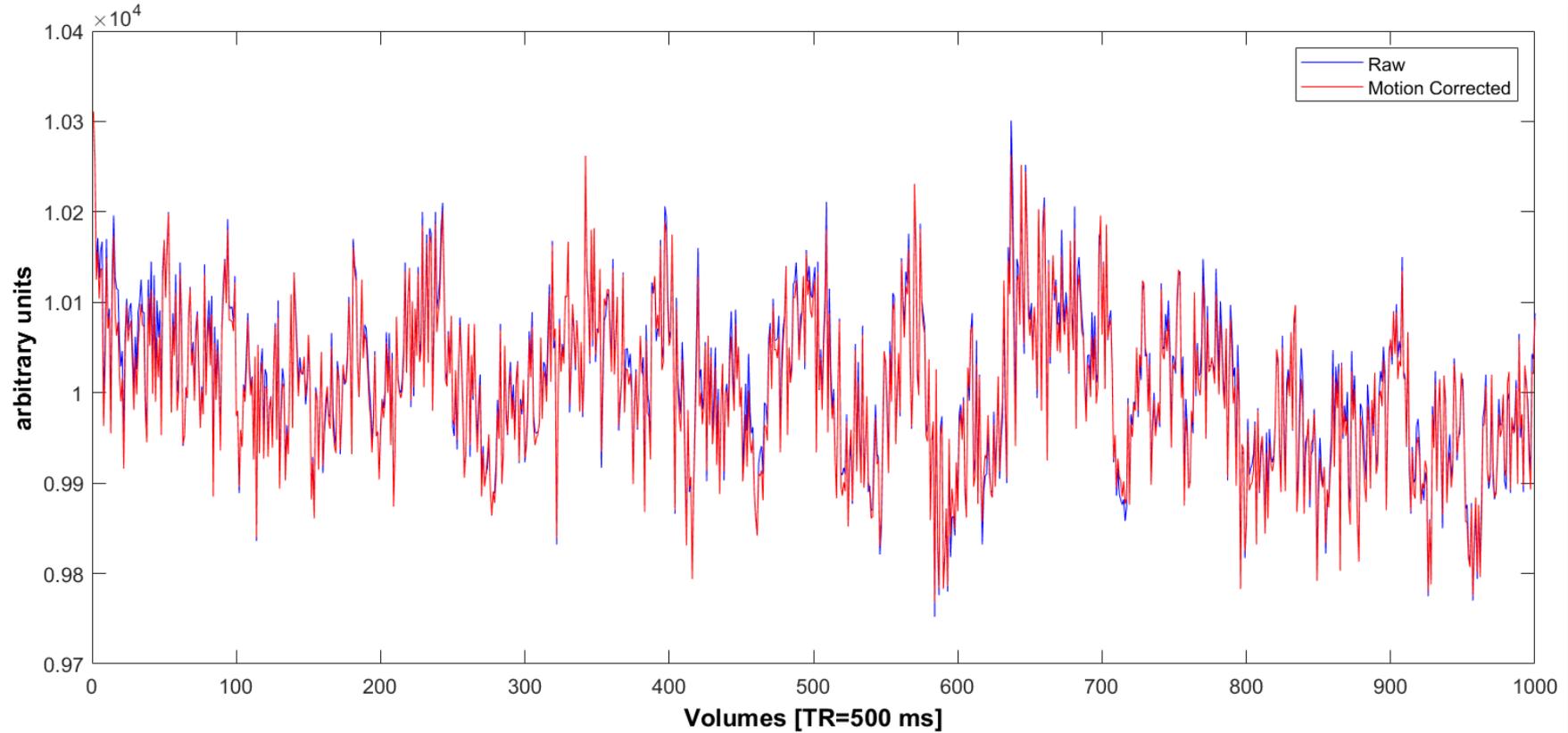
*Hajnal, et al.; Eddy, et al.*

# Frame wise displacement

$$FD = \left| \frac{\partial x}{\partial t} \right| + \left| \frac{\partial y}{\partial t} \right| + \left| \frac{\partial z}{\partial t} \right| + \left| \frac{\partial \alpha}{\partial t} \right| + \left| \frac{\partial \beta}{\partial t} \right| + \left| \frac{\partial \gamma}{\partial t} \right|$$

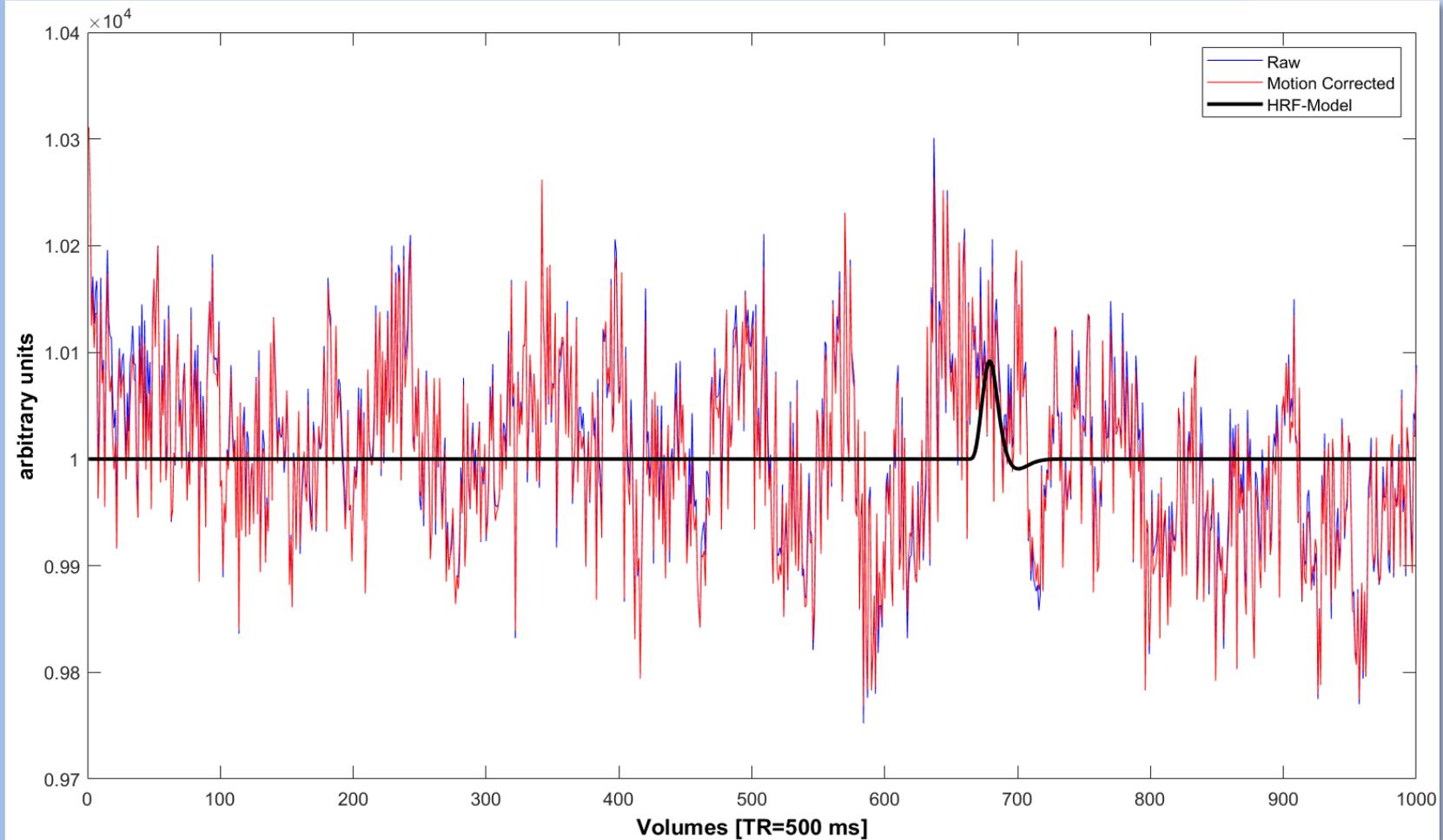


# Understanding fMRI signal



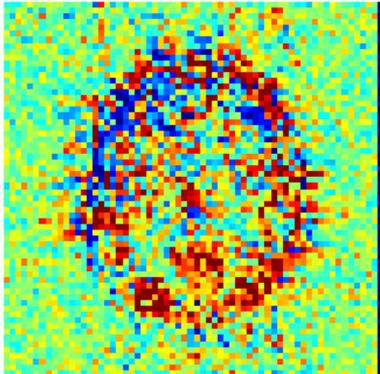
Small motion

# Understanding fMRI signal

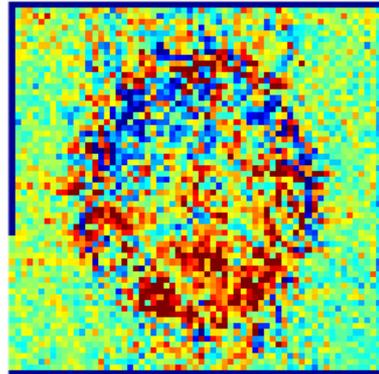


# 3D Head Motion

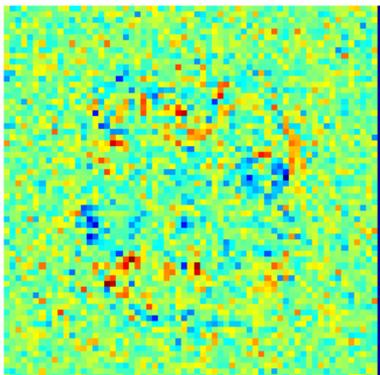
Raw: strong motion



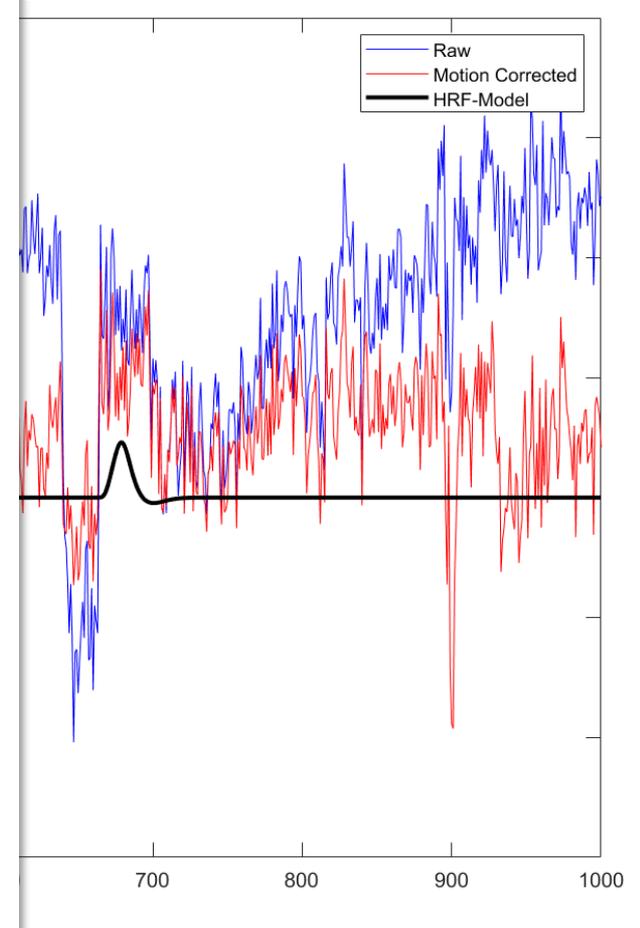
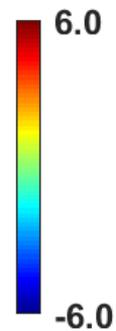
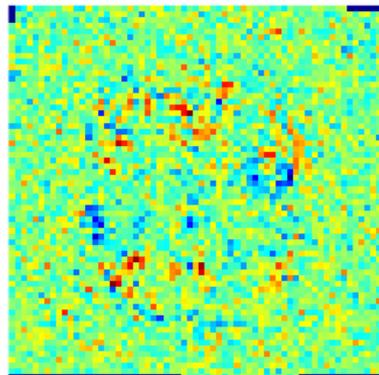
Corrected: strong motion



Raw: low motion

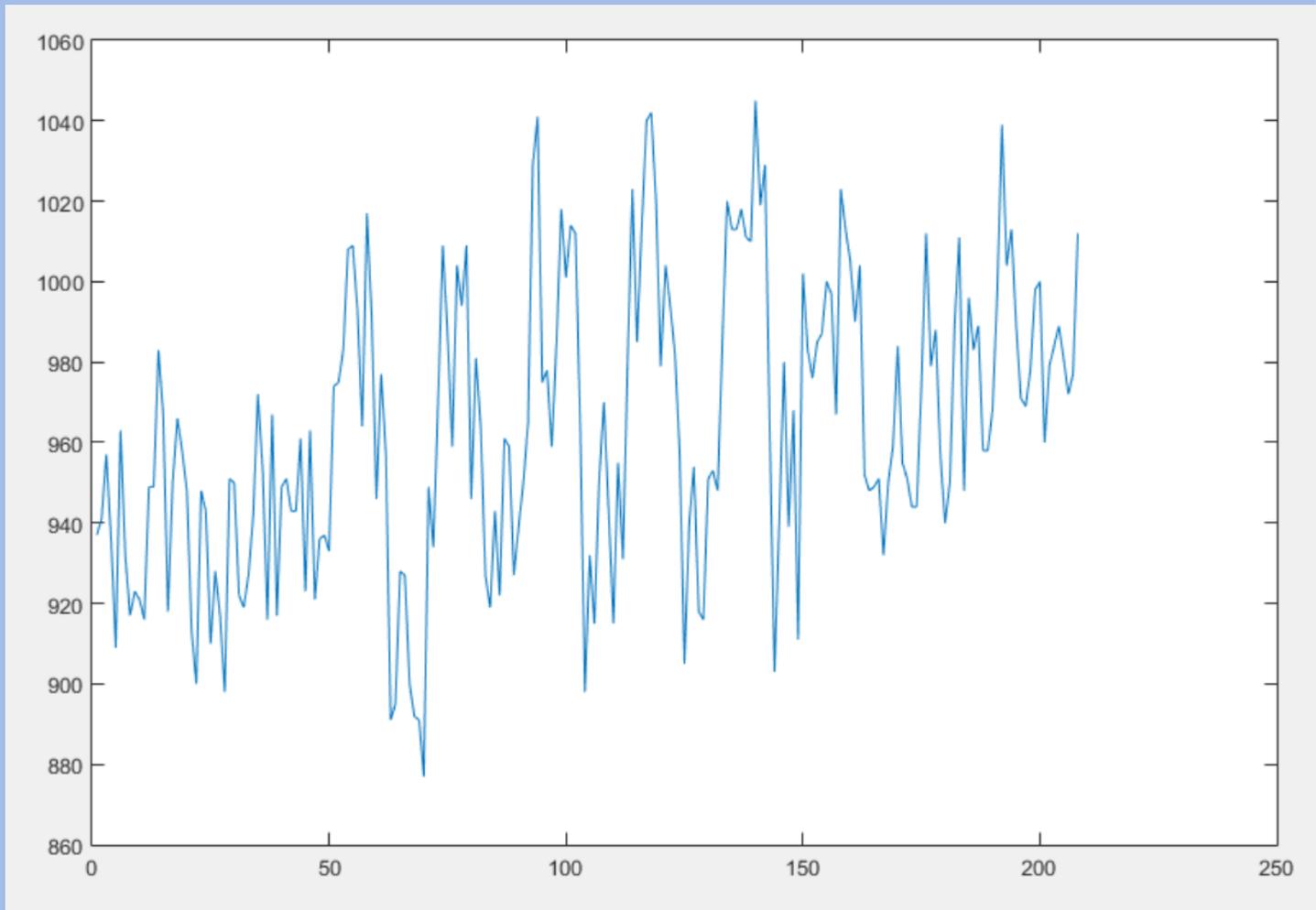


Corrected: low motion

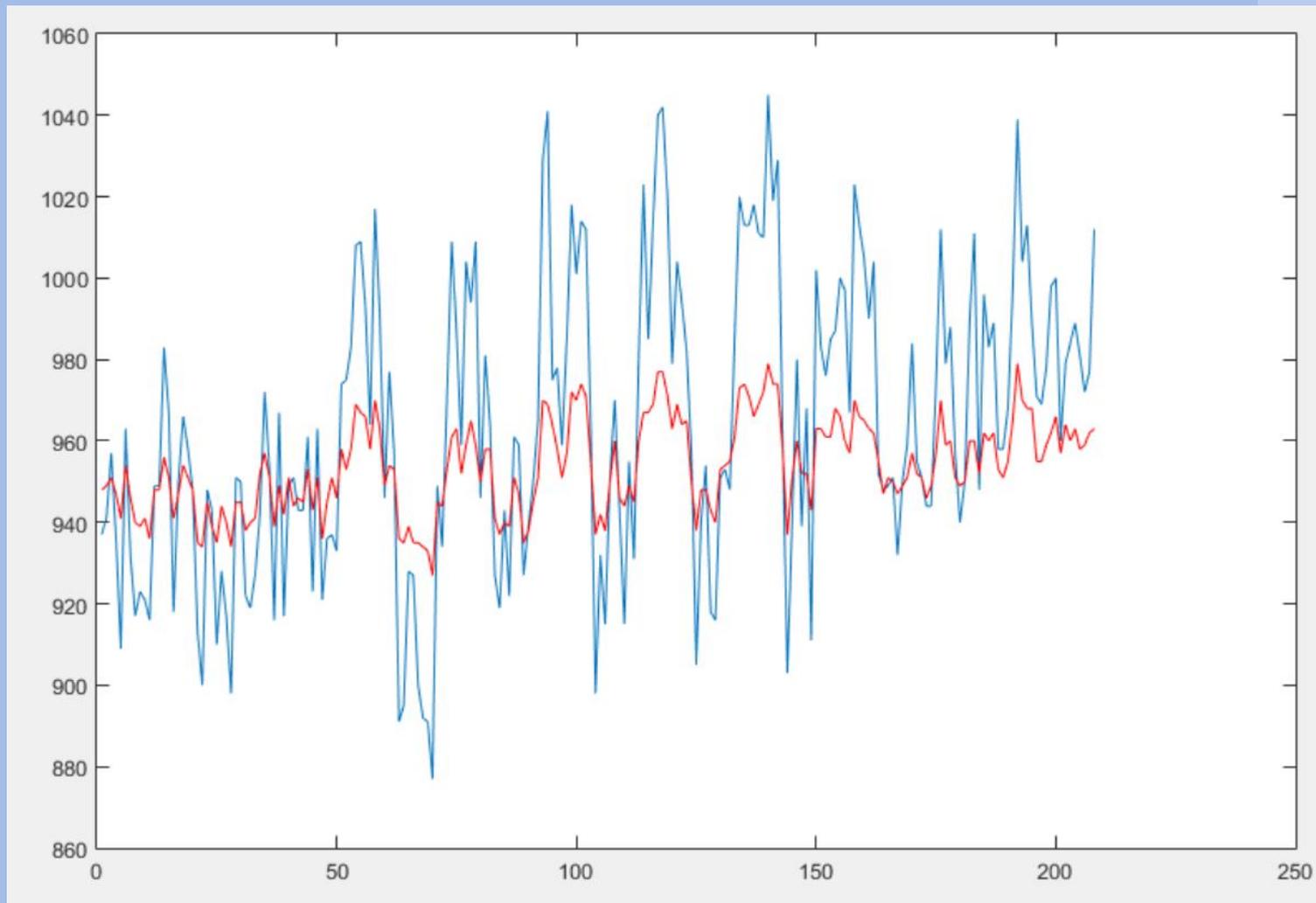


large motion

# Understanding fMRI signal

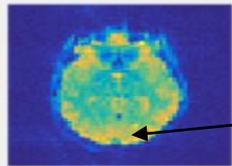


# Understanding fMRI signal

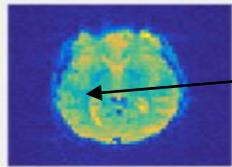


Global Signal regression

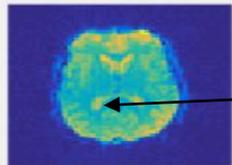
# Understanding fMRI signal



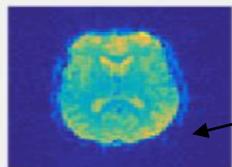
V1



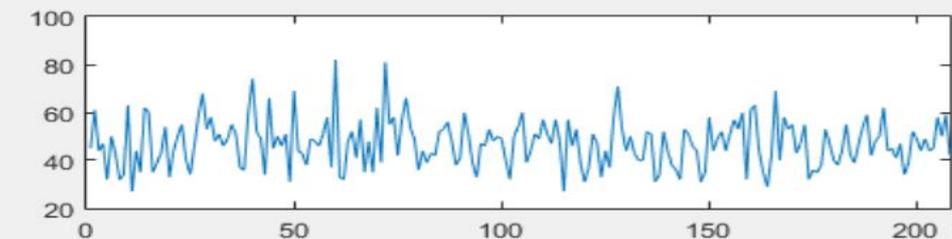
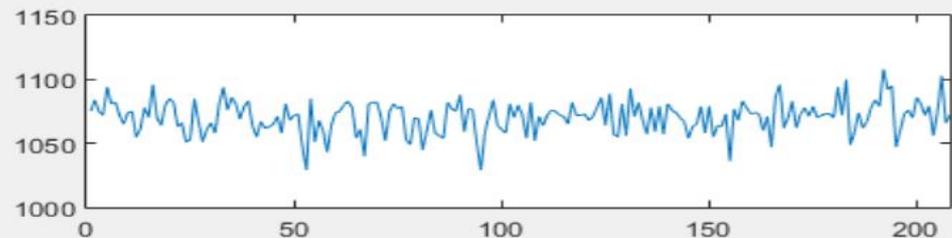
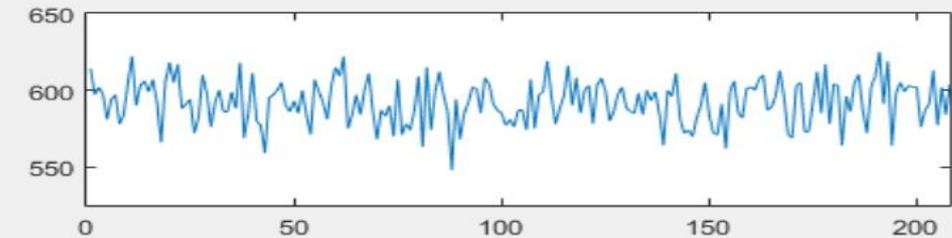
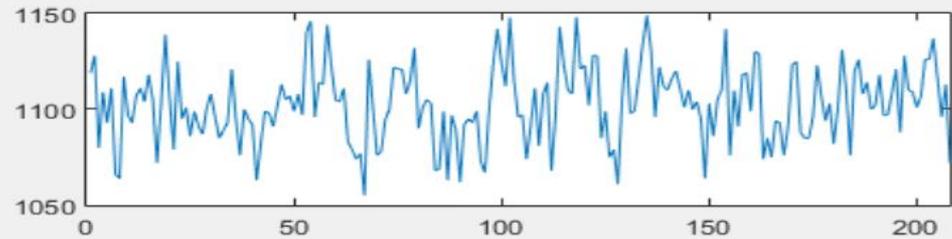
WM



CSF



Outside  
brain



# Understanding fMRI signal

$$BOLD_{signal} = \beta_0 + X_{hrf} * \Theta + \varepsilon$$

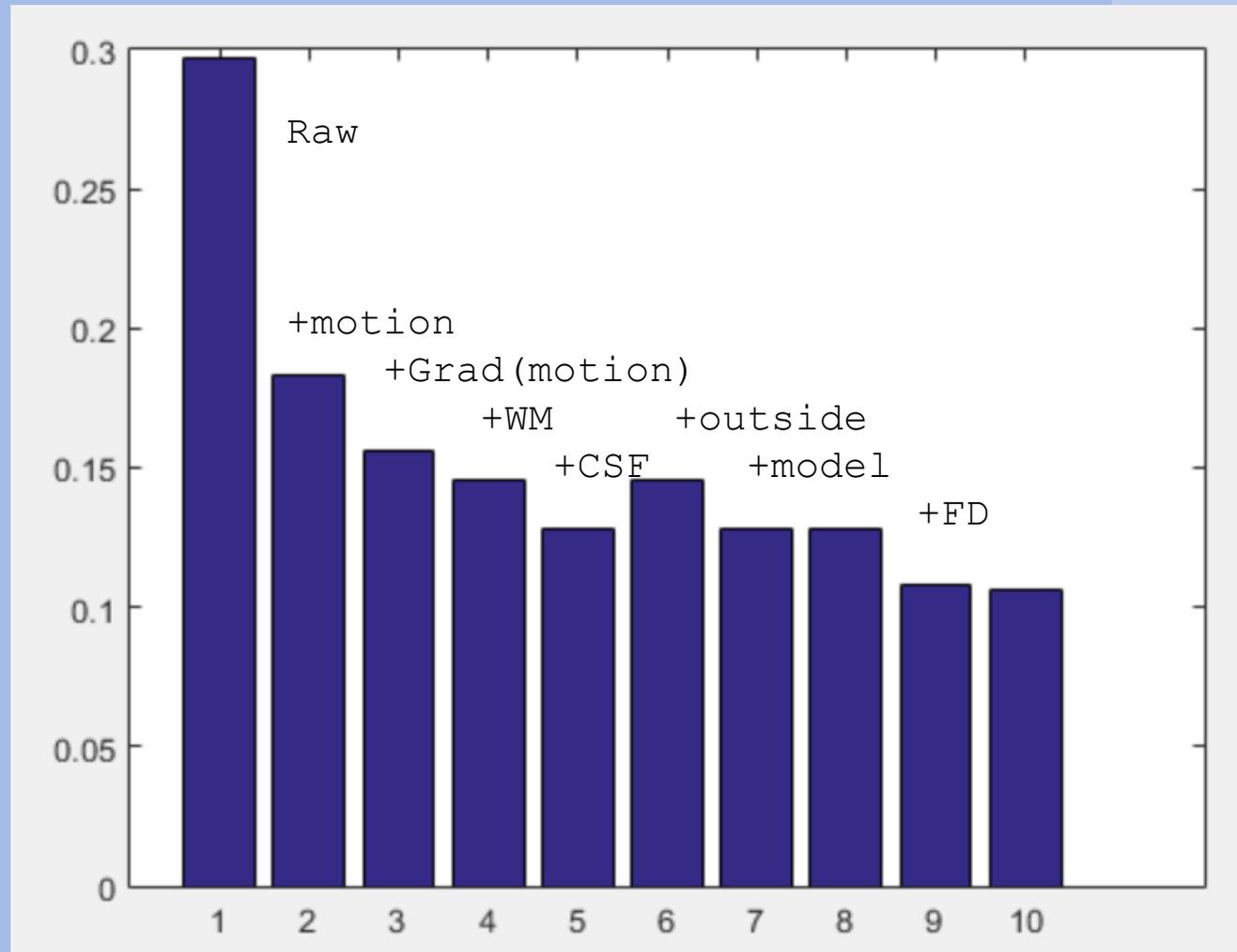
```
[b,dev,stats1] = glmfit(Bold_th,Bold_measure);
```

$$BOLD_{signal} = \beta_0 + X_{hrf} * \Theta + S_{WM} + S_{CSF} + S_{GM} + \dots + \varepsilon$$

```
[b,dev,stats2] = glmfit(  
    [mot dmot fd wm csf n0 Bold_th],Bold_measure);
```

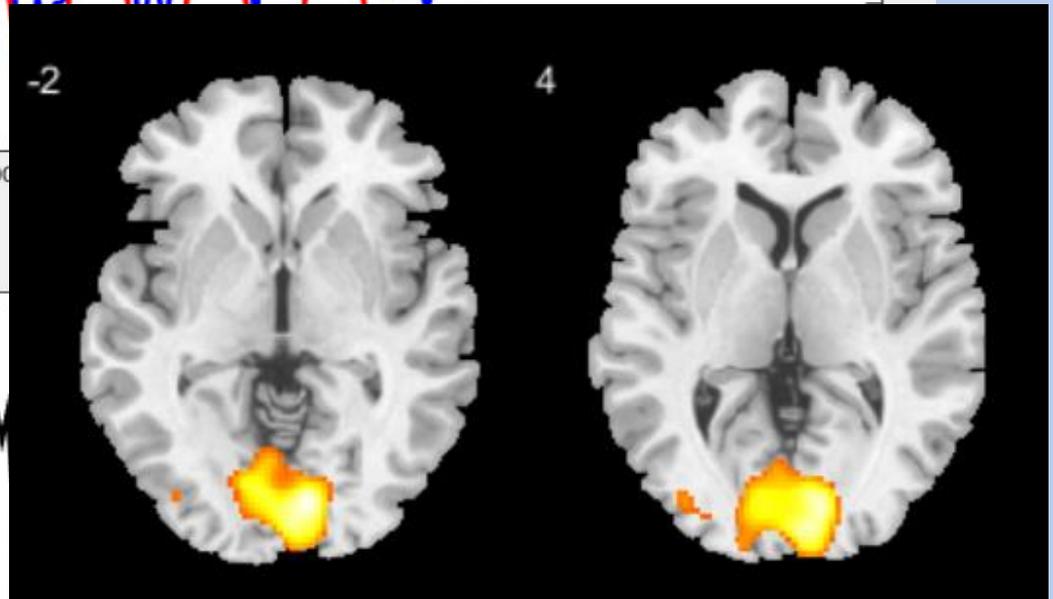
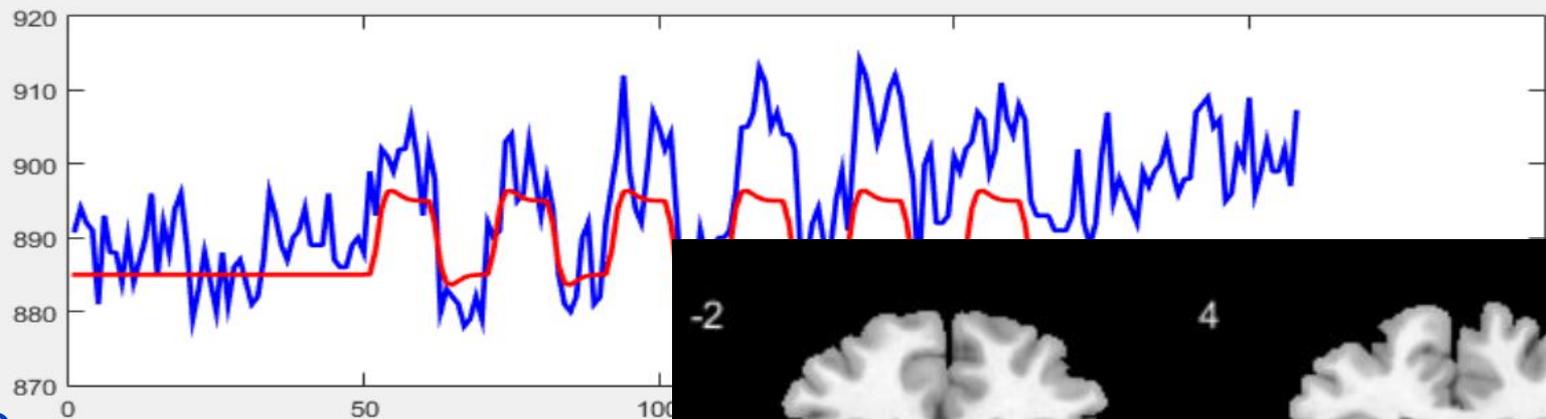
# Understanding fMRI signal

Variance



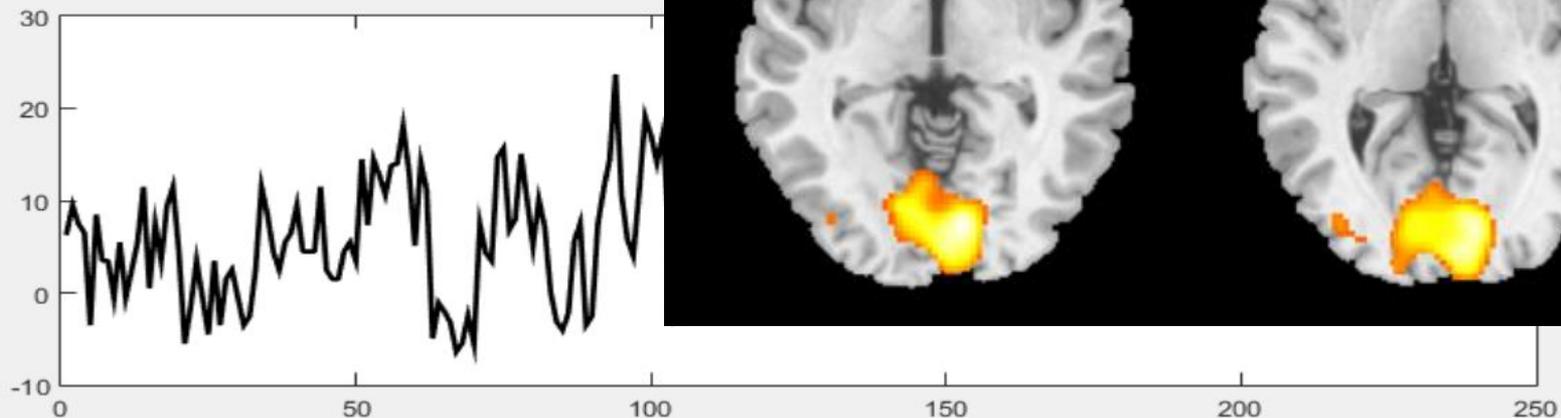
# Understanding fMRI signal

$$BOLD_{signal} = \beta_0 + X_{hrf} * \Theta + S_{WM} + S_{CSF} + S_{GM} + \dots + \varepsilon$$



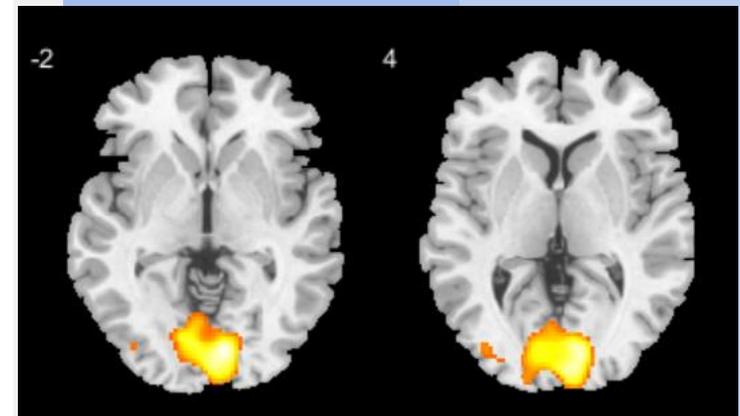
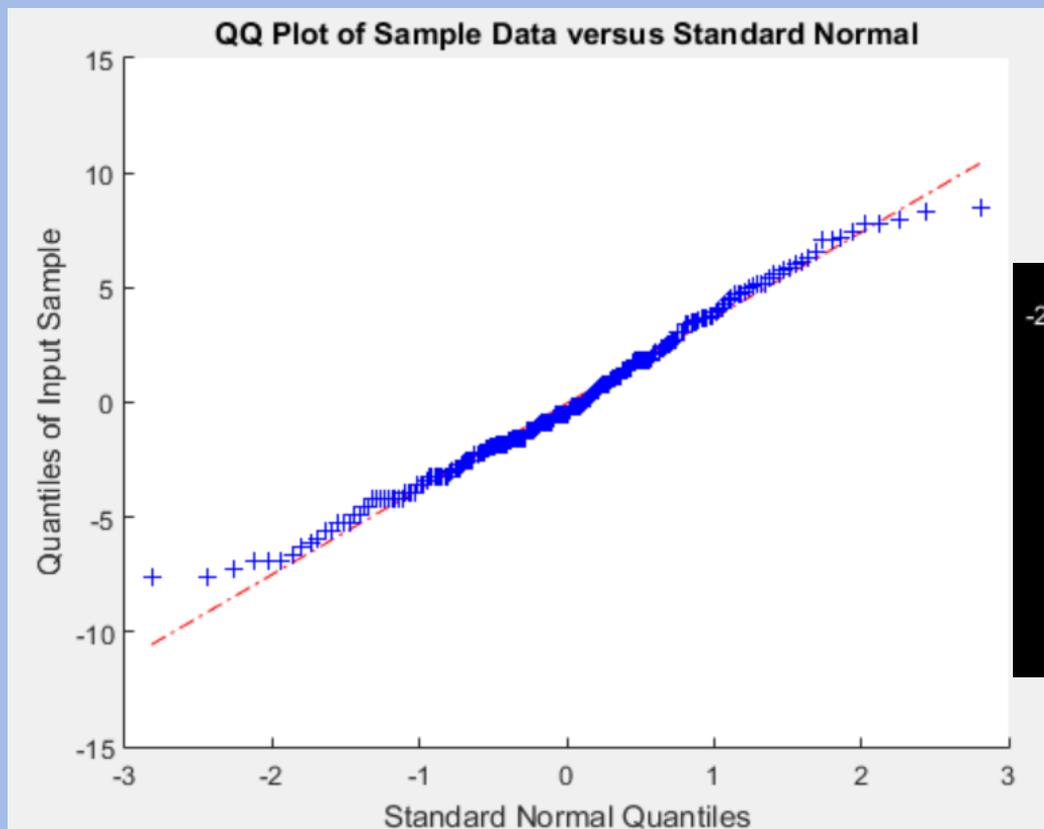
BOLD  
Model

Error



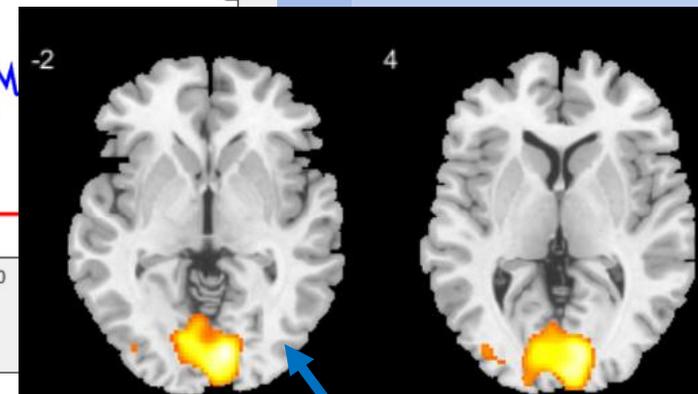
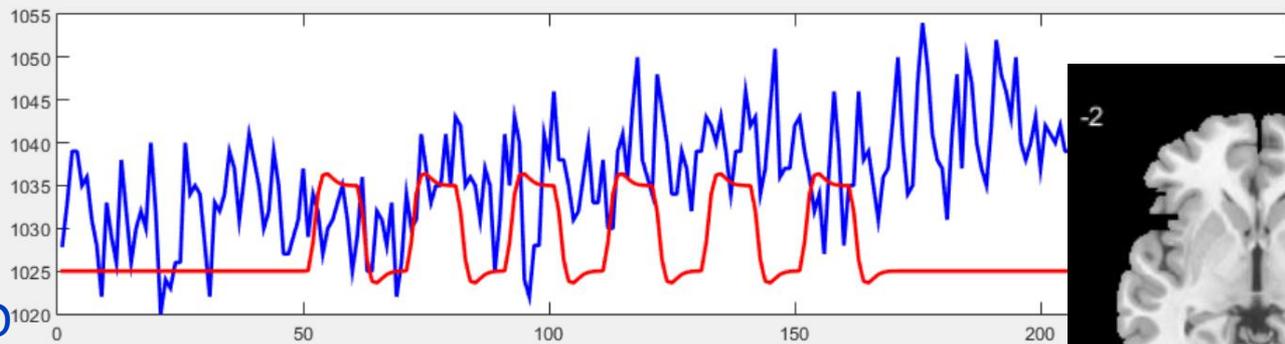
# Understanding fMRI signal

$$BOLD_{signal} = \beta_0 + X_{hrf} * \Theta + S_{WM} + S_{CSF} + S_{GM} + \dots + \varepsilon$$



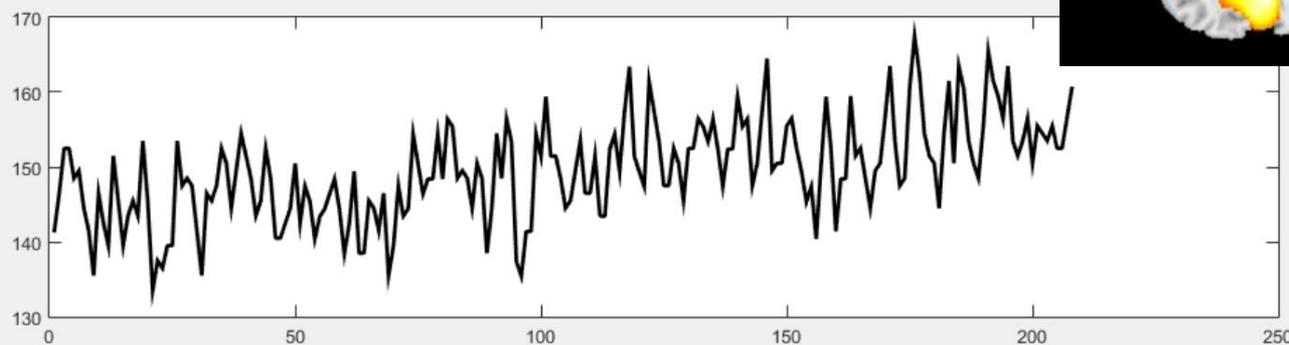
# Understanding fMR signal

$$BOLD_{signal} = \beta_0 + X_{hrf} * \Theta + S_{WM} + S_{CSF} + S_{GM} + \dots + \varepsilon$$



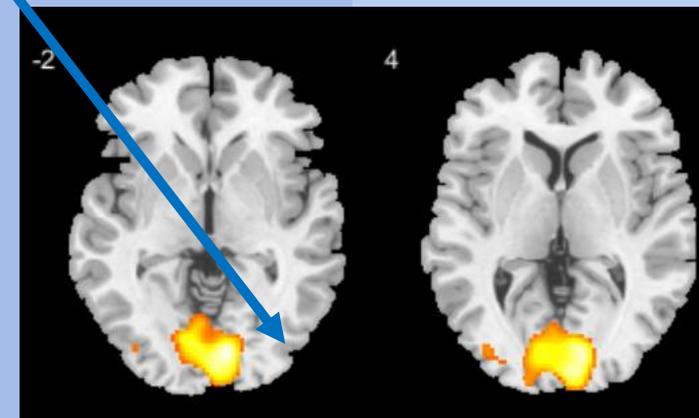
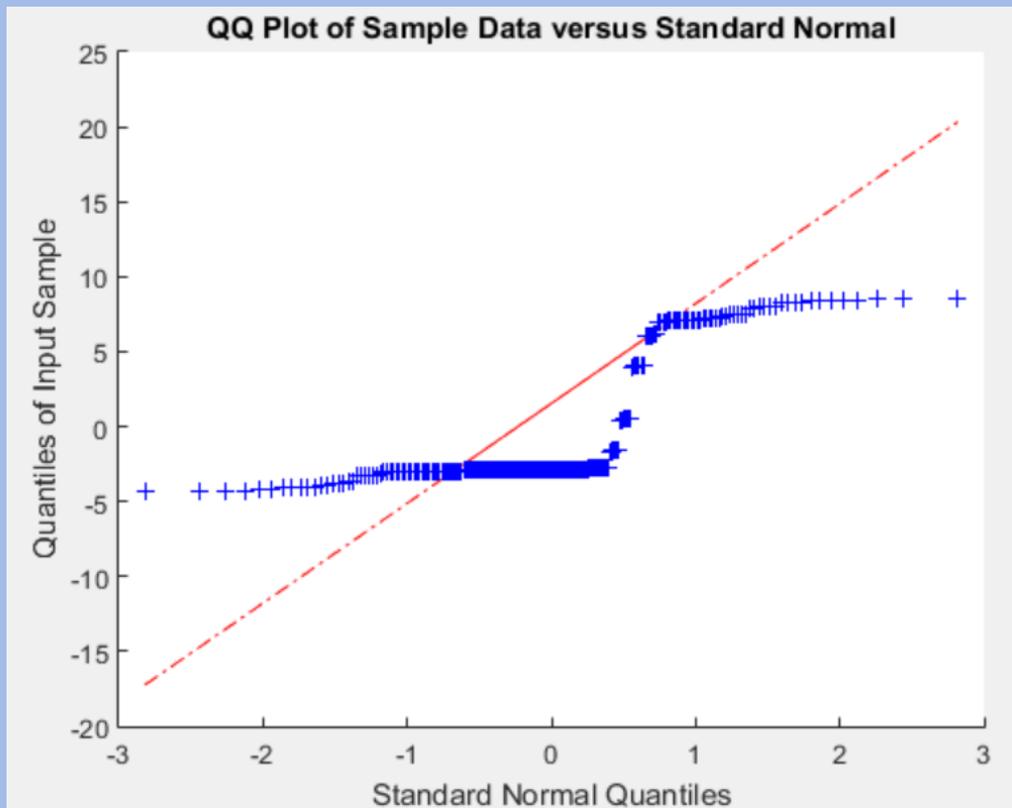
BOLD  
Model

Error

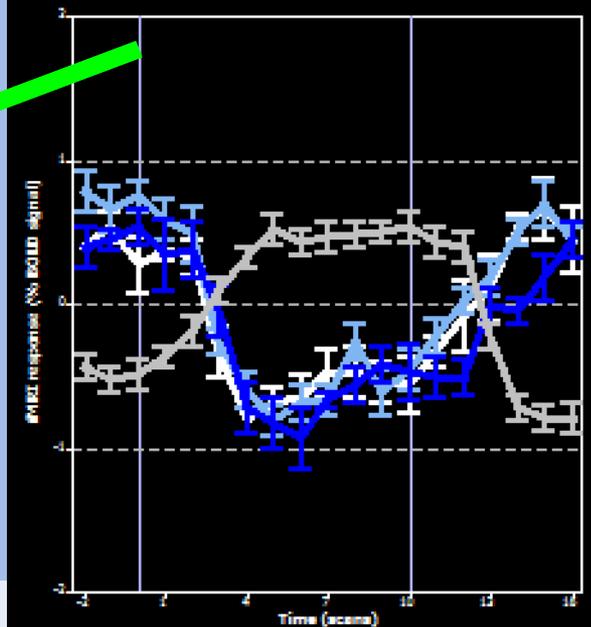
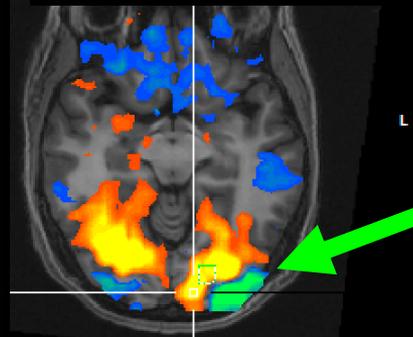
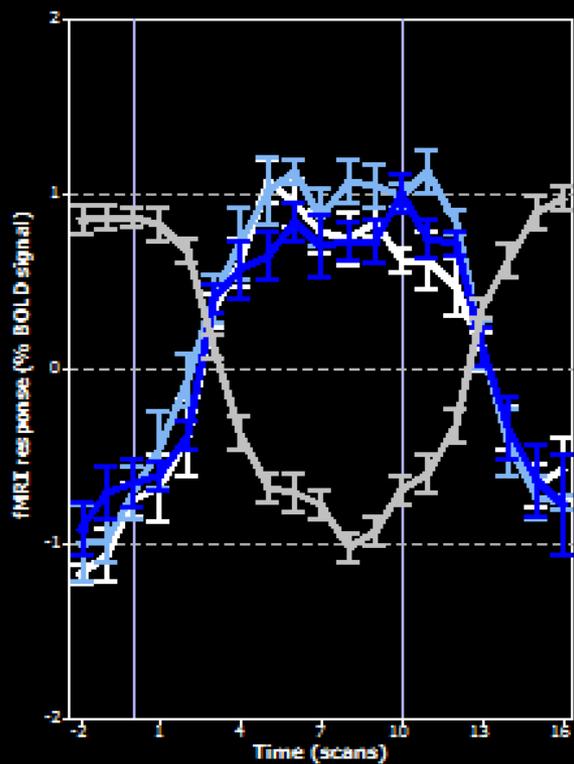
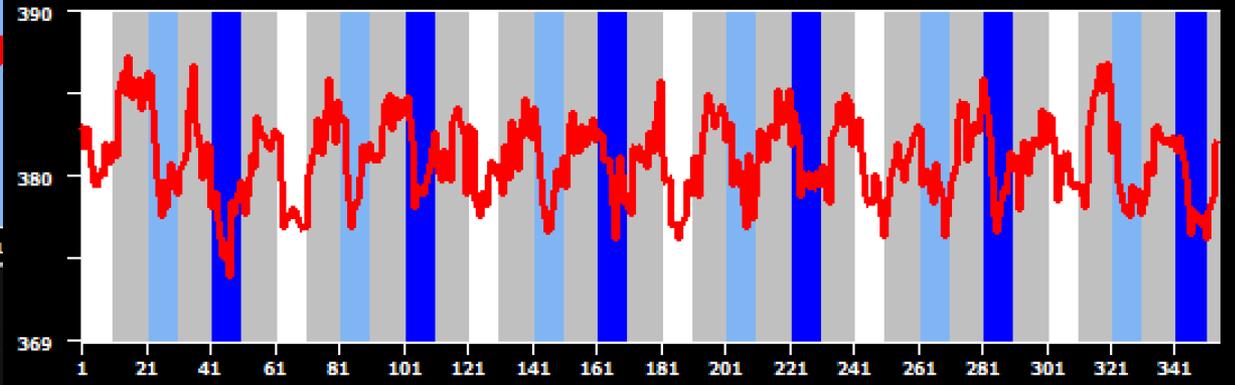
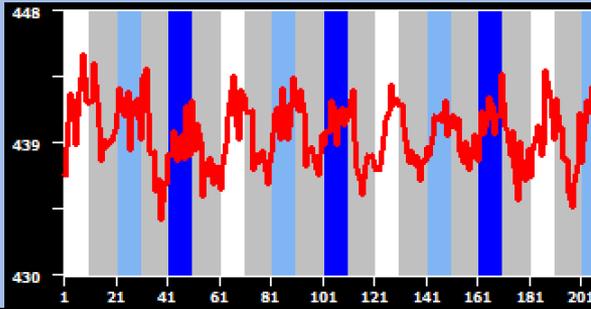


# Understanding fMRI signal

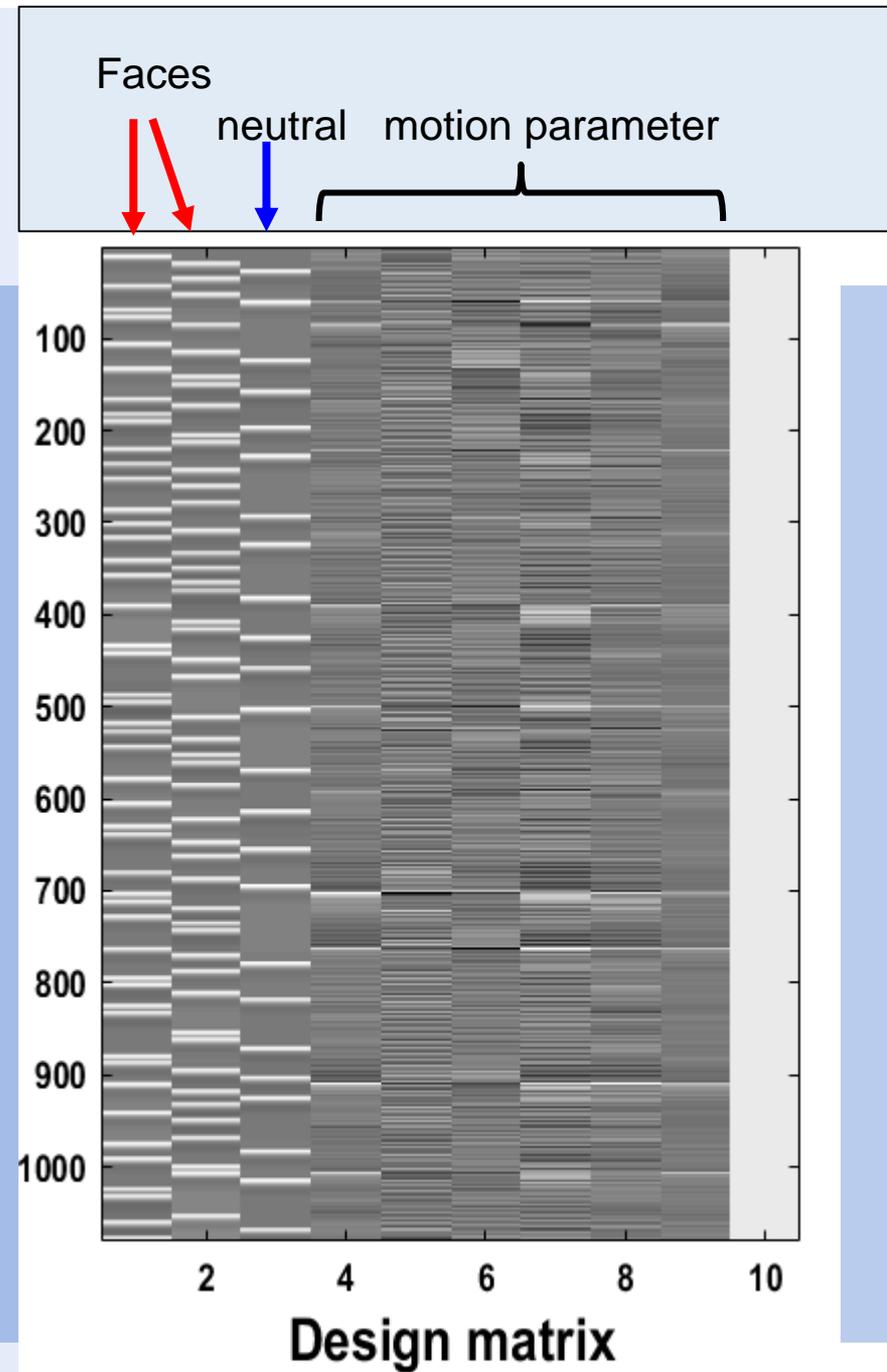
$$BOLD_{signal} = \beta_0 + X_{hrf} * \Theta + S_{WM} + S_{CSF} + S_{GM} + \dots + \varepsilon$$



# Baseline in fMRI signal



# Event related fMRI

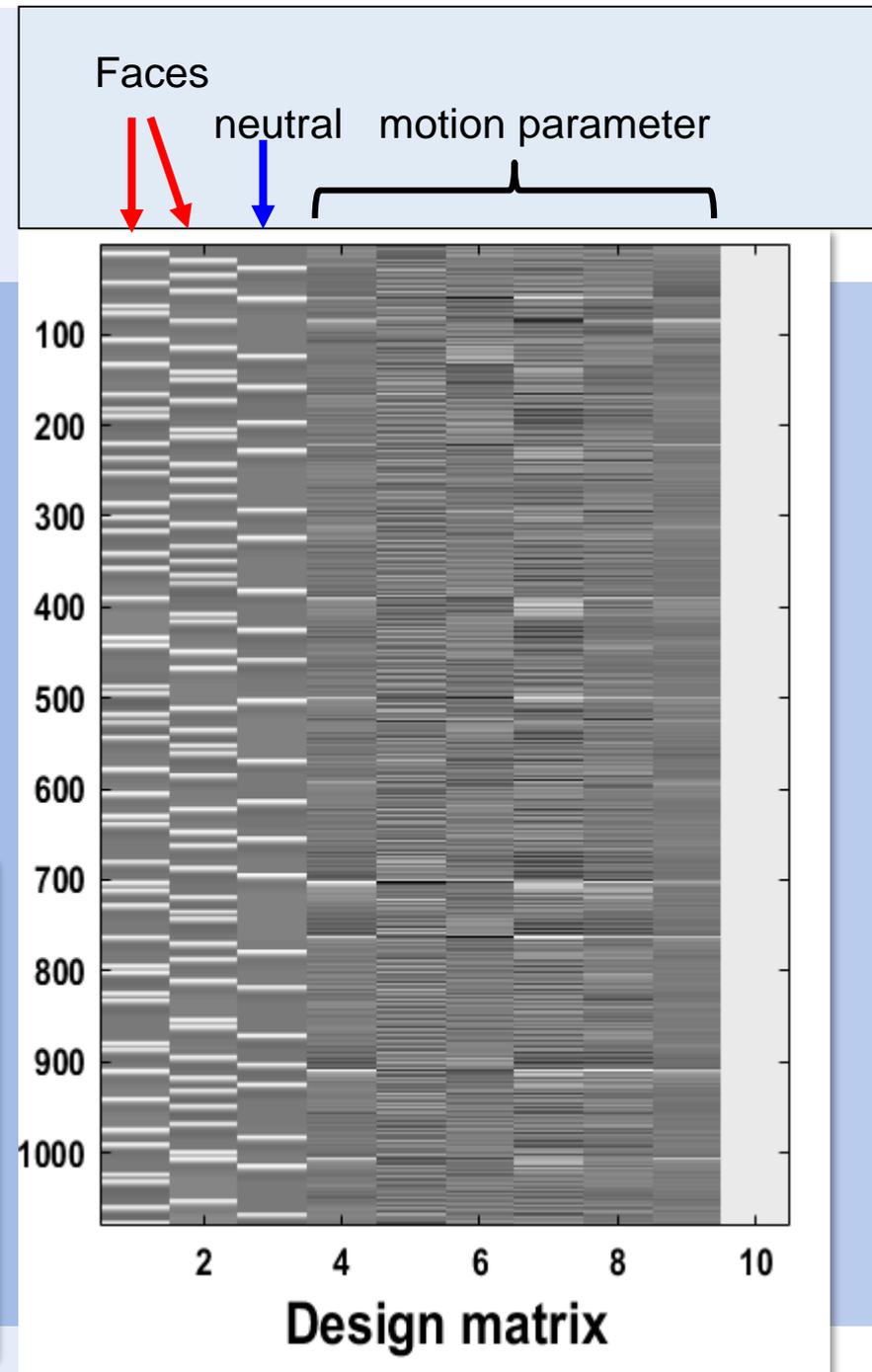
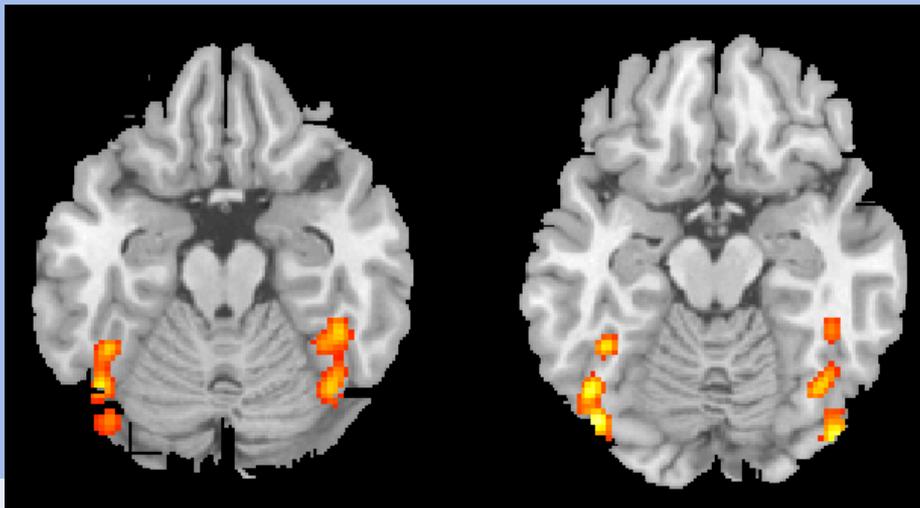


# Event related fMRI

Contrast: faces vs neutral

Question: **Where ?**

Expectation: fusiform gyrus  
/fusiform face area



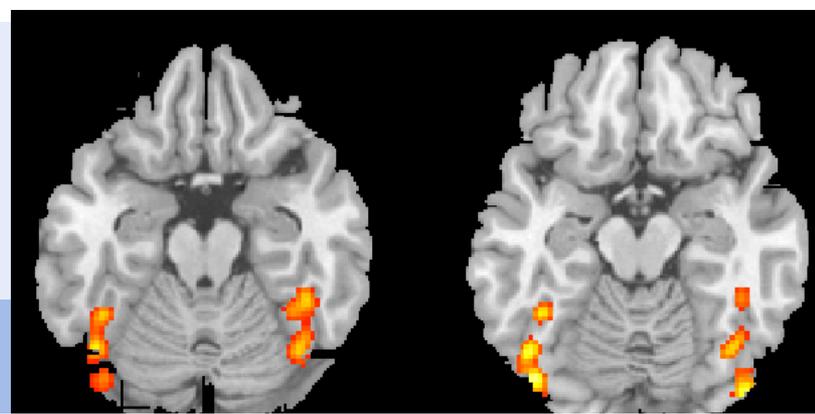
# General Linear Model

$$BOLD_{signal} = b_0 + b_1 * X_{stimulus} + e$$

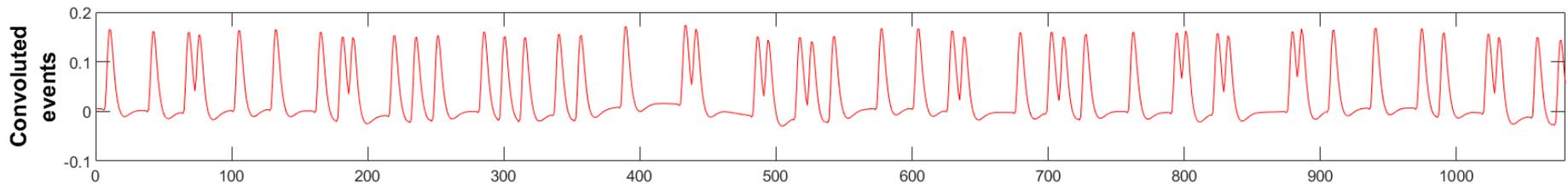
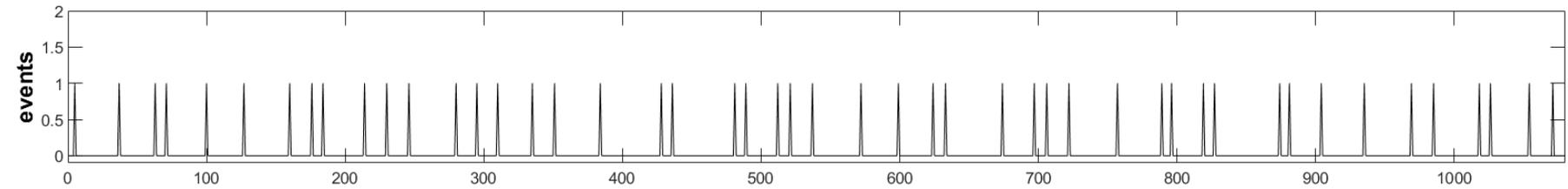
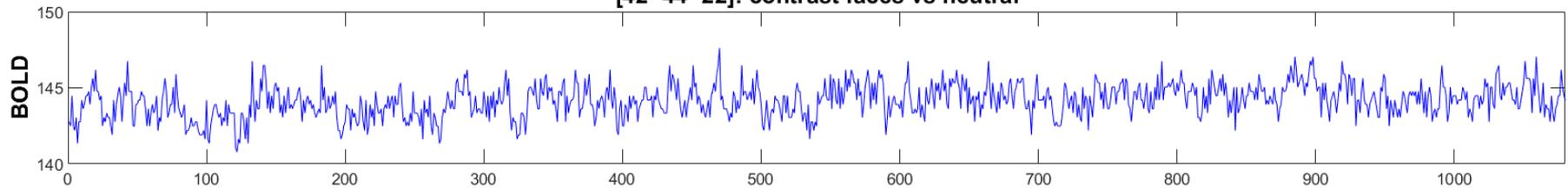
*events*

```
[betas, dev, stats] = glmfit(Bold_model , Bold_signal);
```

# Event related fMRI

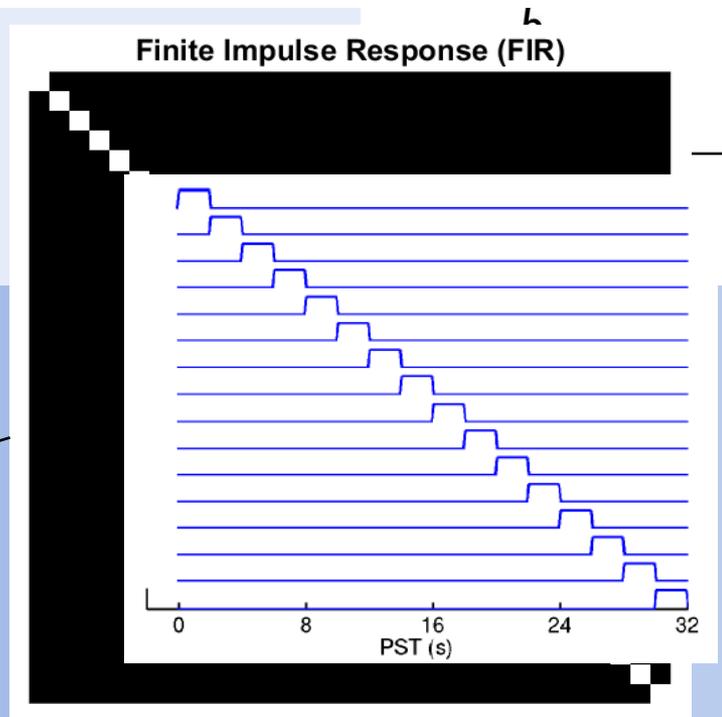


[42 -44 -22]: contrast faces vs neutral

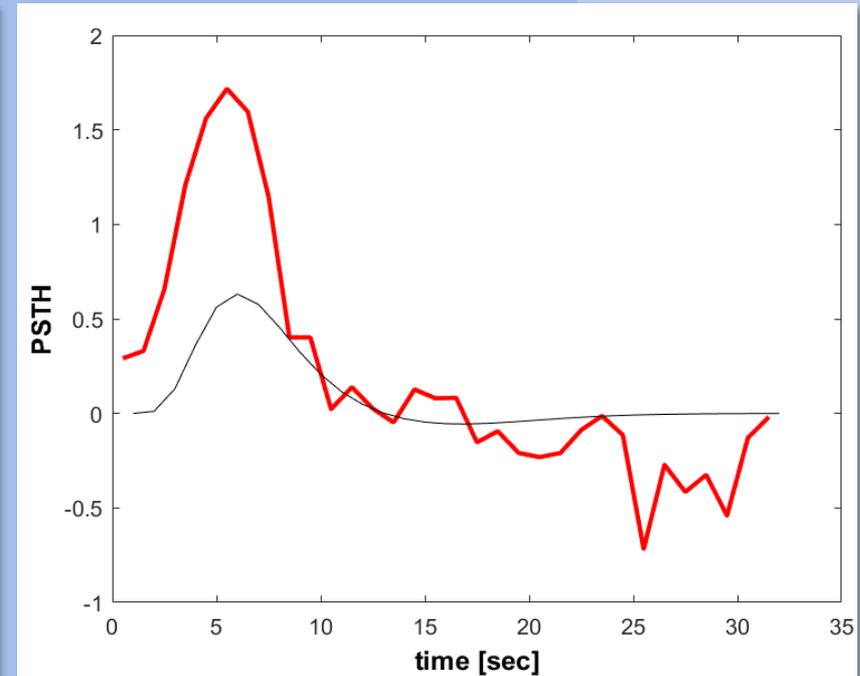
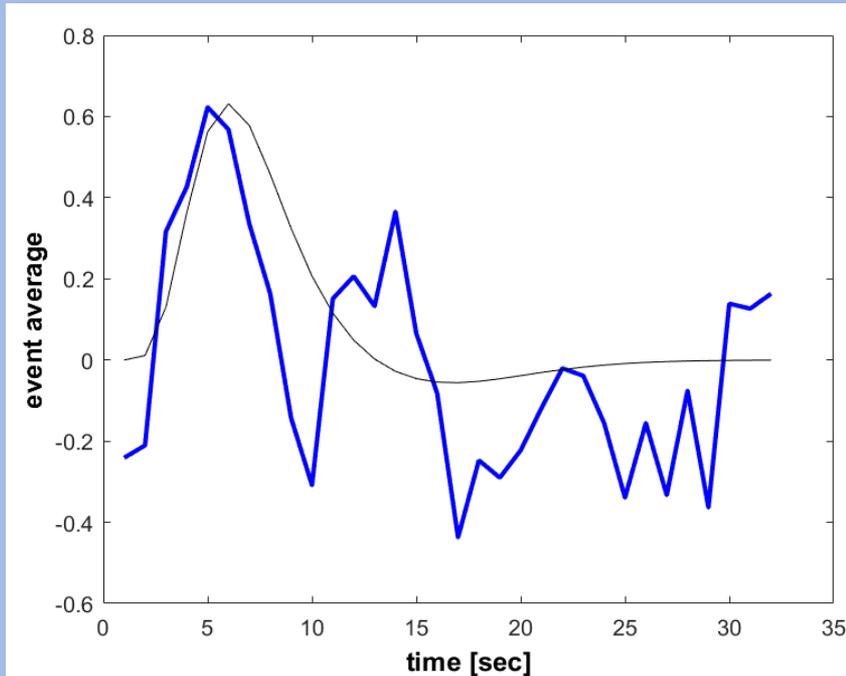


# Event related fMRI

```
N=size(Events,1)
FIR=eye(32);
estimated_HRF=zeros(N,32);
for n=1:N
    Twindow=single(yBOLD(Events(n)+2:Events(n)+33));
    for time=1:32
        [b,de,st]=glmfit(FIR(time,:),Twindow);
        estimated_HRF(n,time)=b(2);
    end
end
estimated_HRF=mean(estimated_HRF,1);
```

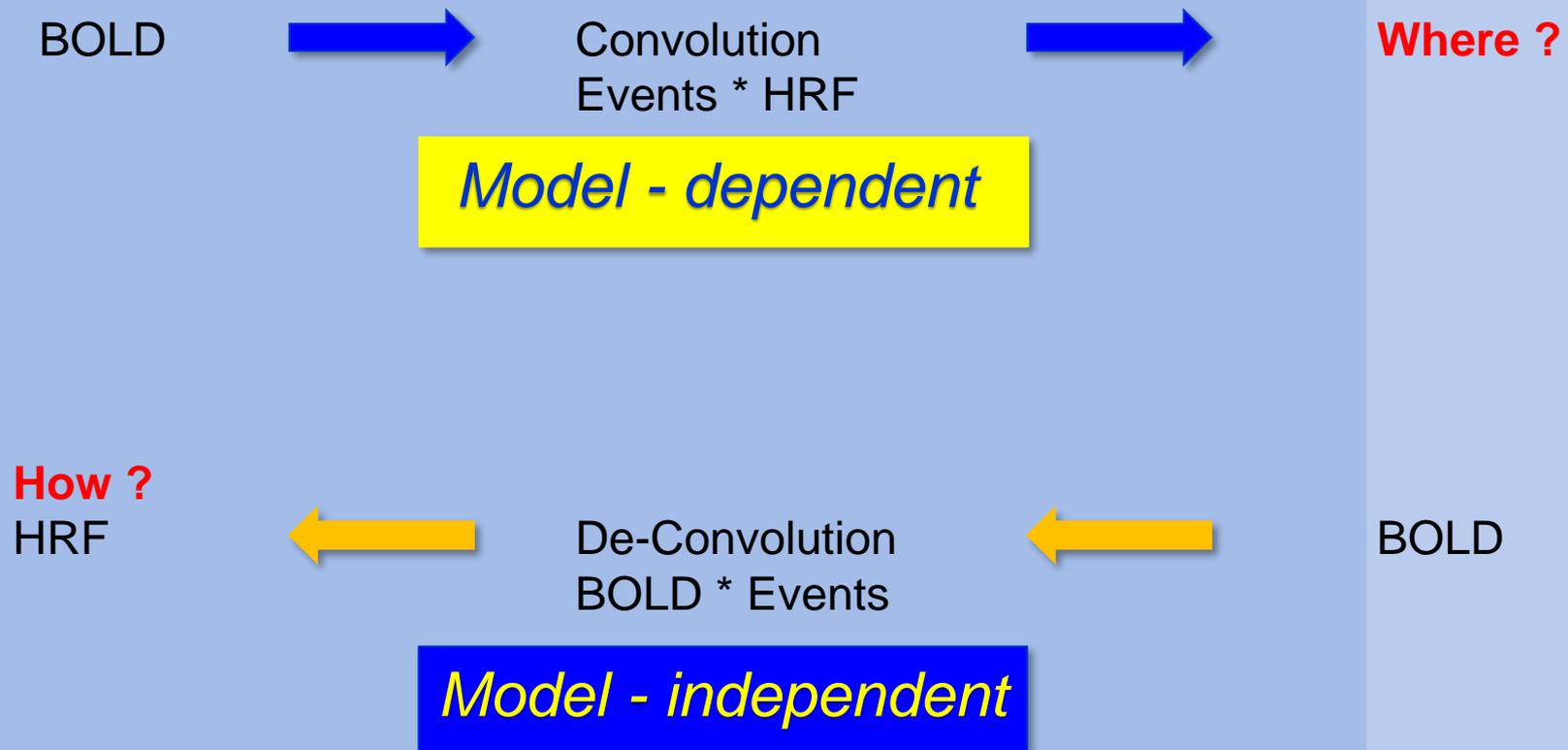


# How sure can you be ?

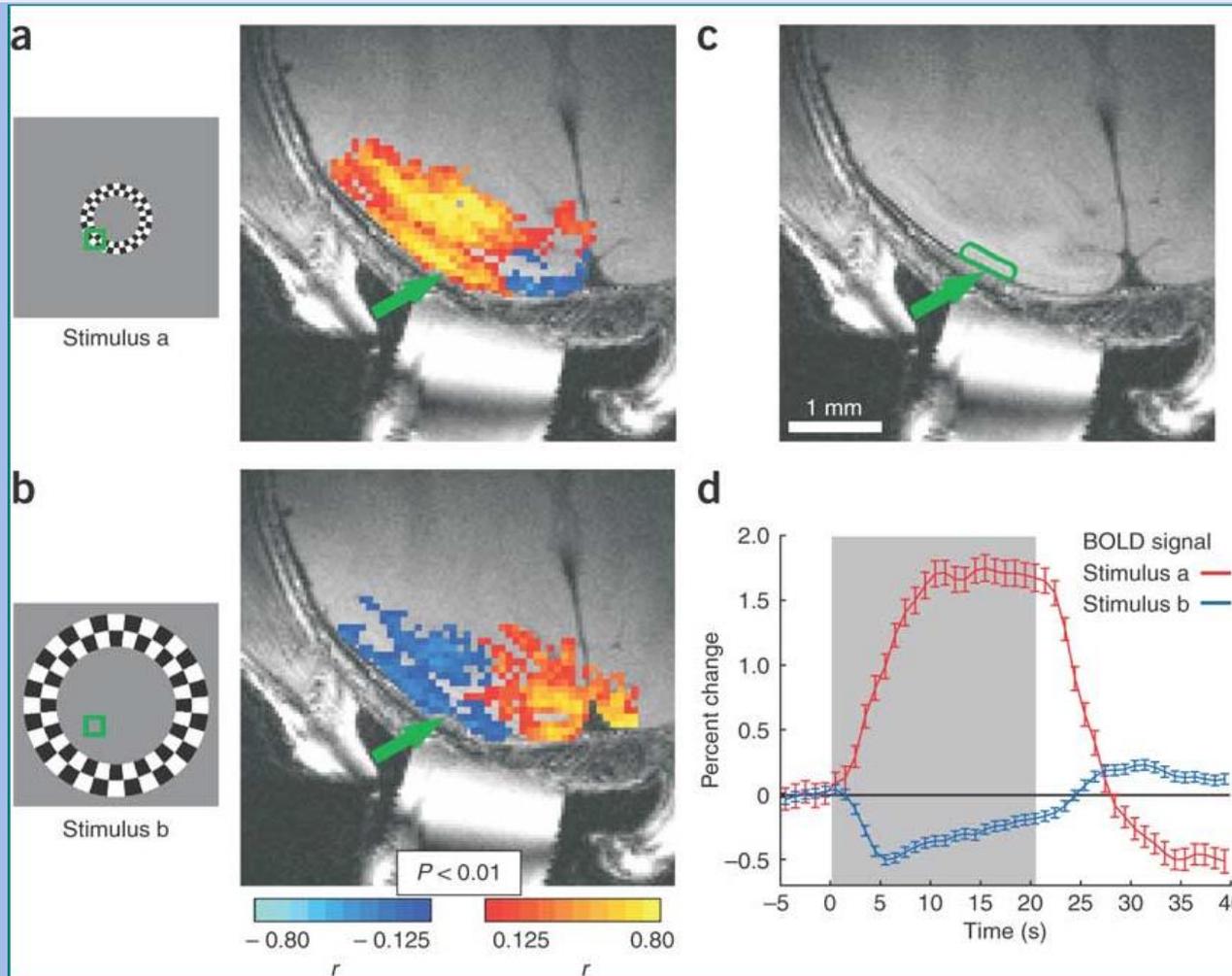


estimated\_HRF

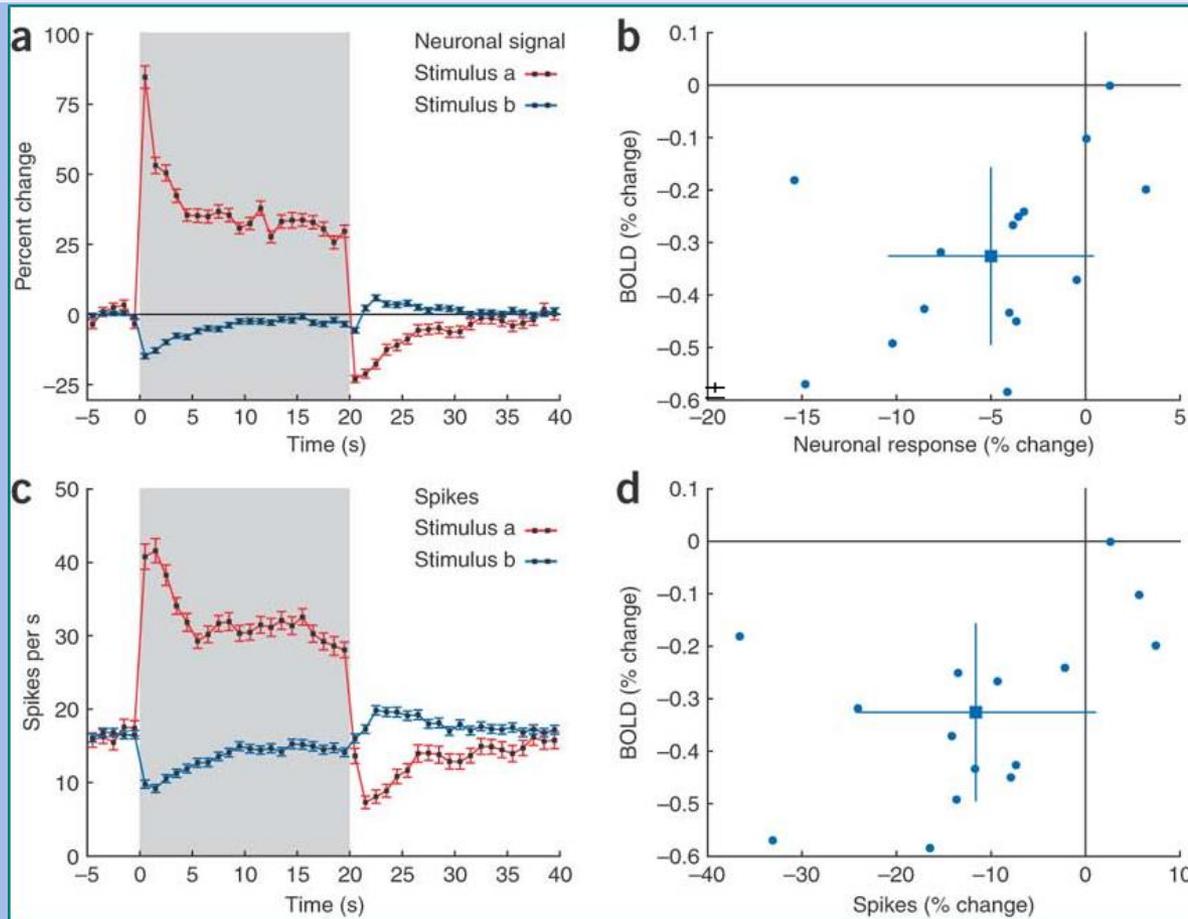
# Event related fMRI: check results



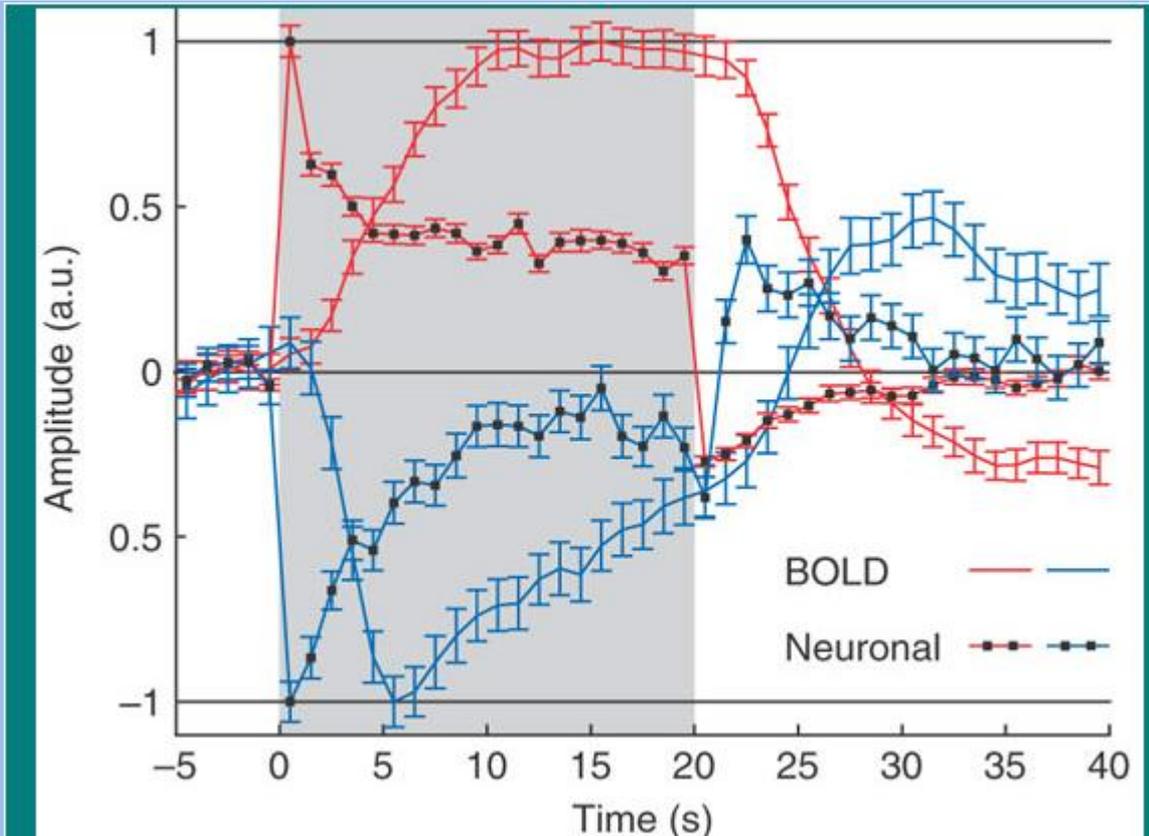
# Negative BOLD: challenging



# Negative BOLD: challenging



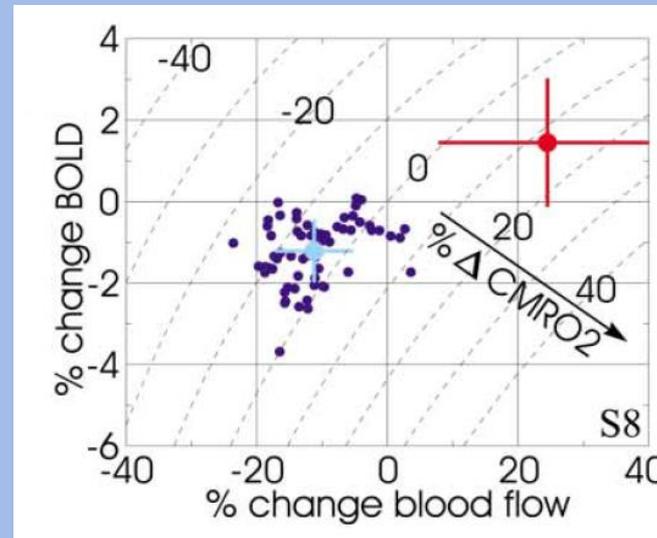
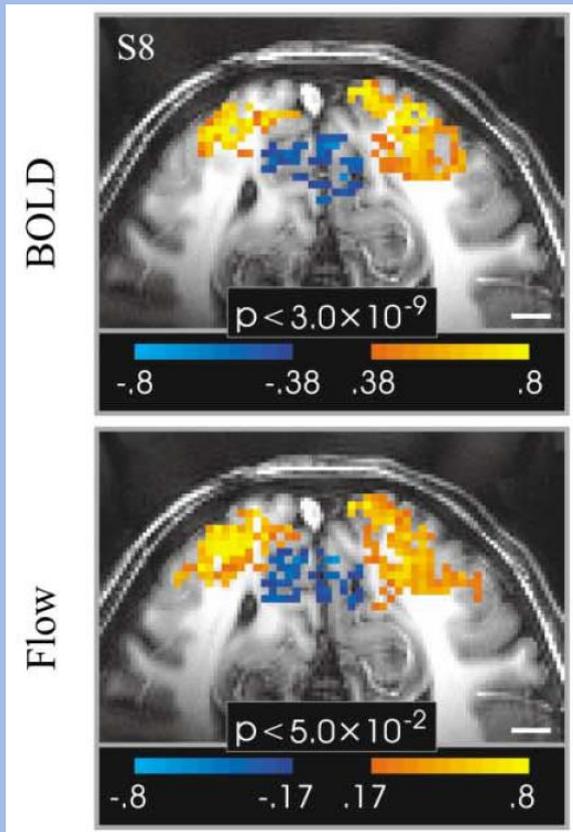
# Negative BOLD: challenging



Could NBR be originated by  $\downarrow$  CBF ? Caused by hypoxia  
Could NBR be originated by «vascular steal» ?

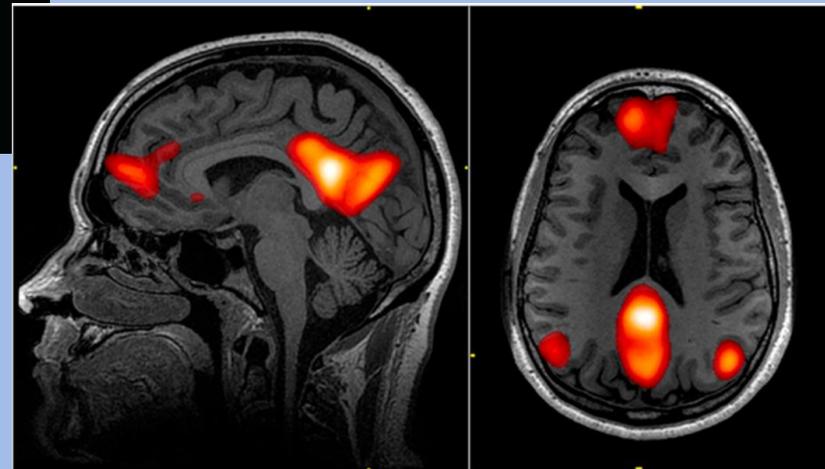
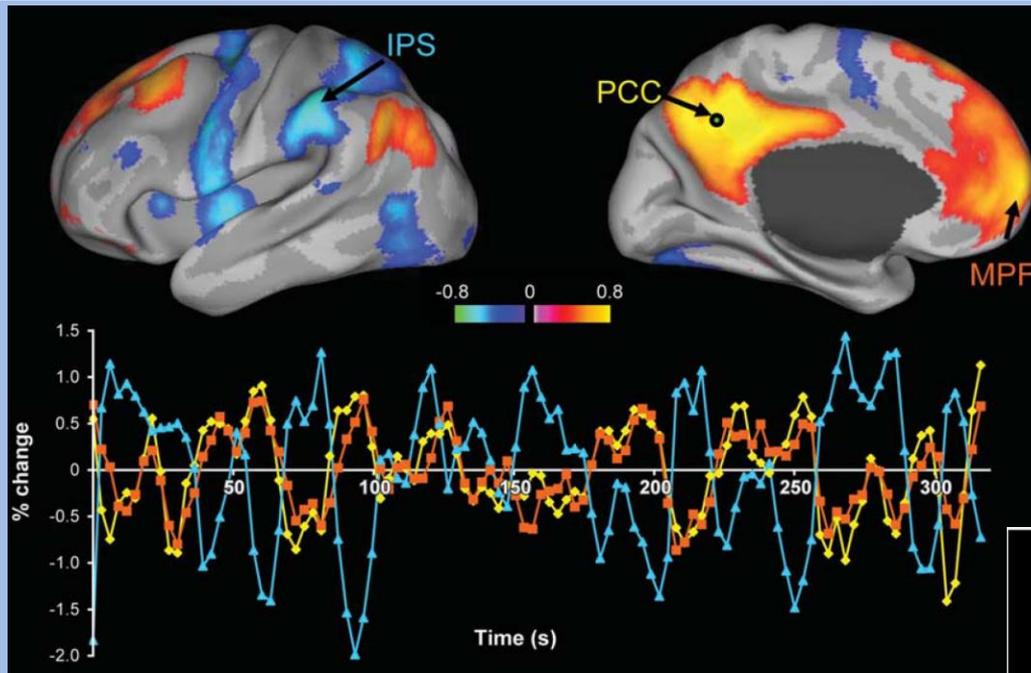
# NBR: BOLD and CBF\* measure

\* Cerebral Blood Flow (CBF)



**Conclusion: NBR is associated with a decreased CMRO<sub>2</sub>**

# Resting state/functional connectivity

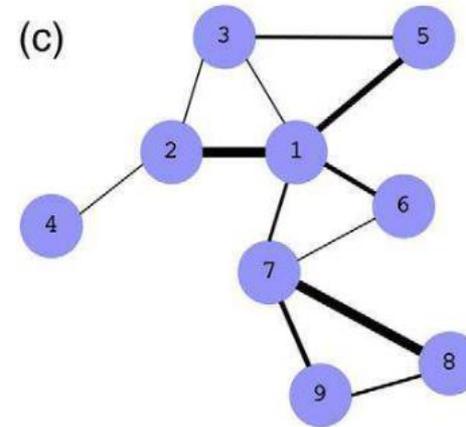
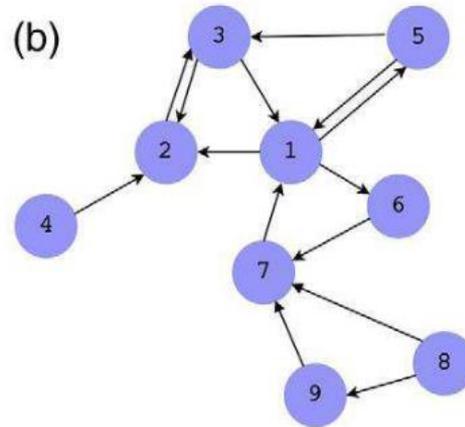
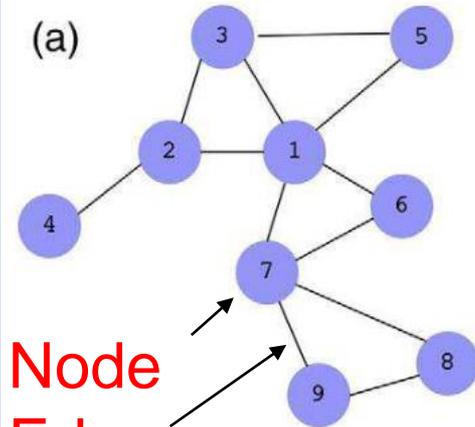


# Network analysis

undirected

directed

weighted



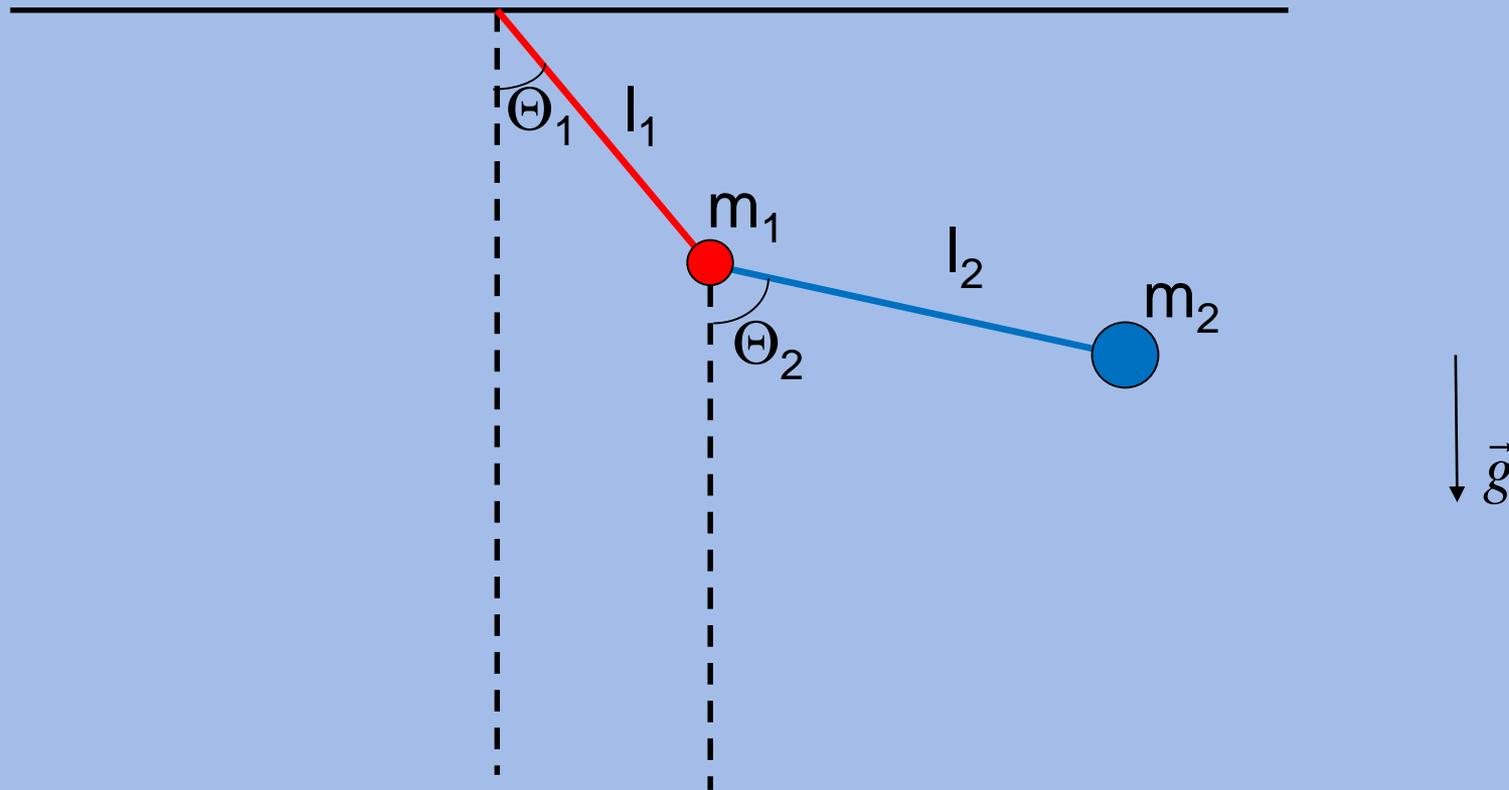
Node  
Edge

	1	2	3	4	5	6	7	8	9
1	0	1	1	0	1	1	1	0	0
2	1	0	1	1	0	0	0	0	0
3	1	1	0	0	1	0	0	0	0
4	0	1	0	0	0	0	0	0	0
5	1	0	1	0	0	0	0	0	0
6	1	0	0	0	0	0	1	0	0
7	1	0	0	0	0	1	0	1	1
8	0	0	0	0	0	0	1	0	1
9	0	0	0	0	0	0	1	1	0

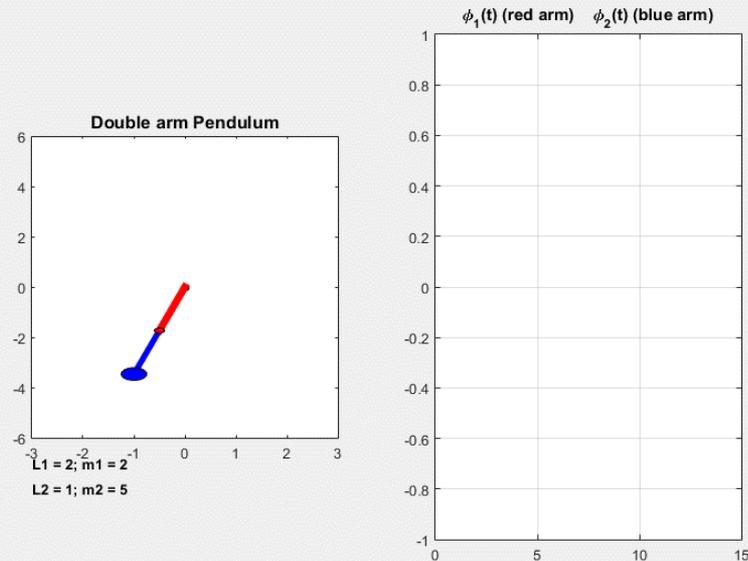
	1	2	3	4	5	6	7	8	9
1	0	1	0	0	1	1	0	0	0
2	0	0	1	0	0	0	0	0	0
3	1	1	0	0	0	0	0	0	0
4	0	1	0	0	0	0	0	0	0
5	1	0	1	0	0	0	0	0	0
6	0	0	0	0	0	0	1	0	0
7	1	0	0	0	0	0	0	0	0
8	0	0	0	0	0	0	1	0	1
9	0	0	0	0	0	0	1	0	0

	1	2	3	4	5	6	7	8	9
1	0	1	0.2	0	0.8	0.5	0.4	0	0
2	1	0	0.2	0.2	0	0	0	0	0
3	0.2	0.2	0	0	0.4	0	0	0	0
4	0	0.2	0	0	0	0	0	0	0
5	0.8	0	0.2	0	0	0	0	0	0
6	0.5	0	0	0	0	0	0.2	0	0
7	0.4	0	0	0	0	0.2	0	1	0.5
8	0	0	0	0	0	0	1	0	0.4
9	0	0	0	0	0	0	0.5	0.4	0

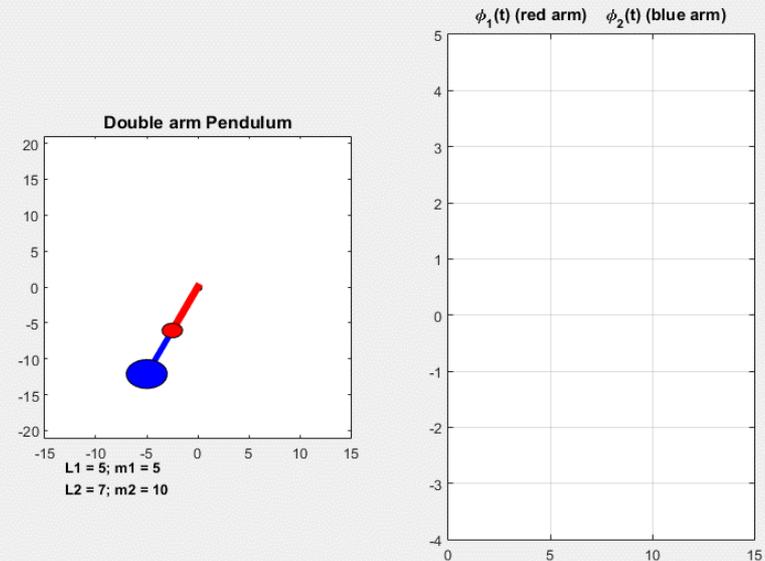
# Double Pendulum: approaching connectivity



# Functional coupling



$r=0.6166$ ;  $p<0.001$



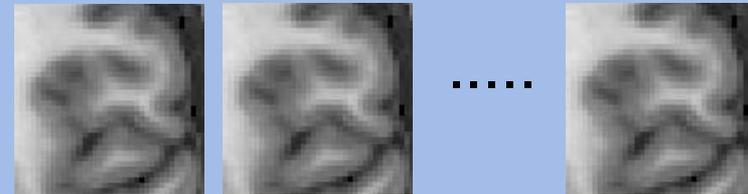
$r=-0.0106$ ;  $p=n.s.$

- > coupling present
- > connectivity is visible in the angle of both arms
- > interaction of red-to-blue arm

# Functional connectivity: assumptions

- >  $\approx$  homogeneous medium in GM
- >  $\approx$  homogeneous medium in WM
- >  $\approx$  micro vasculature
- >  $\approx$  nerve conduction velocity
- >  $\approx$  oxygen extraction fraction
- >  $\approx$  neuro vascular coupling
- >  $\approx k[\text{ATP}]$
- > .....

roi<sub>1</sub>    roi<sub>2</sub>    .....    roi<sub>n</sub>



# Functional connectivity: features

Transport: energy, information

## fast

- > large diameter axons
- > high NCV
- > extracellular
- > U-shape fibers

## slow

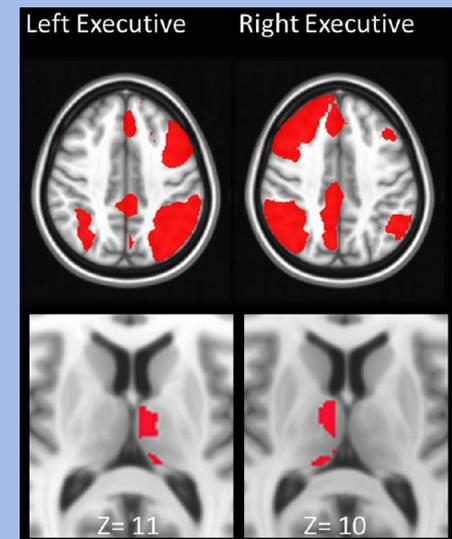
- > small diameter axons
- > low NCV
- > intracellular
- > frontal regions

# Thalamo-cortical Network



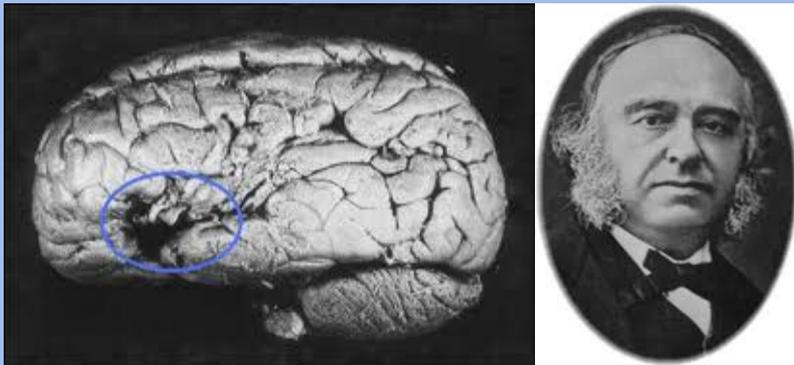
Thalamus as „seed“ ROI

Network analysis (independent components):  
i.e. each IC corresponds to a specific Network

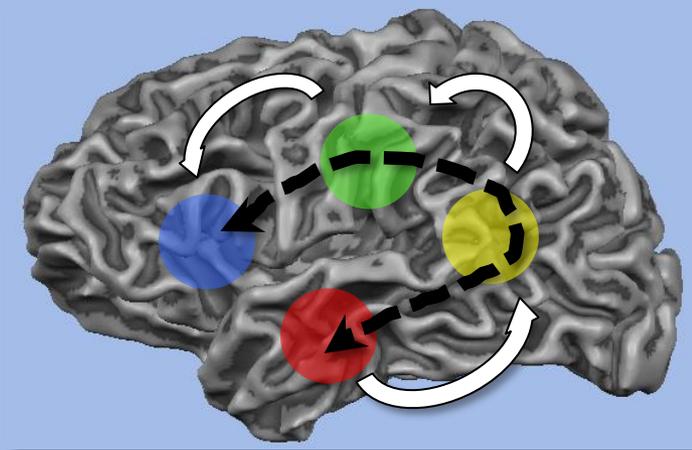


# Ideas behind RSN

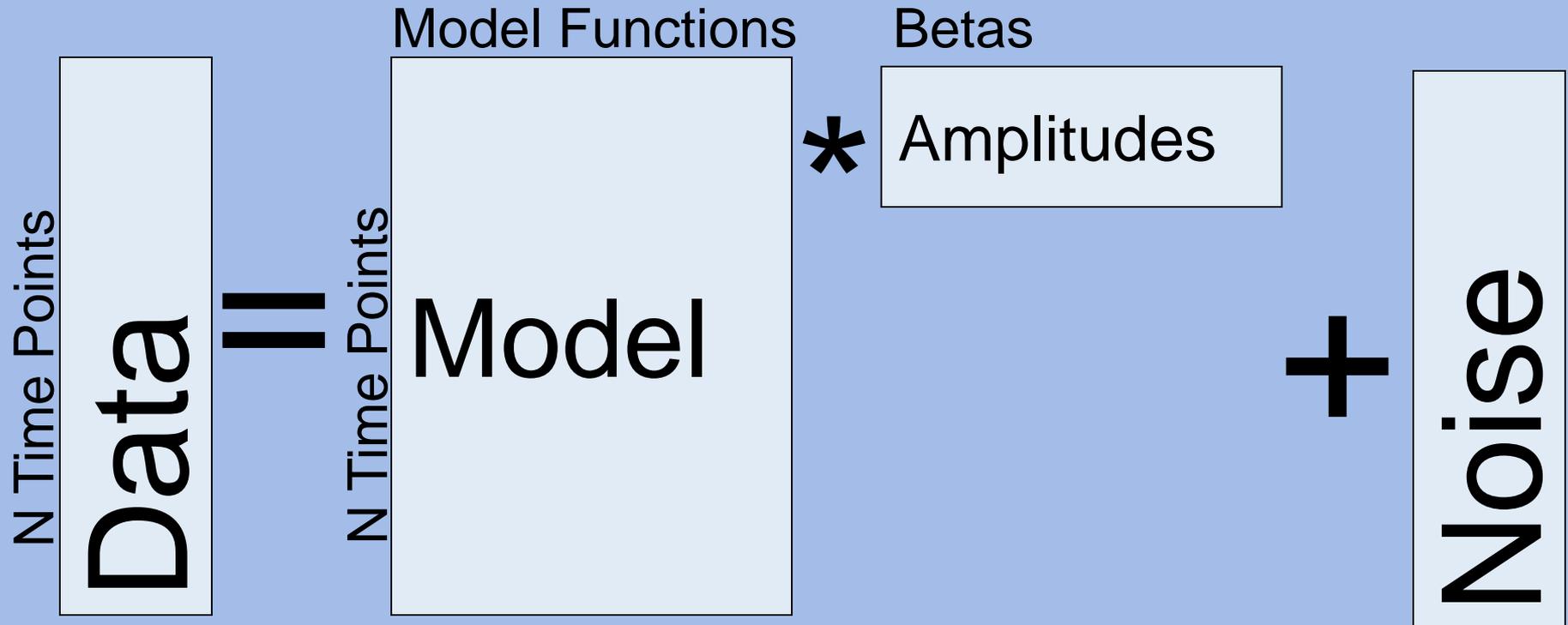
Localization <> Causality



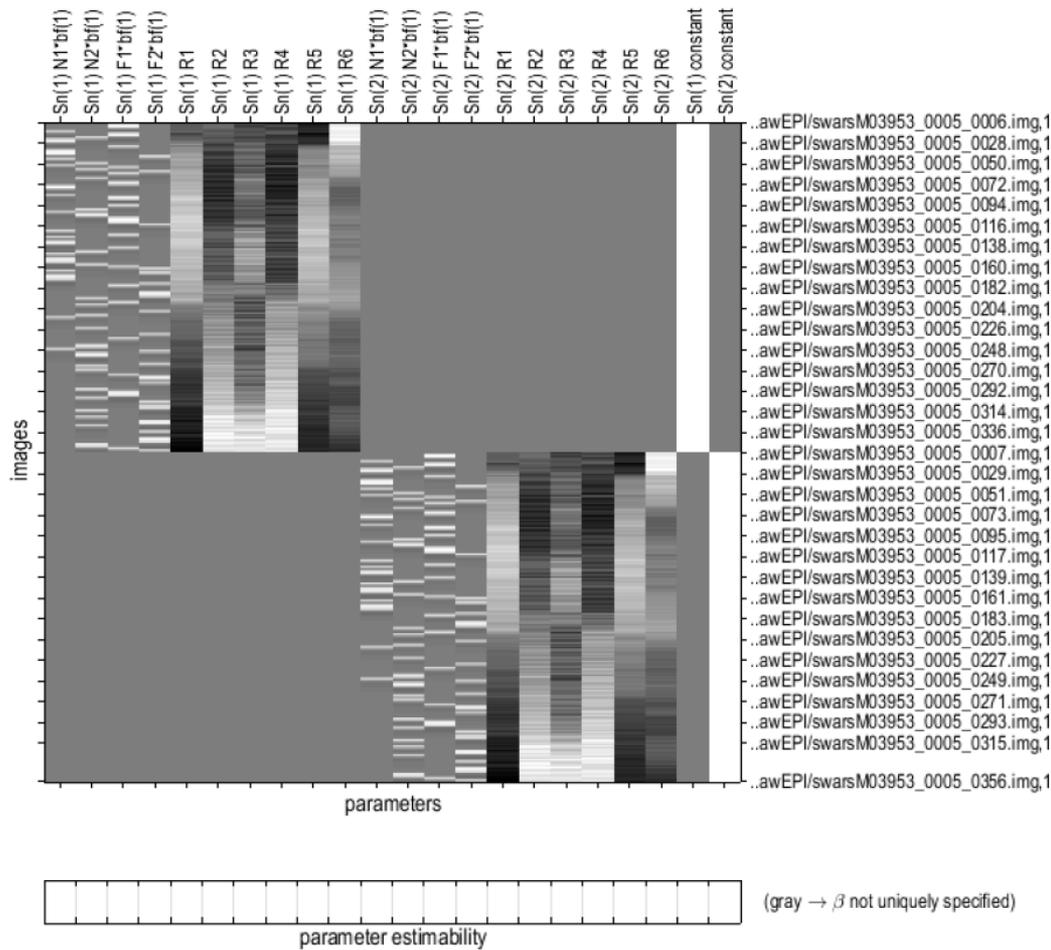
functional integration



# Statistical steps: GLM



$$Y = M\beta + \varepsilon$$



## Design matrix

### Design description...

Basis functions : hrf  
 Number of sessions : 2  
 Trials per session : 4 4  
 Interscan interval : 2.00 {s}  
 High pass Filter : [min] Cutoff: 128 {s}  
 Global calculation : mean voxel value  
 Grand mean scaling : session specific  
 Global normalisation : None

# Level of statistics

- 1. Level:** subject's level  
task performance, motion, etc.
  
- 2. Level:** between subjects and within subjects  
Group comparison  
result-generating statistics

# Different pathes of analysis in fMRI

Fixed Effects Analysis – (FFX)

concatenating all the subjects runs

Random Effects Analysis – (RFX)

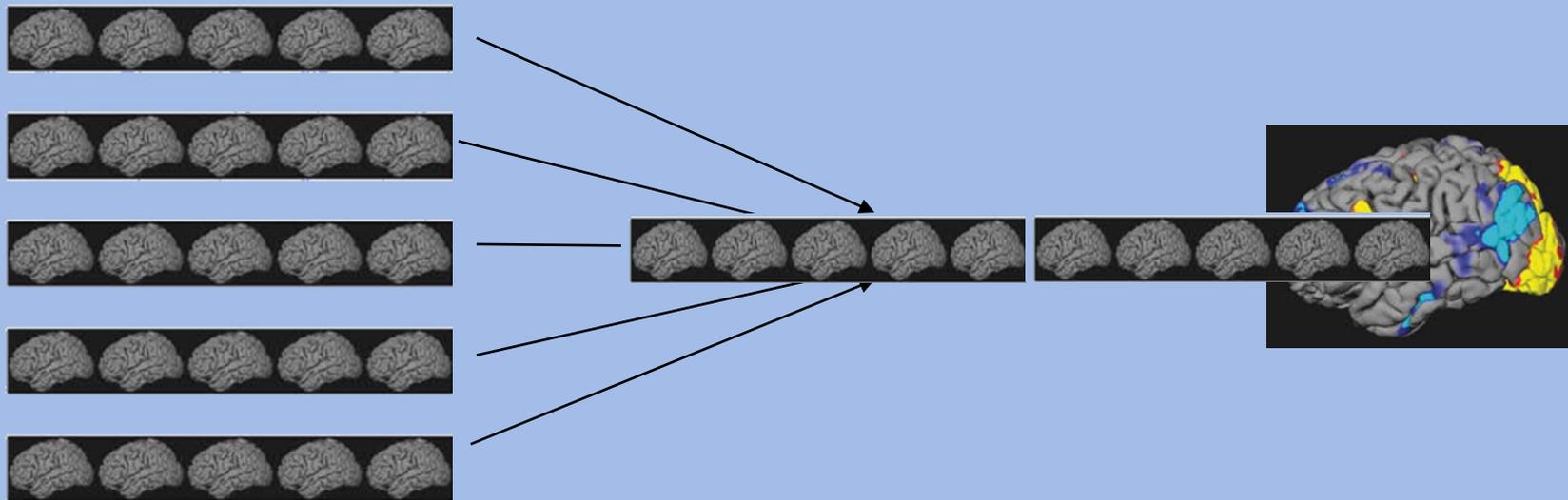
generalization to the population level

# Fixed Effects Analysis FFX

concatenate subjects

degree of freedom „big“

Allows inference to subject's sample

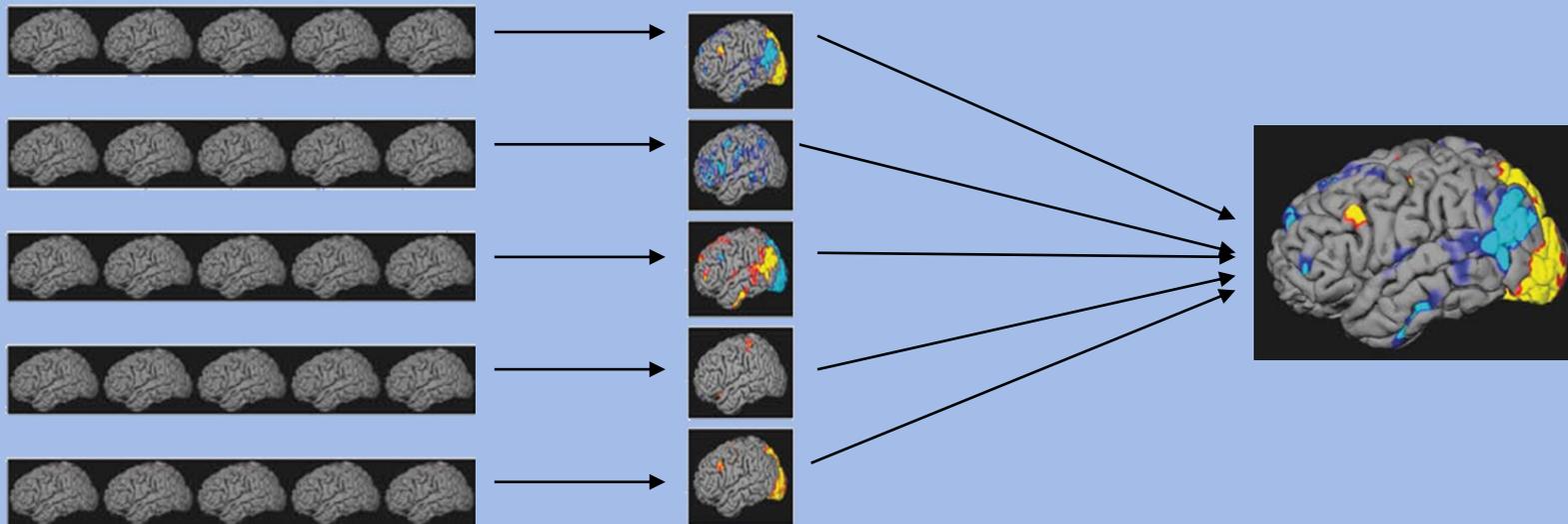


# Random Effects Analysis RFX

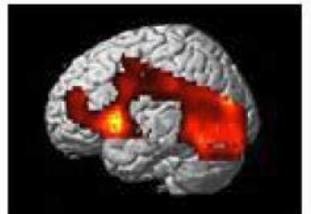
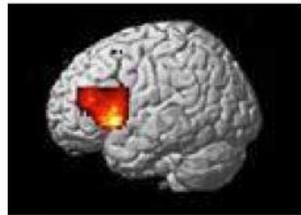
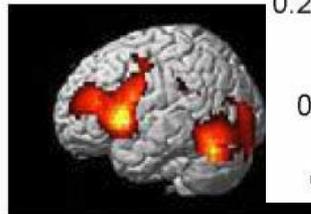
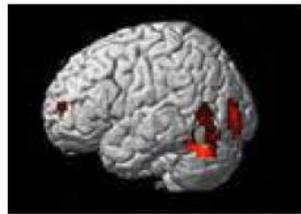
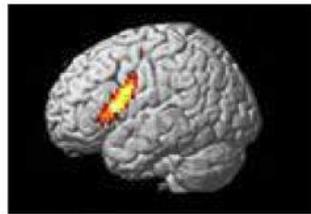
concatenate subjects

degree of freedom „small“

Allows inference to population from the sample cohort

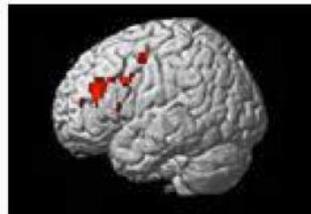


# FFX vs RFX

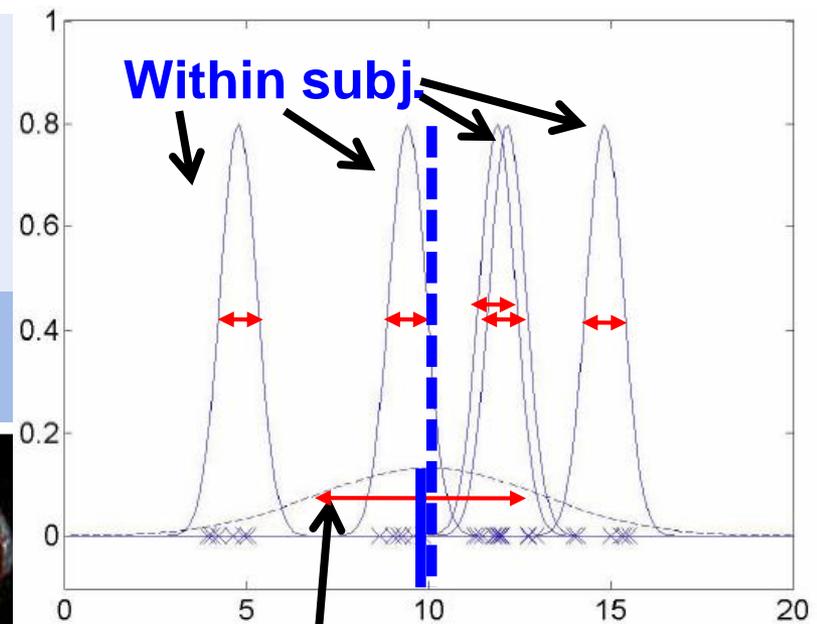


(a) Individual subject activations

(b) Group Fixed Effects



(c) Random Effects



Between subj.

↔ variance

| average

# That's science: it's all about assumptions

Procedure	Assumptions
Design	Previous literature
Record data	Patient/Control; Drugs; Circadian rhythm; age; gender; social status; etc... MR scanner; Resolution ( $\vec{x}, t$ ); Temperature; Pressure; etc...
Preprocess data	Gaussian distribution; serial correlation; Coregistration; Normalize Template; Smooth; etc...
1-level statistics	Gaussian distribution; linear trend; GLM residuals; etc...
2-level statistics	Gaussian distribution; variance; independent data; GLM residuals; etc.
Inference statistics	Correct p for multiple comparison; <b>Random Field Theory; Smooth Field; Spatial autocorrelation</b> ; etc...

# Multiple testing

## Cluster failure: Why fMRI inferences for spatial extent have inflated false-positive rates

Anders Eklund<sup>a,b,c,1</sup>, Thomas E. Nichols<sup>d,e</sup>, and Hans Knutsson<sup>a,c</sup>

*PNAS* 2016 Jul 12;113(28):7900-5

## What's all about ?

- $H_a$ : alternative hypothesis      There is activation!
- $H_0$ : null hypothesis      No effect
- $\alpha$ : false positive rate  
probability to *reject*  $H_0$   
when  $H_0$  is TRUE      } type I error
- $\beta$ : false negative rate  
probability to accept  $H_0$   
when  $H_a$  is TRUE

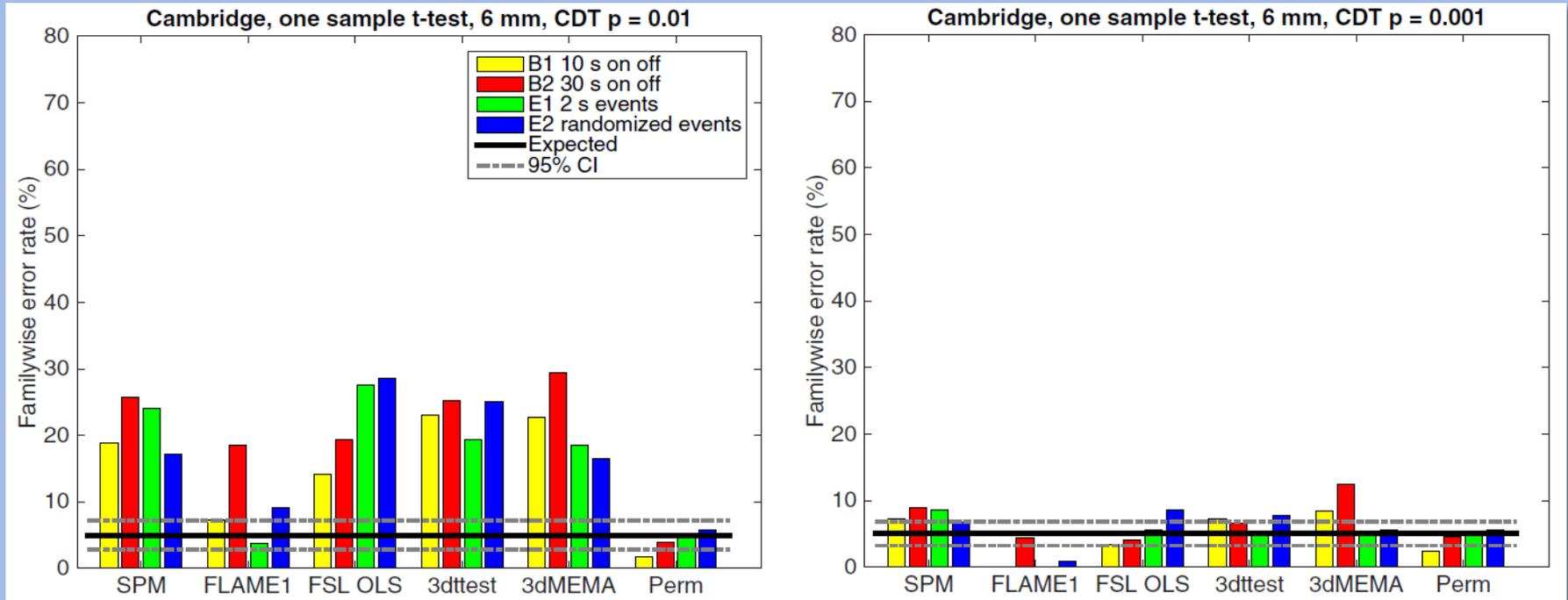
The probability to make at least one type I error [Family Wise Error Rate (FWER)]

# Multiple testing

FWER (voxels): find  $p_{\alpha} = 0.05$   
so that there is a 5 %  
chance to find at least  
1 activated voxel

FWER (cluster): with  $p_{\alpha} = 0.05$  find the  
# voxels in a cluster  
so that there is a 5 %  
chance in the cluster to  
find at least 1 activated  
voxel

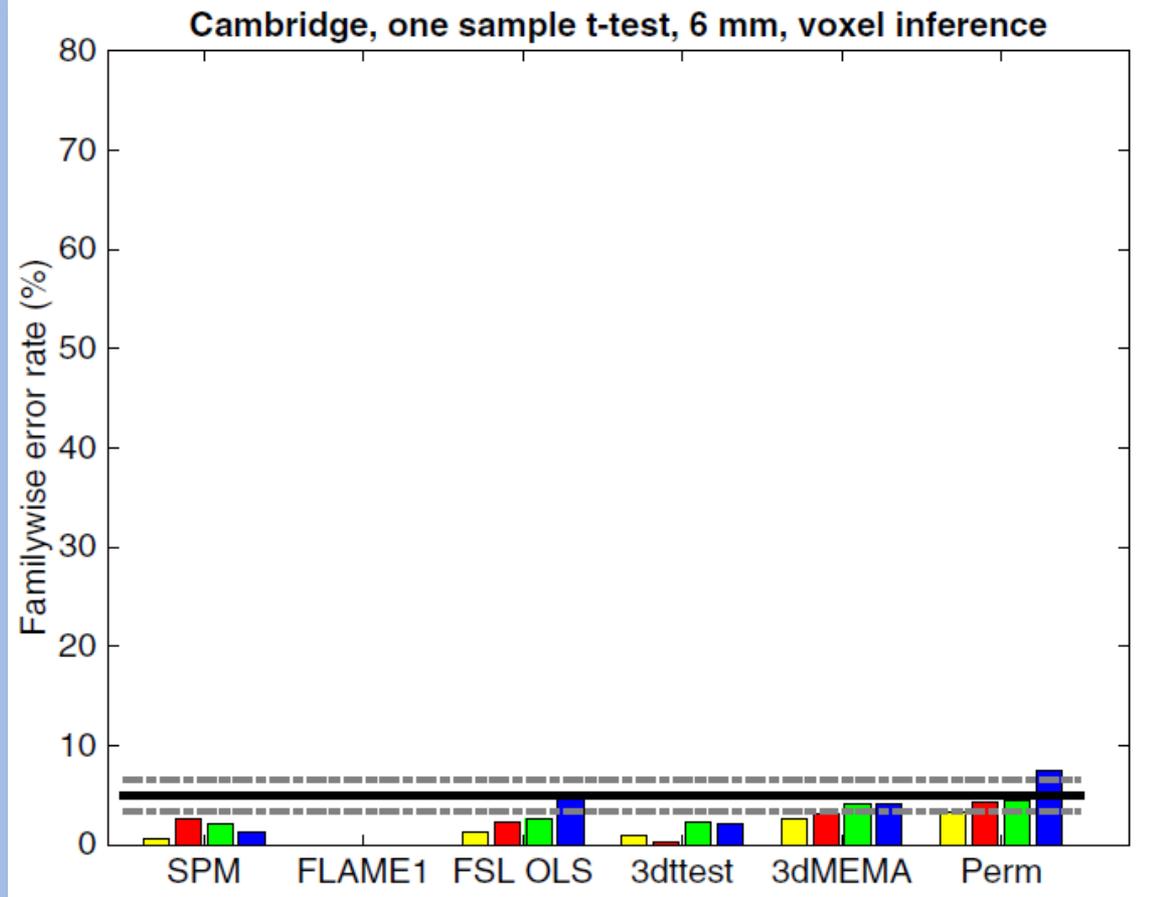
# SPM, FSL, AFNI, non-parametr.



Cluster wise threshold: **Not OK** for  $p < 0.01$

~OK for  $p < 0.001$

# Voxel wise threshold OK



## Multiple testing (Bonferroni)

If we have  $64 \times 64$  voxles  
we do 4096 test:

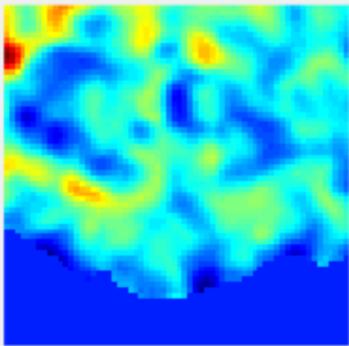
$$p \leq \frac{0.05}{4096} = 0.0000122$$

Example:

$$\begin{aligned} df &= 30; \\ p &= 0.0000122 \\ t &= 1 - \text{tinvs}(0.05/4096, 30); \end{aligned}$$

$$= 5.9834$$

too conservative !



no correcction

$t > 5.98$

# Random Field Theory

mathematical model:

- estimate the # RESEL in your search volume
- estimate the # cluster (thresholded at some level)
- and correct the thresholded level

# Random Field Theory

Independent data: data of one voxel should be independent of its neighbourhood

in fMRI spatial correlation is present !

Smoothness: should be constant over the brain  
how to check this ?

*Problems when:*

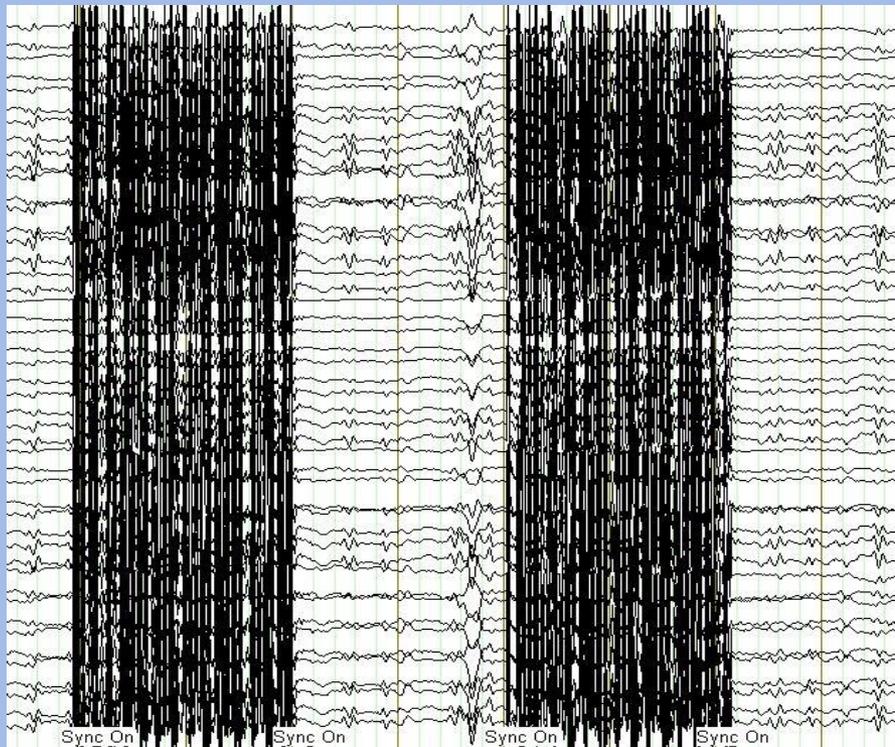
- Small sample size
- errors/residuals not normally distributed and not smooth

## Take home

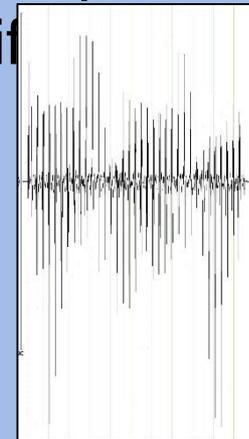
- Check your data carefully (assumptions Y/N ?)
- Investigate into Signal and Noise in your data !
- Careful interpretation of results;  
especially when dealing with (large) clusters
- Non-parametric SnPM may be an optimal choice
- COBIDAS\* White paper with guidelines «best-practices»

# DATA PRE-PROCESSING

## Scan-Puls artifact correction



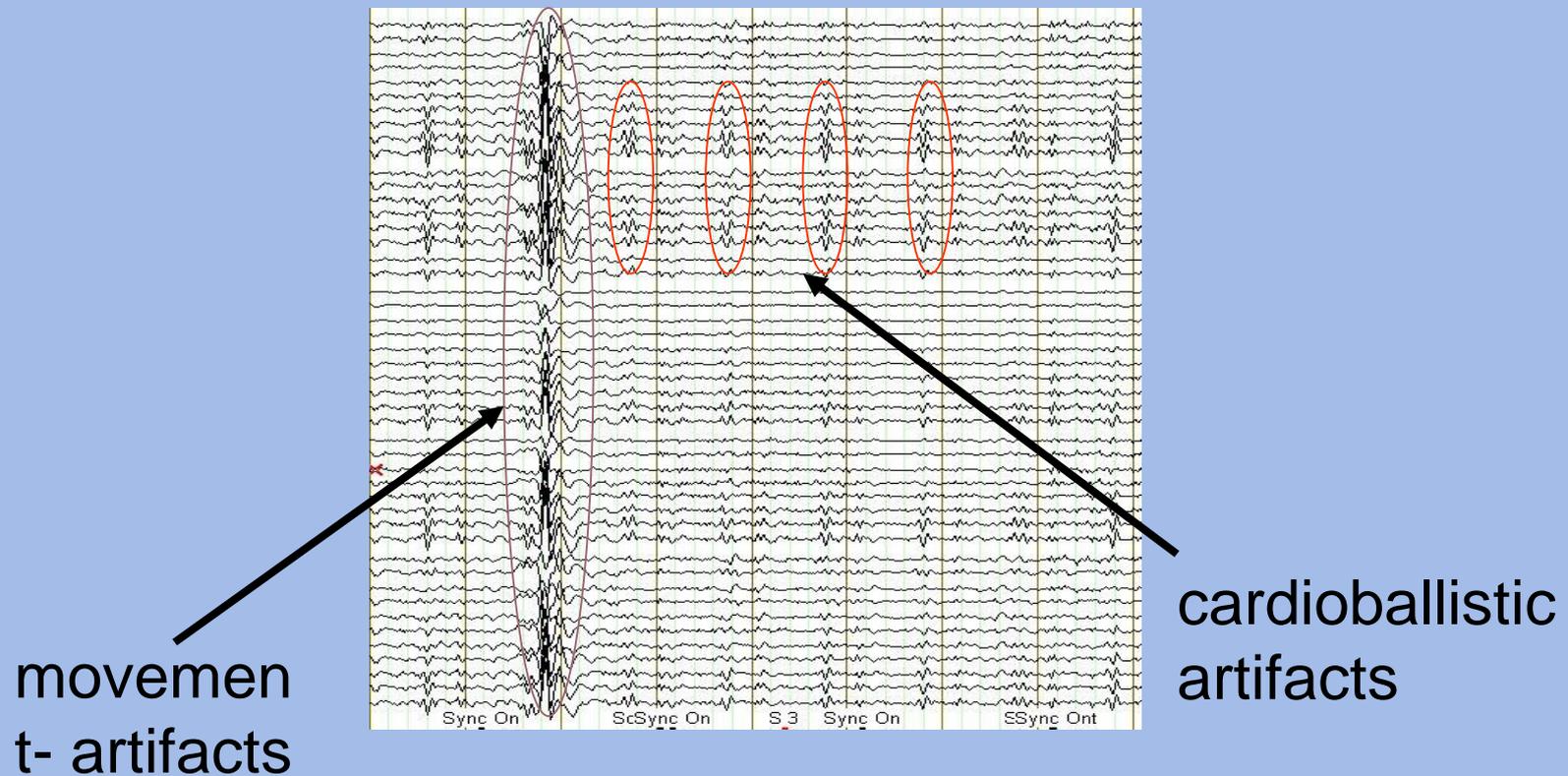
Subtraction of  
a template-  
artifact



EEG in 3T MRT

# DATA PRE-PROCESSING

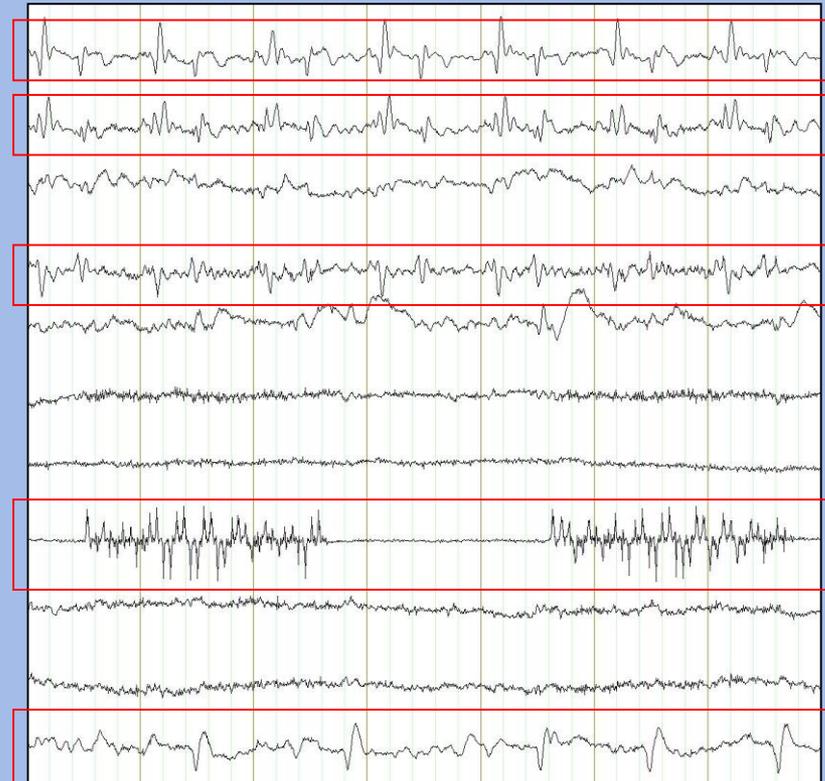
## After Scan-Puls artifact correction



## Independent Component Analysis

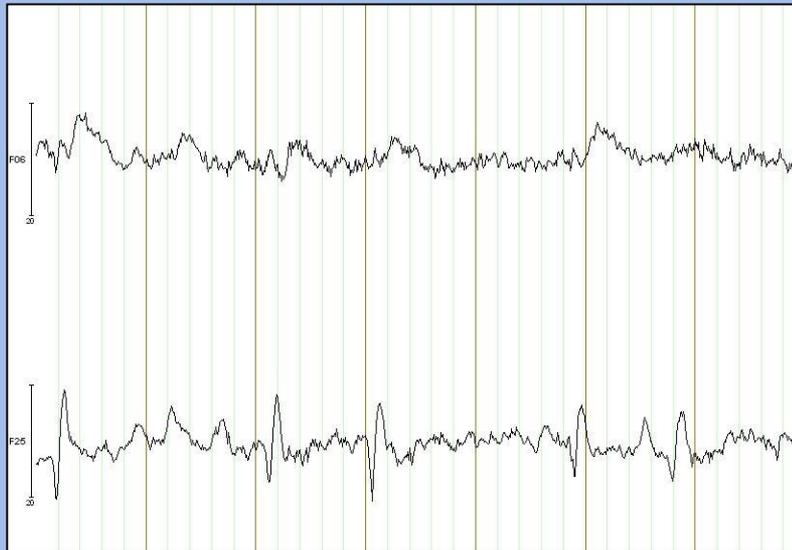
Decomposition of EEG data into independent factors.

- Scan-Puls artifact
- Cardioballistic artifact
- Eye blinks
- Epileptiform activity
- Other

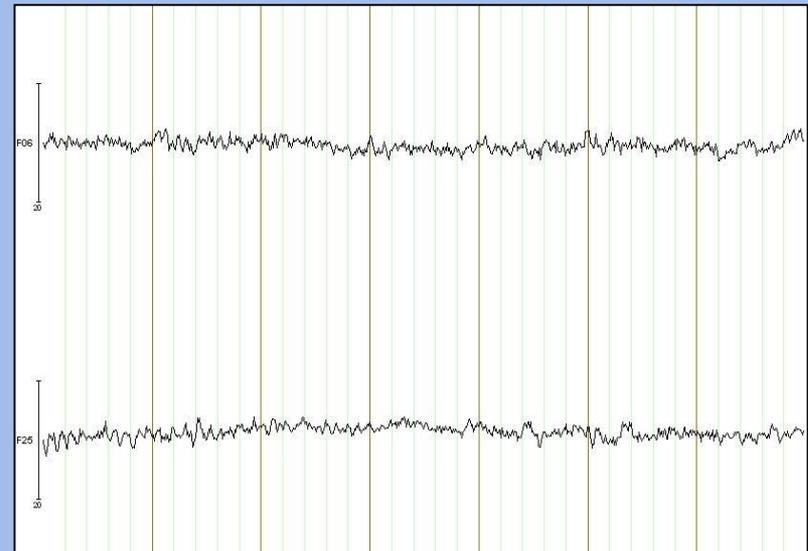


# ICA FACTORS (EXAMPLE)

## Example factors coding for epileptiform activity



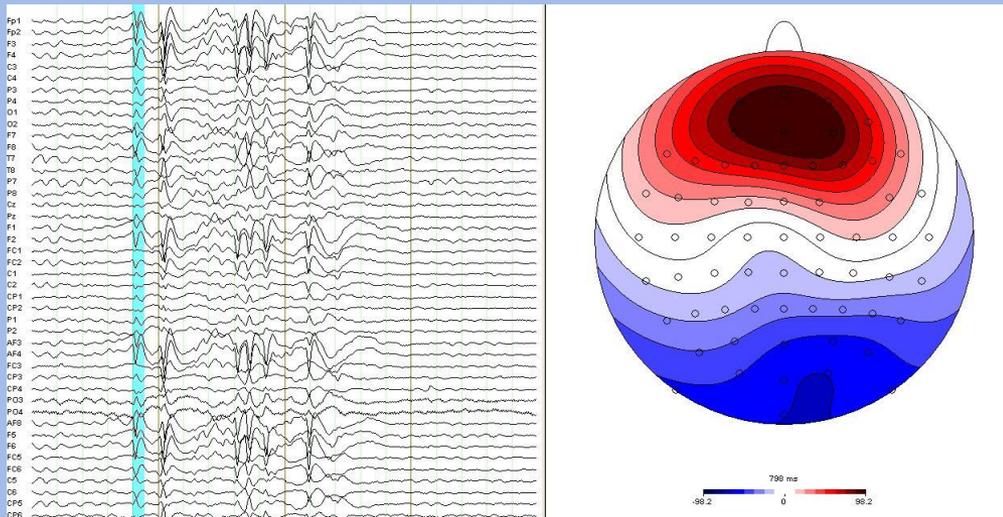
Epoch **WITH** interictal spikes.



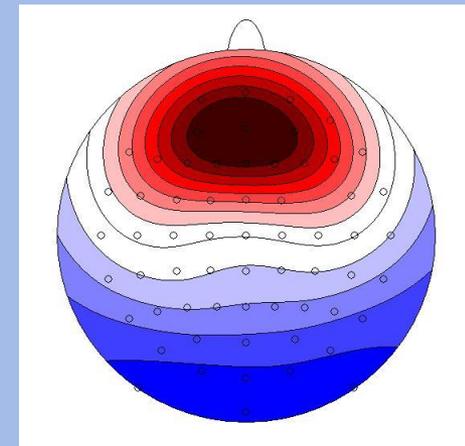
Epoch **WITHOUT** discharges.

# ICA FACTORS (EXAMPLE)

Selection of factor further dependent on topography



EEG topography.

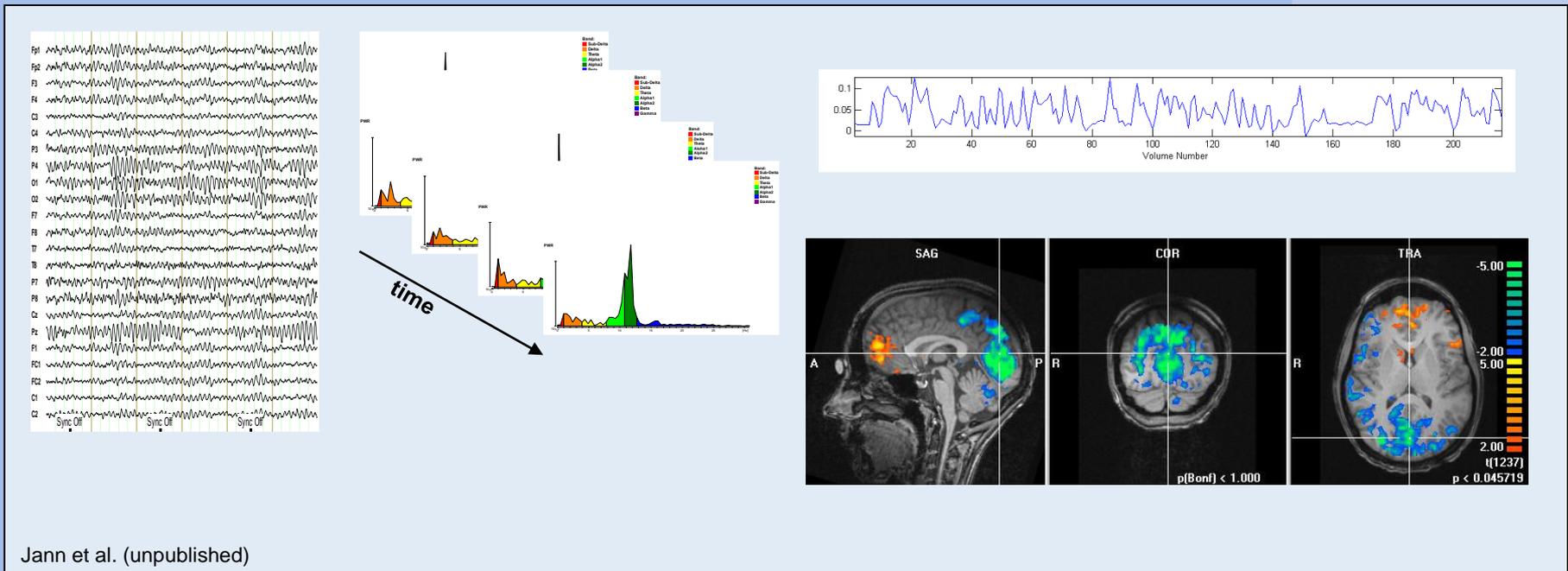


ICA factor topography.

# Features extracted from EEG

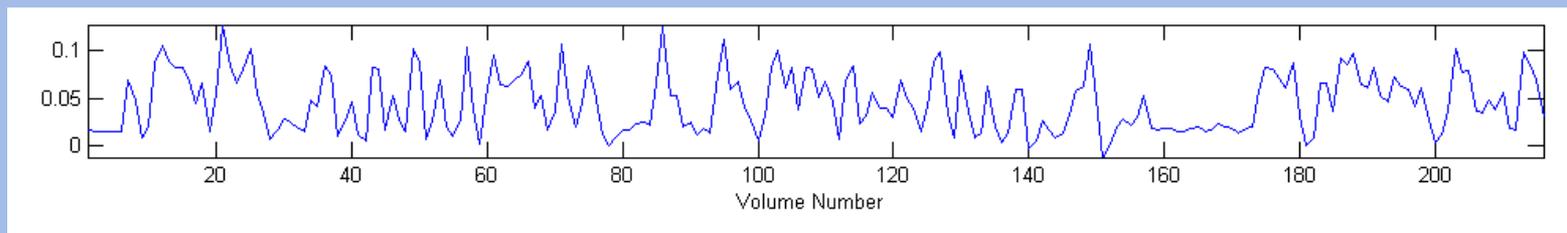
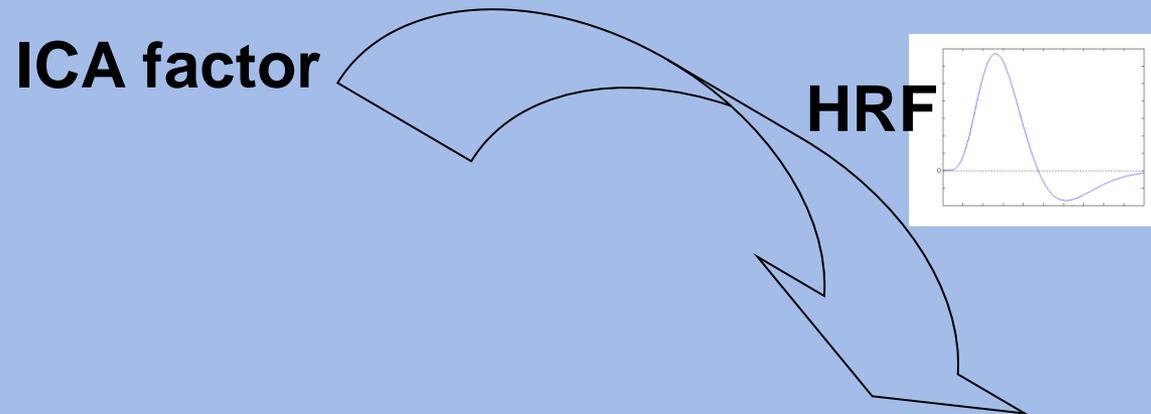
## Spontaneous Activity

- Frequency domain



# DATA PROCESSING

## Convolution with a 'Hemodynamic Response Function' (HRF)

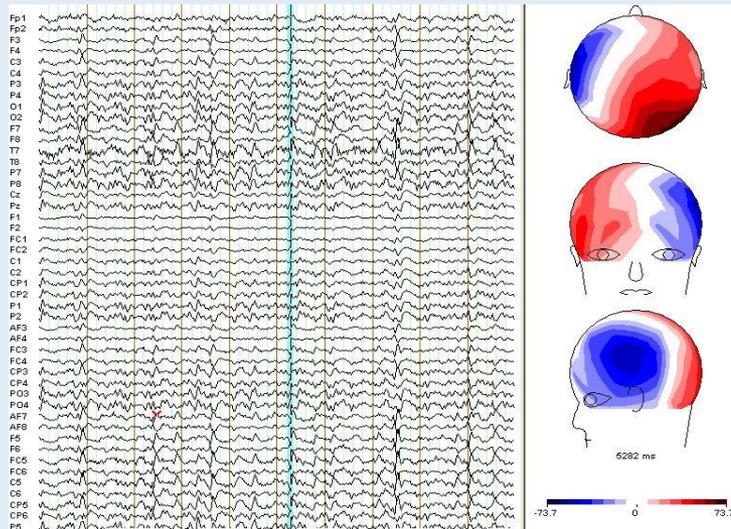


**Predictor for fMRI BOLD signal**

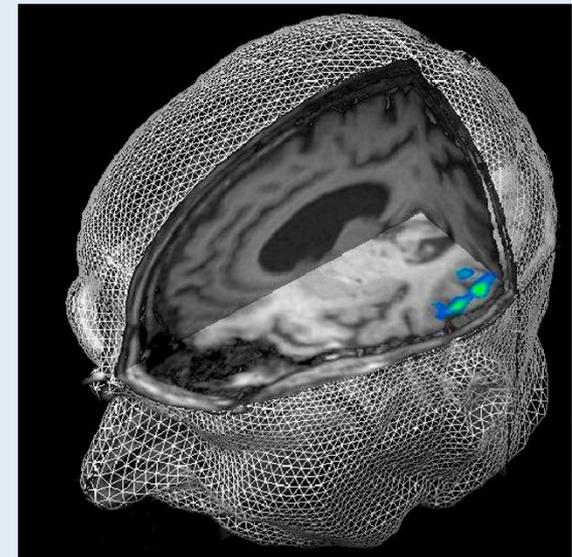
## Spontaneous Activity

- Single events

e.g. Epilepsy: unpredictable events



Occurrence of  
IEDs as  
predictor for  
BOLD



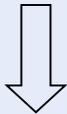
Adapted from Jann et al., Neuroimage, 42 (2008), p635-648

# fMRI study

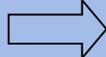
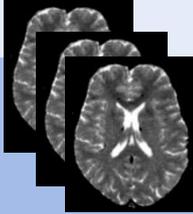
$u^b$

<sup>b</sup>  
UNIVERSITÄT  
BERN

Experimental  
setup



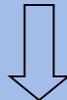
Recorded time  
series (EPI)



Coregistration of  
functional and  
anatomical data



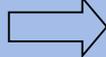
Anatomical  
images



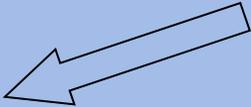
Segmentation  
Cortex reconstruction

Spatial  
normalisation

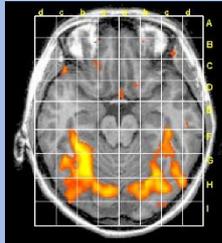
Preprocessing



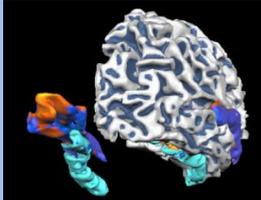
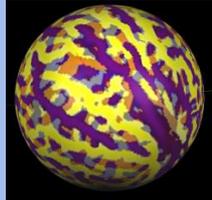
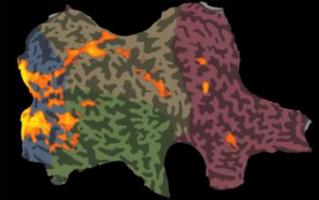
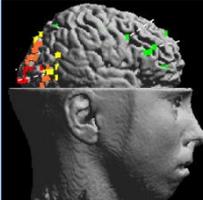
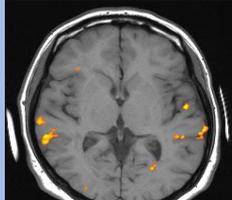
Statistical localization of brain  
activation, functional maps



Single subject analysis



Group analysis I  
Talairach space



Group analysis II – Surface-based alignment