



### Noise Models and Correction for fMRI

11111

### - an Introduction to the PhysIO Toolbox

Matthias Müller-Schrader

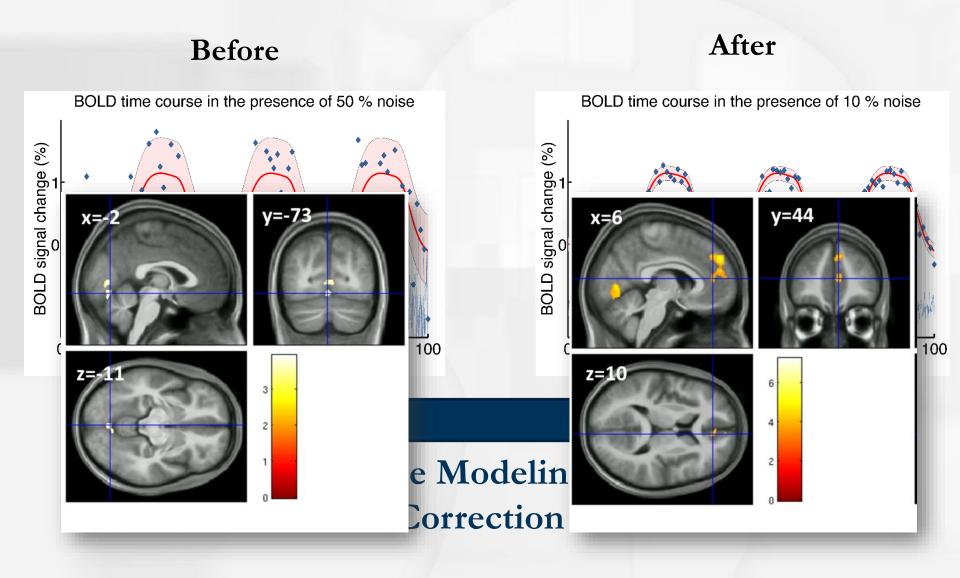
Nov 19<sup>th</sup>, 2019

Generous slide courtesy

Lars Kasper

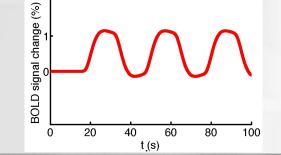
### The Goal of Noise Correction



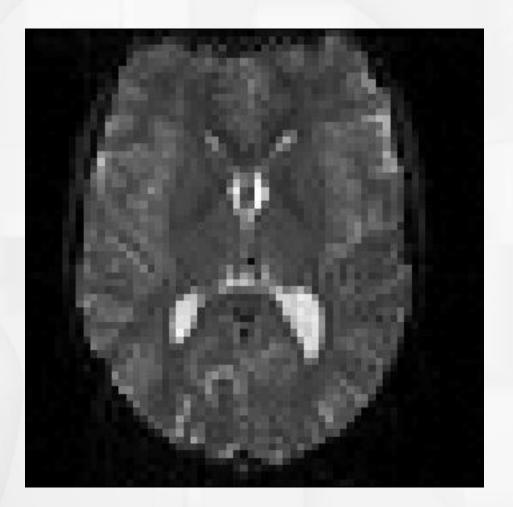


### Reminder: fMRI Data is noisy...







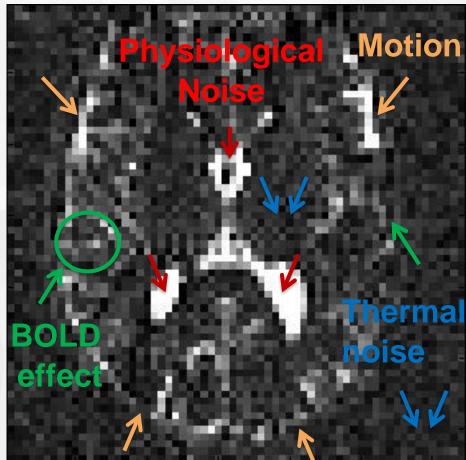


### fMRI Data is noisy...



# Interest in fluctuations only: Subtract the mean

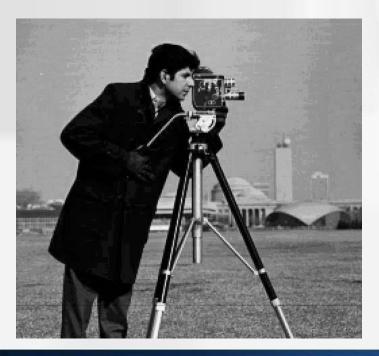


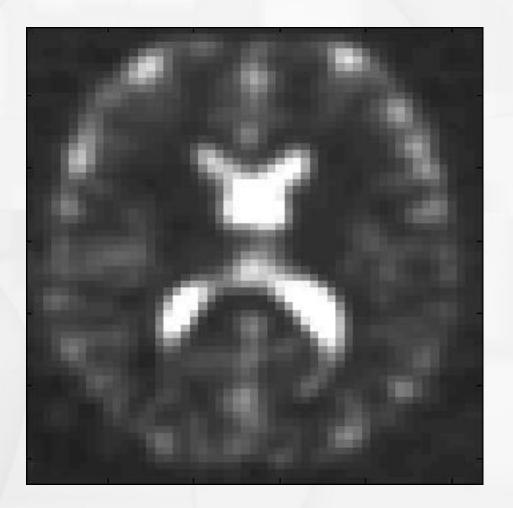


# Previously...



- How we ended 6 weeks ago (after preprocessing)
- After smoothing...still some fluctuation





# **Recap: Imaging pathway**



See blackboard.

# Physiological noise



- Noise: «Fluctuations we are not interested in.»
  - Often random
- Sources of noise
  - MR-System
    - Heating of gradient coils
    - Noise in amplifiers
    - Spikes in coils
  - Subject in the Scanner
    - Motion
    - Physiological noise
      - Cardiac cycle
      - Breathing cycle
    - Not the BOLD-signal

### Outline



- Why denoising?
- Pathways of physiological noise
  - Recap: MR image encoding
  - Cardiac effects
  - Respiratory effects
- Noise correction approaches
  - Method
     Preprocessing vs modeling
  - Input
     fMRI data vs. peripheral measure
- Effects of noise correction
- Limitations

### Outline



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### Limitations

# fMRI = Acquiring Movies

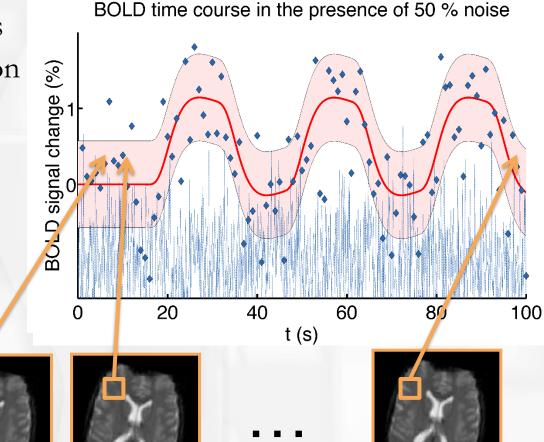


- The Localized Time-series is the Fundamental Information Unit of fMRI
- Signal: Fluctuation through Blood oxygen level dependent (BOLD) contrast

Noise: All other fluctuations

Run/Session: Time Series of Images

2019-11-19



time

Matthias Müller-Schrader: fMRI Noise Models & Correction

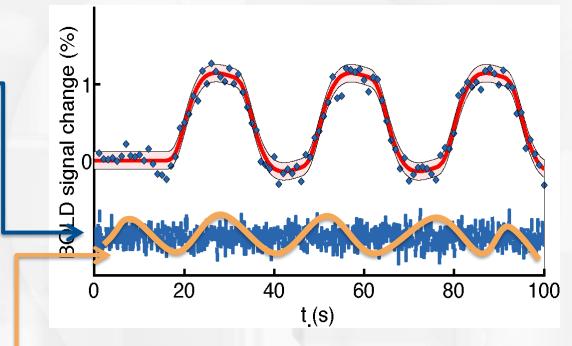
scan 1

scan N

### **Noise Categories**



- Thermal Noise
  - temporally uncorrelated
  - reduced SNR → risk of false negatives
  - Remedy: Spatial Smoothing
- Noise: All other fluctuations
- "Structured" Noise
  - temporally correlated
  - reduced SNR → risk of false negatives
  - correlated with task  $\rightarrow$  risk of false positives
  - Remedy: Noise modeling (e.g. GLM)

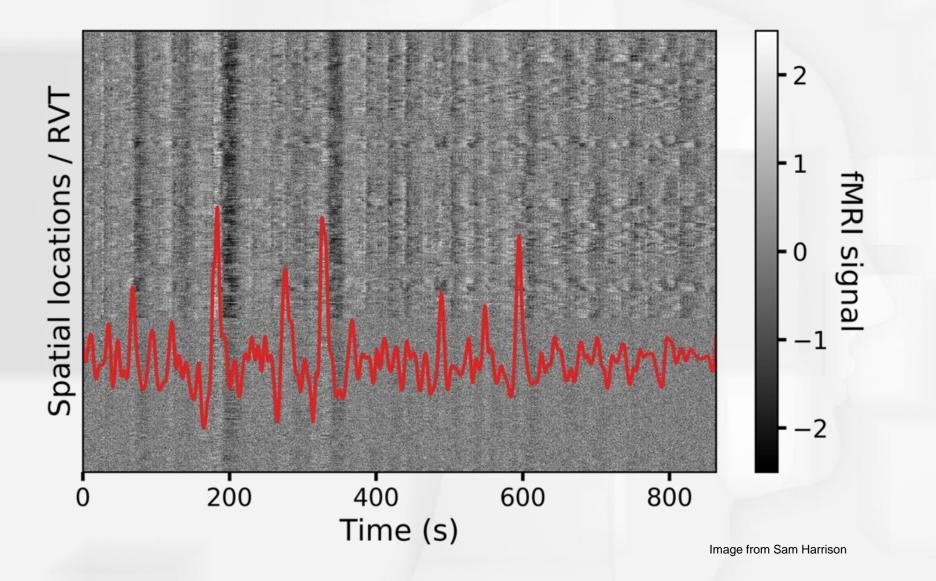


Inference = Signal-To-Noise

$$t = \frac{\beta}{\sqrt{\sigma_{\varepsilon}^2 (X^T X)^{-1}}} = \frac{\beta \| \boldsymbol{x} \|}{\sigma_{\varepsilon}}$$

$$F = \frac{N - M}{M_1} \cdot \frac{(\sigma_S^2 + \sigma_N^2) - \sigma_N^2}{\sigma_N^2}$$

# False positives in resting state fmri



### Outline



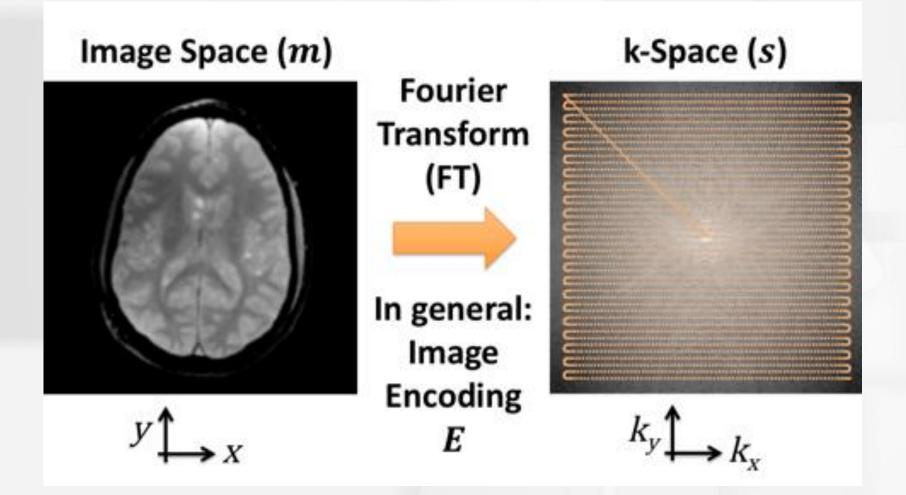
### Why denoising?

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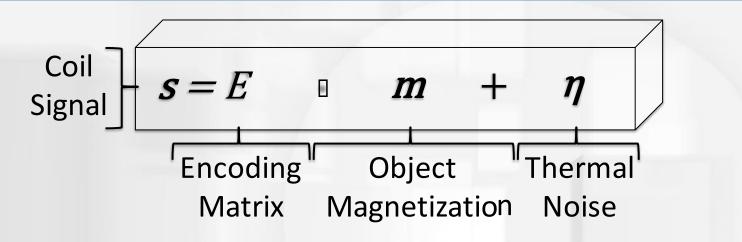
### **Recap: MR Image Encoding**





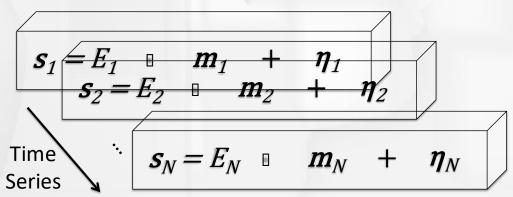
### Image Reconstruction & Noise





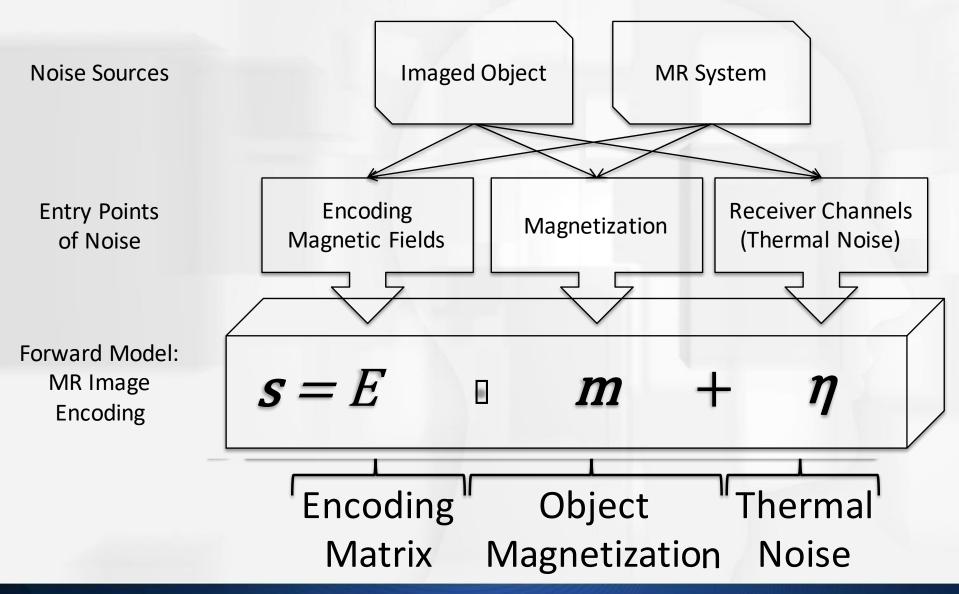
- Image reconstruction is also a *huge* GLM,  $\sim 10^5$ - $10^6$  rows
  - <sup>3</sup> mm slice, 8 chan: 64<sup>2</sup>\*8 = 512k
  - 1 mm slice, 32 ch:  $256^{2*}32 = 2M$
- Any change between volumes in encoding matrix (field), object magnetization and thermally induces image noise

$$\widehat{m} = (E^H E)^{-1} E^H s$$



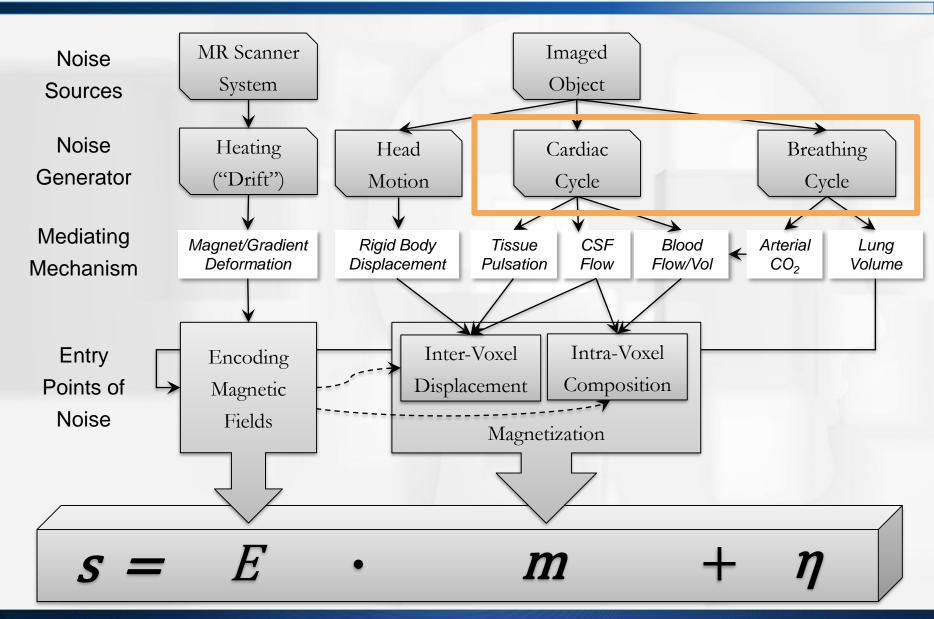
### What fluctuates?





### **Structured Noise in MRI**





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### Limitations

### Cardiac effects: CSF pulsation

### • CSF pulsation:

#### Systole:

- Blood pumped into brain, vessel
   volume increases: pulsatile vessels
- CSF pushed down: pulsatile CSF
- Diastole:
  - Vessel volume decreases
  - CSF flows back into "void" brain volume

#### A Cardiac Cycle in the Brain

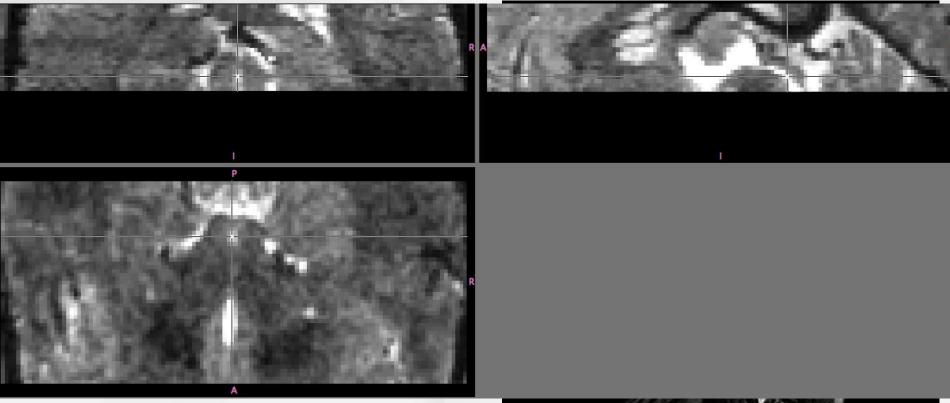






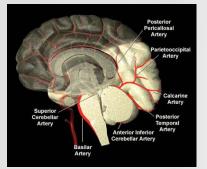
#### **Triggered High-Resolution fMRI**

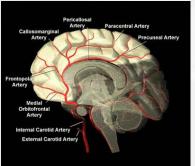
#### A Cardiac Cycle in the Brain

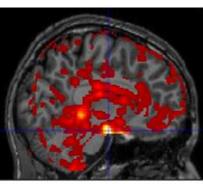


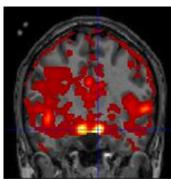
### **Cardiac effects**

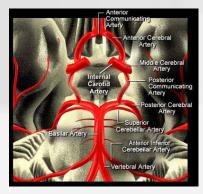




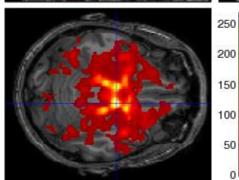








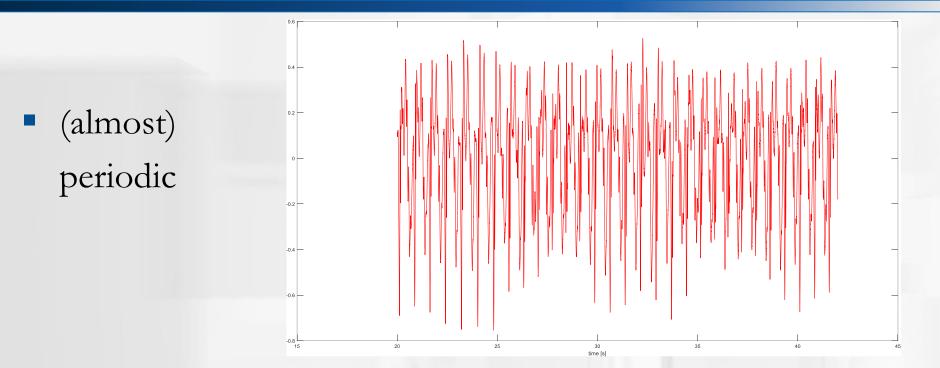
#### Vessel Anatomy



Locations of Fluctuations

# **Cardiac signal**





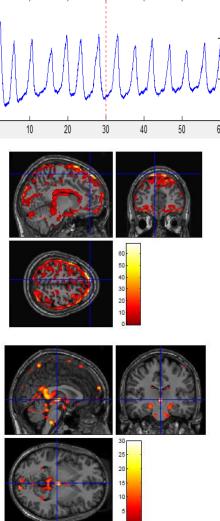
### But:

 Fluctuations in BOLDlevel due to heart-rate variability (HRV)

### **Respiratory effects**







- Changes in lung volume change encoding magnetic field for MR
  - Chest (&head) moves with respiratory cycle
  - Geometric distortion/scaling
- Change in oxygenation of blood
- Respiratory-sinus arrhythmia
  - Heart beats faster during inhalation

### Outline



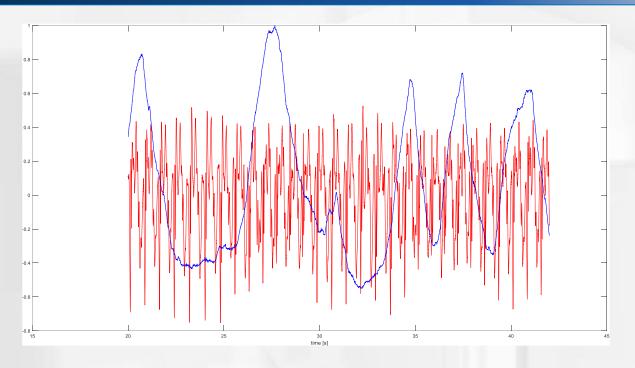
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### Limitations

### Why not just filtering?

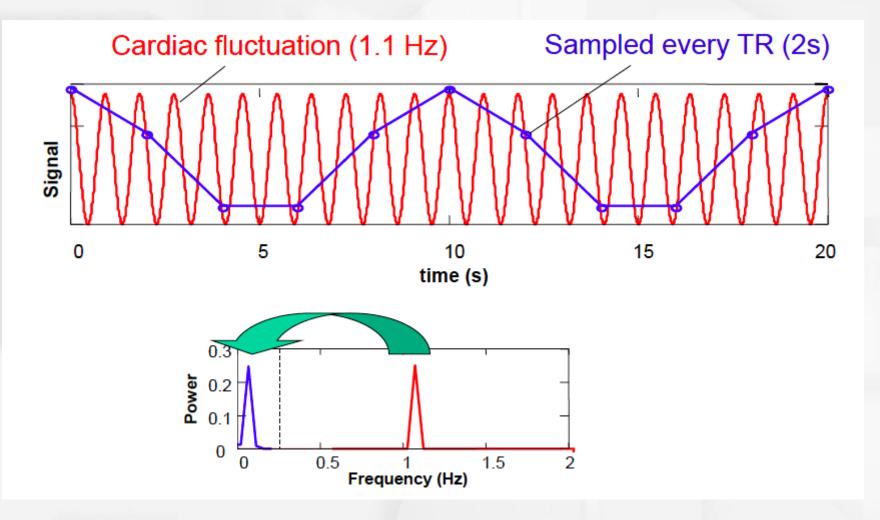




- Cycles are (almost)periodic
   Problem: Aliasing
  - Could filter frequencies
    - 0.2-0.4 Hz respiration
    - 0.8 1.2 Hz heart beat (main)

### **Aliasing of Physiology**





Courtesy: R. Birn, HBM 2015

# **Modeling VS Preprocessing**



- Modeling:
  - Filters, projections (e.g. to independent components) etc. are all linear operations
  - Combination in one design matrix, together with task
  - Simple test of correction efficacy: F-test on nuisance regressors
- Preprocessing:
  - The data y entering the GLM is altered  $\Rightarrow y' = X\beta + \varepsilon$
  - For non-linear changes of y or inter-voxel dependencies, alteration outside GLM necessary

# The Problem with Preprocessing 50

- Problem: No inherent measure of efficacy (F-test in GLM), correlation with task regressors undetected
- "Advantage": No loss of degrees of freedom (sensitivity of F-test)
  - But it it only a hidden loss, statistics for inference is biased, if performed modeling is not incorporated
- Modeling via GLM recommended, if possible
  - Drifts, Motion Regressors
  - RETROICOR, HRV, RVT
  - aCompCor, (ICA)

# Motion: Preprocess & Modeling

- Correction for motion artifacts is actually a combination of Preprocessing and modeling
- Preprocessing cannot correct spin-history effects, intravolume movements (non-rigid!), small partial volume effects
- Preprocessing:
  - Realignment
  - Motion "Scrubbing"
- Modeling (from estimated realignment parameters)
  - Retrospective Modeling: Motion Regressors
  - Motion Censoring

# Retrospective Motion Correction

- Best: Avoid subject motion in the first place
- Better: Use Prospective Motion Correction
- Standard: Perform rigid-body realignment, use parameters as nuisance regressors
  - 6 parameters: translation+rotation
  - 12 parameters: include derivatives (for temporal shifts)
  - 24 parameters: include squared regressors
- 24-parameter model known as Volterra expansion

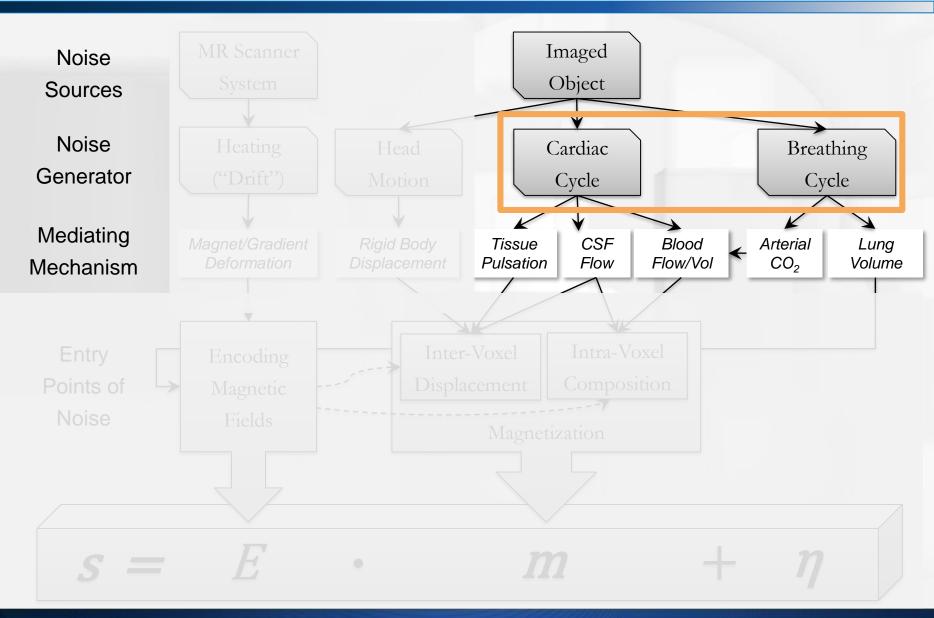
Friston, MRM, 1996

# Motion Censoring = "Scrubbing"

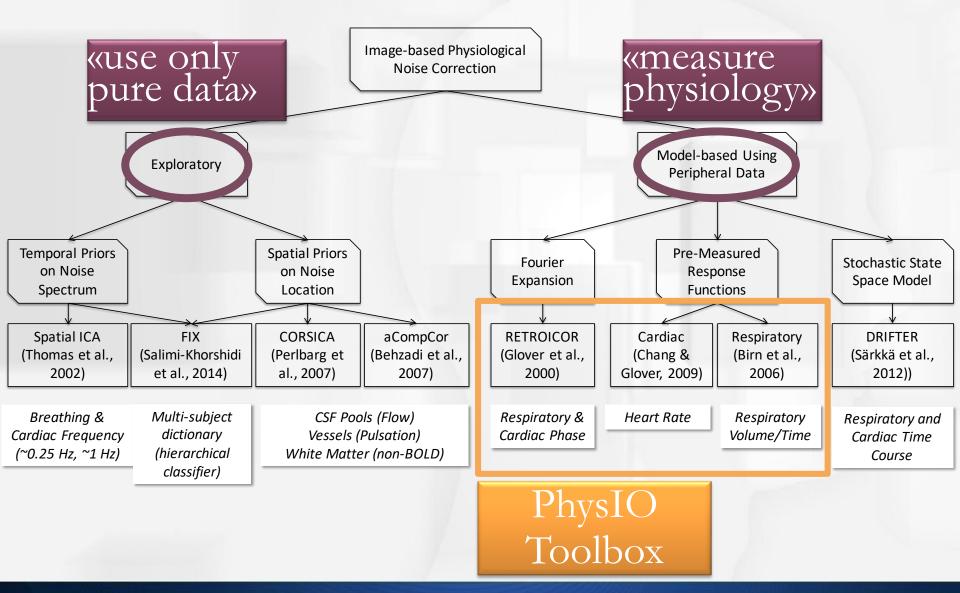
- Detect outlier volumes (strong movement, but also spikes, RF flip angle fluctuations)
- Inform the GLM of these bad volumes via stick regressors (zero everywhere else, 1 at volume)
  - Will absorb all variance of that volume
- Problem: Temporal filtering before GLM might create
   Gibbs ringing of outliers into neighbors
- Alternative: censoring during preprocessing
  - interpolate faulty volume by neighbors

### **Noise Correction Targets**





### **Image-based Noise Correction**



# **Noise Modeling**



 RETROspective Image CORrection
 Cardiac Response Function
 Respiratory Response Function

 • Cardiac/respiratory
 • Heart Rate
 • Resp. Volume

phase  $\varphi_c \quad \varphi_r$  per Time

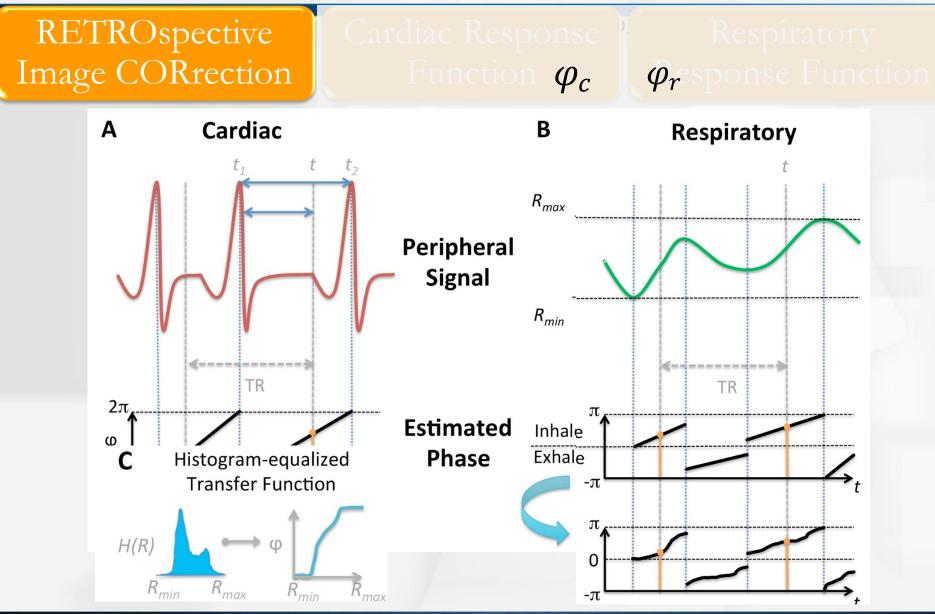
Fourier expansionconvolved withconvolved with(cosine/sine)CRFRRF

evaluated at 1 time point (slice) per volume =

regressor

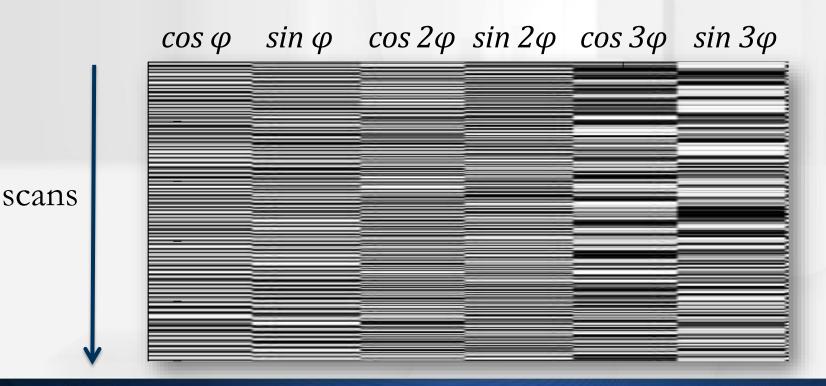
# **Noise Modeling**





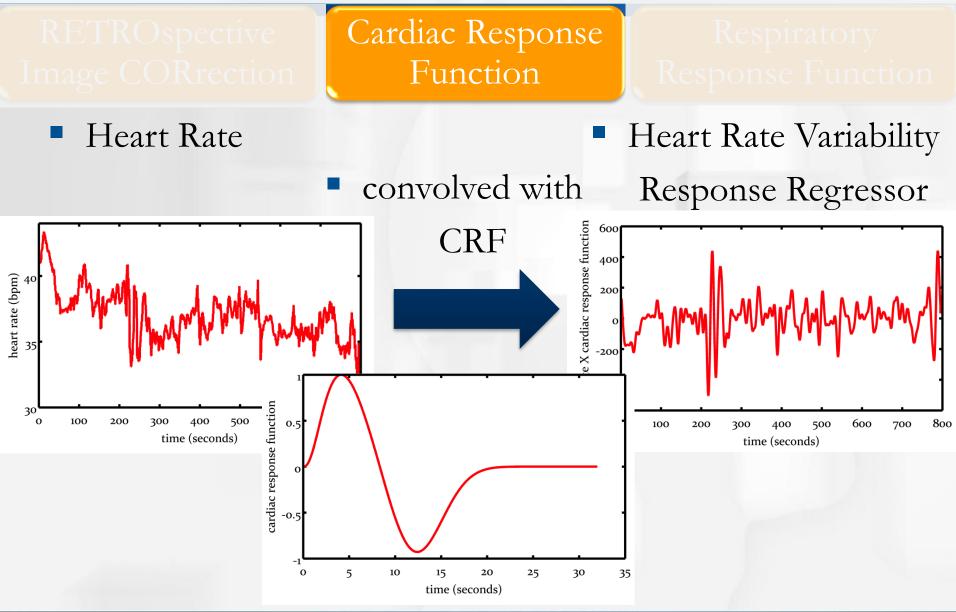
# Model: Fourier Phase Expansion

- Cosine and sine to allow for constant phase shifts per voxel
- Higher model orders to account for under-sampling of physiological frequencies with typical TR in fMRI



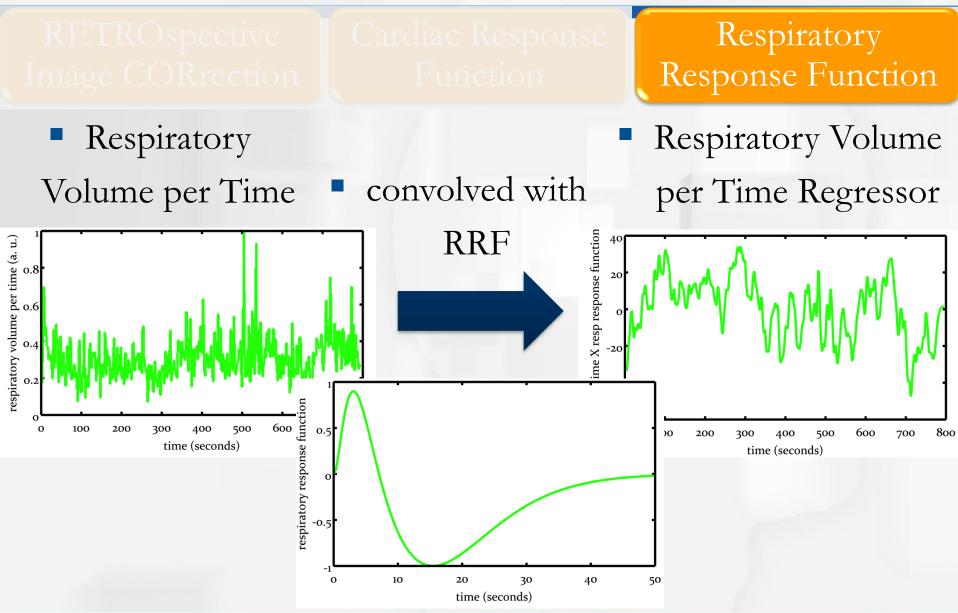
## **Noise Modeling**



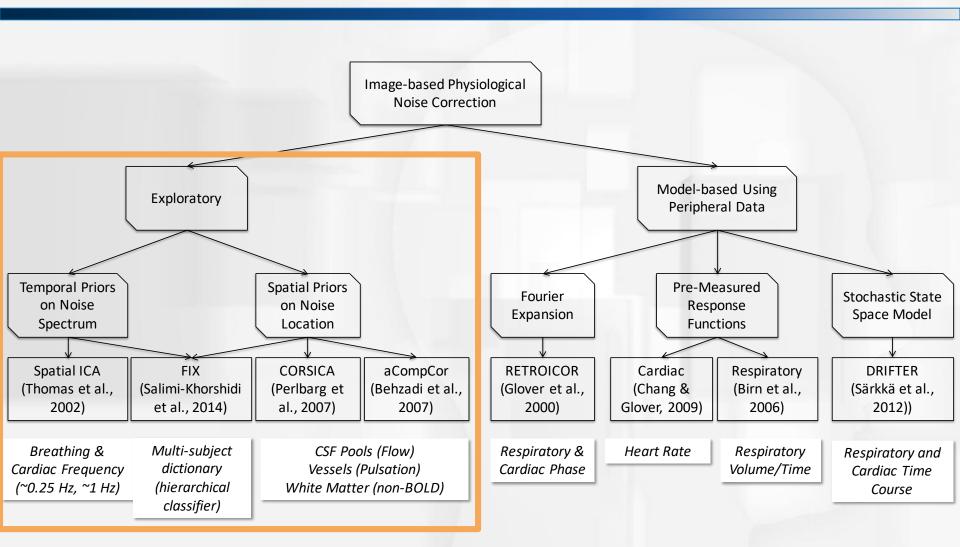


# **Noise Modeling**





#### **Exploratory Phys Noise Correction**



#### **Noise Component Modeling**



- Use priors about physiological noise to identify noise components (time series)
  - Spatial Priors: Mechanisms of physiological noise implicate physiological noise in CSF, blood vessels
  - Temporal Priors: Knowledge about typical physiological frequency contents (heart ~ 1Hz, breathing 0.2-0.4 Hz)
    - Note that simple filtering is impossible (cf. aliasing)
  - Population Priors: Use dictionary learning from manually labelled training set of subjects (FIX)

¥		V	
Spatial ICA	FIX	CORSICA	aCompCor
(Thomas et al.,	(Salimi-Khorshidi	(Perlbarg et	(Behzadi et al.,
2002)	et al., 2014)	al., 2007)	2007)
Breathing &	Multi-subject	CSF Pools (Flow) Vessels (Pulsation) White Matter (non-BOLD)	
Cardiac Frequency	dictionary		
(~0.25 Hz, ~1 Hz)	(hierarchical		
· · · ·	classifier)		· · ·
	, ,		

## PCA VS ICA



- Methods to extract components (i.e. summarize ROIs/spectra) differ:
  - Maximum variance time series: Principal Component Analysis (PCA) from region of interest (aCompCor, Behzadi 2007)
  - Maximally independent time courses/sites: spatial/temporal ICA, FSL MELODIC, FIX
- aCompCor is basically identical to a seed-based correlation analysis in resting-state fMRI
  - Here: seed is in region-of-no-interest and correlated time series regressed out
  - See previous talk (resting state analysis) for more details

## Other Physiological Corrections

- Non-linear models
  - DRIFTER: Kalman Filter, Bayesian, *Joint* Stochastic State-space model of peripheral physiology and BOLD
- Identify noise via task test-retest reproducibility
  - PHYCAA: e.g. via high-freq. autocorrelation, anatomy
  - GLMDenoise: PCA of noise regressors
- MEICA: Multi-Echo ICA
  - Use diff. TE-images to decompose proton density from T2\* changes

Särkkä, Neurolmage, 2012 Churchill, Neurolmage, 2012/13 Kay, Front. Neurosc., 2013 Olafsson, Neurolmage, 2015

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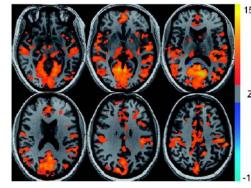
#### When? – Literature Evidence



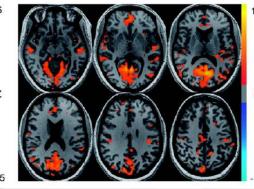
#### Resting-state:

- Birn, R. M. "The Role of Physiological Noise in Resting-state Functional Connectivity." *NeuroImage 62*, 2012
- Birn, R. M., et al. "Separating Respiratory-variation-related
  Fluctuations from Neuronal-activityrelated Fluctuations in fMRI." *NeuroImage 31*, 2006

C Resting-state correlation

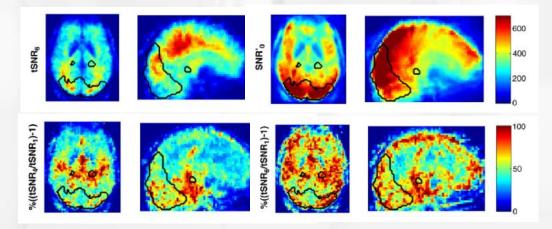


D Rest-state corr – after RVTcor



#### Task-based:

Hutton, C., et al. "The Impact of Physiological Noise Correction on fMRI at 7 T." *NeuroImage 57*, 2011:



#### All these methods, but...



 Physiological noise correction not a default preprocessing step in task-based fMRI

Reasons

- Impact on group level fMRI
  - no reports for non-trivial paradigms
- Existing Toolboxes lack...
  - robust, automatic implementation
  - dealing with variable peripheral data quality

# Paradigm: Learning from Advice

- Hierarchical learning of trustworthyness of advisor over time
- Contrasts: Prediction and Prediction Error about advice



recommendations of adviser were **veridical** (pre-recorded videos from behavioural study)

**volatility of advice** (changing intentions of adviser through incentive structure)

interactive, gender-matched (**40** male subjects)

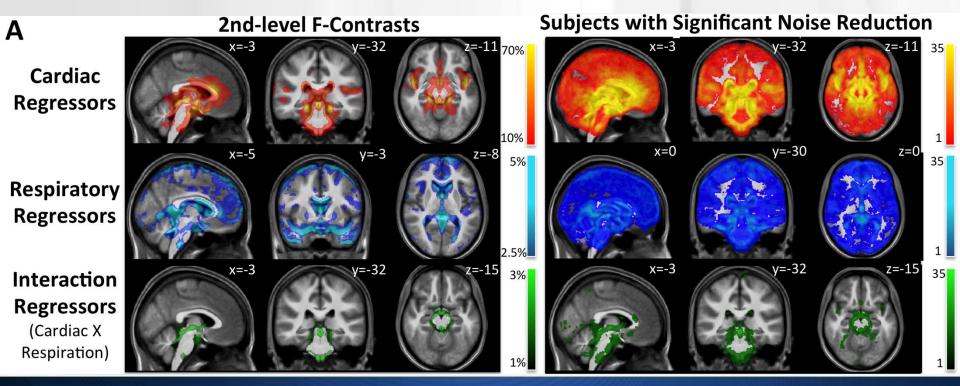
**fMRI design**: Philips Achieva 3T TR/TE 2500/36ms, 2 x 2 x 3 mm<sup>3</sup>

Diaconescu et al, 2014, PLoS Comp. Biol.

## **Group Level Impact PhysIO**

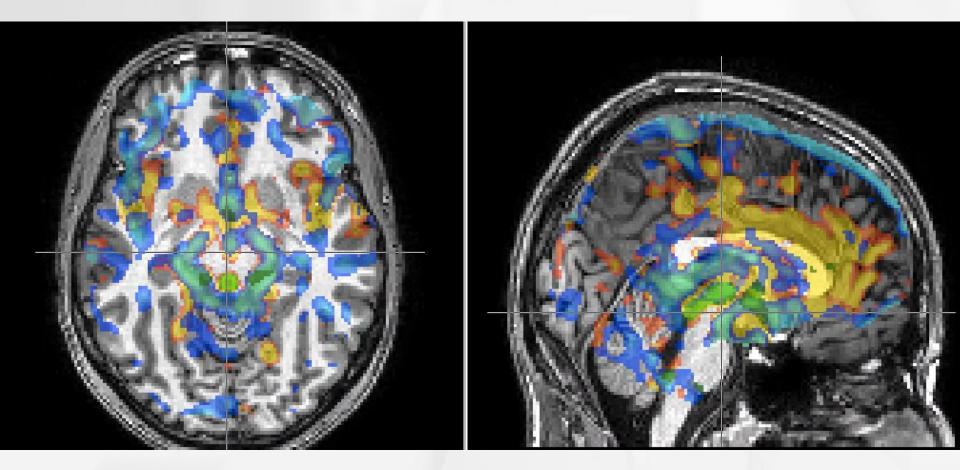


- Andreea Diaconescu (TNU): Social Learning Experiment 2012-2014, (N=35)
- F-contrast: Where does physiological noise model explain significant variance?



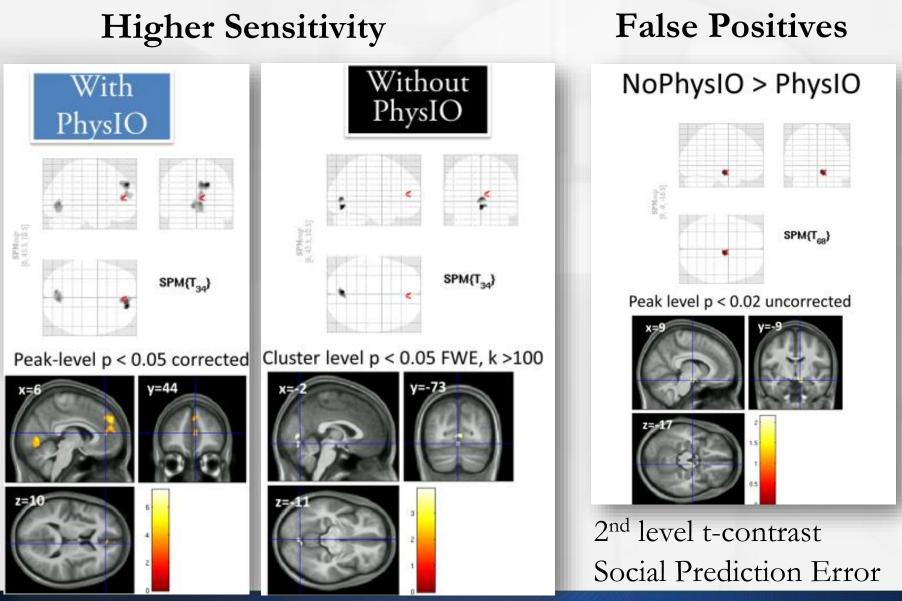
#### Relevance for Neuromodulation

#### cardiac (red), respiratory (blue), cardXresp (green)



## **Effects on Group Contrasts**





Matthias Müller-Schrader: fMRI Noise Models & Correction

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#### **Limitations of Noise Modeling**

- Degrees of freedom, sensitivity reduced by too many ineffective regressors
  - F-test informative
- Intrinsic correlations of functional areas of interoception and peripheral physiology
  - E.g. Amygdala, Insula, ACC
  - Controversial reading:
     <u>fMRI of the Amygdala: All In Vein? Neuroskeptic</u>
  - Alternative: Masking, Pure anatomical priors removing CSF, angiography (vessels)

#### Conclusion

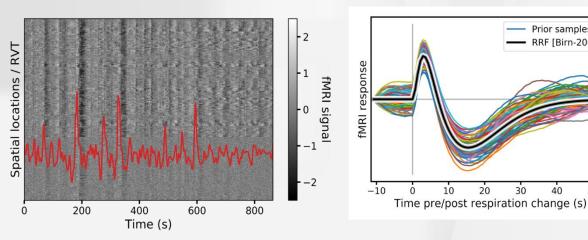


- MRI Time Series and Physiological Noise
- Image-Based Correction in the GLM
- Noise Modeling Prospects: Group FX

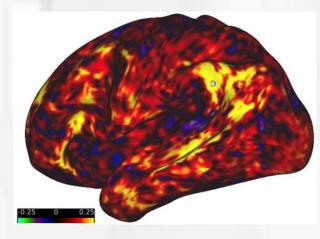
The PhysIO Toolbox

- Structured noise through cardiac/resp cycle (70%)
- Nuisance regressors from
   Fourier expansion, response
   functions
- Increase group sensitivity (low inter-subject variability), fewer false positives
- Correction in SPM/Matlab in practice => NOW!

- "Subject-specific physiological noise removal with deep Bayesian inference"
  - Semester / Master thesis
  - Start early next year
  - harrison@biomed.ee.ethz.ch









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Prior samples

RRF [Birn-2008]

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#### Thank you for your attention



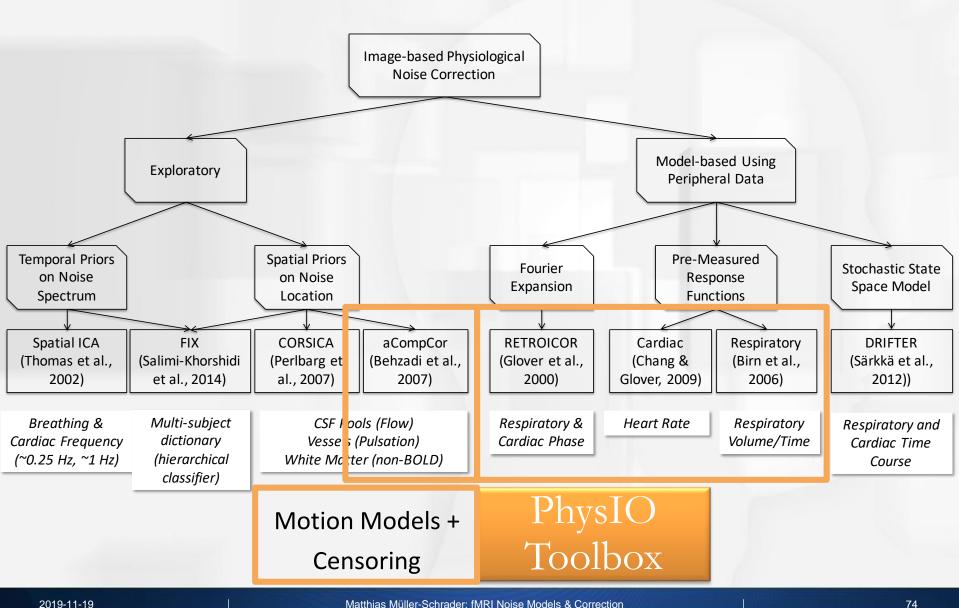
- ... and
  - TNU
  - Lars Kasper





- Demo: The PhysIO Toolbox for Physiological Noise
   Correction in fMRI
  - Features and Workflow
  - Image-based physiological noise correction in the GLM
    - RETROICOR, HRV, RVT
    - Noise-ROIs
  - Practical Demo (SPM Batch)
    - Estimating different Models
    - Understanding the Preprocessing Plots
    - Automatic Model Assessment, Diagnostics on Contrast

#### **Image-based Noise Correction**



## The PhysIO Toolbox

- Developed at the Translational Neuromodeling Unit (TNU) since 2008
  - Lead programmer: Lars Kasper (TNU)
  - Contributors: Jakob Heinzle (TNU), Steffen Bollmann (KiSpi Zurich)
- Part of the TNU «TAPAS» software suite
- Used at the TNU, in Zurich and beyond by ~50 researchers
  - Iglesias 2013, Neuron; Kasper 2014, NeuroImage; Bollmann 2014, PhDThesis; Sulzer 2013, NeuroImage; Hauser 2014, NeuroImage; Grueschow 2015, Neuron
- Download & Example Data:
  - <u>https://www.tnu.ethz.ch/en/software/tapas.html</u>
  - <u>https://www.tnu.ethz.ch/en/software/tapas/data.html</u>

# **Installing TAPAS**



Download as zip:

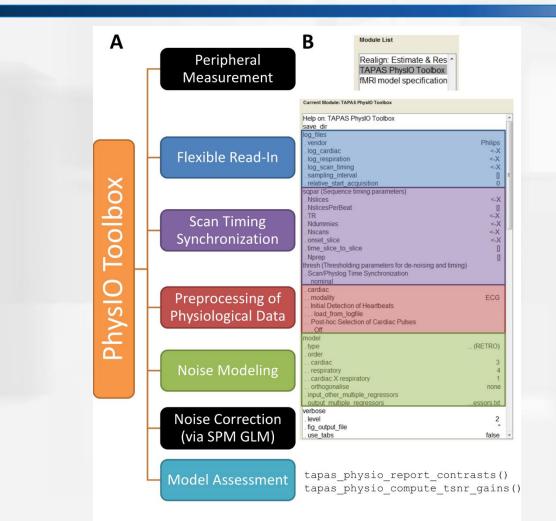
https://translationalneuromodeling.github.io/tapas/#do wnload

• Git/SVN:

https://github.com/translationalneuromodeling/tapas.git

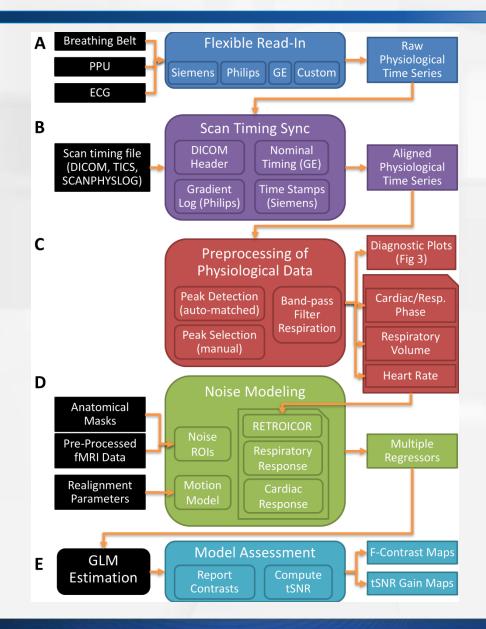
- Run tapas/tapas\_init.m (adds tapas folder to path)
- Run tapas\_physio\_init.m (for SPM integration)

## Workflow of the PhysIO Toolbox **C**

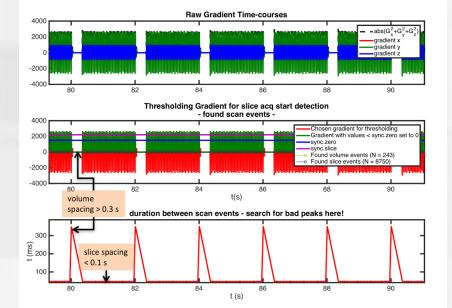


#### Flowchart of Noise Correction

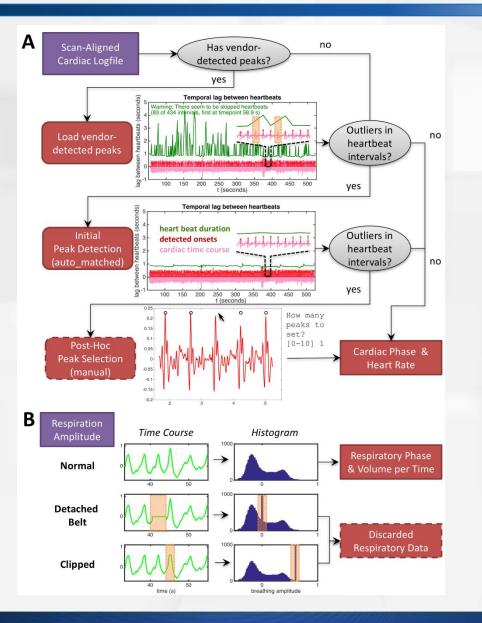




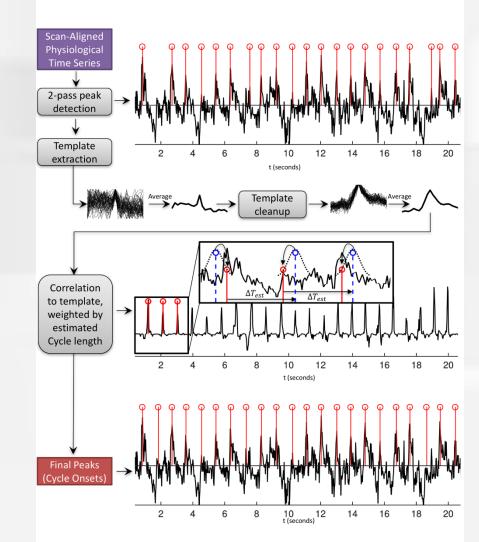
## Scan Sync with Philips Gradients



#### **Data Preprocessing Overview**

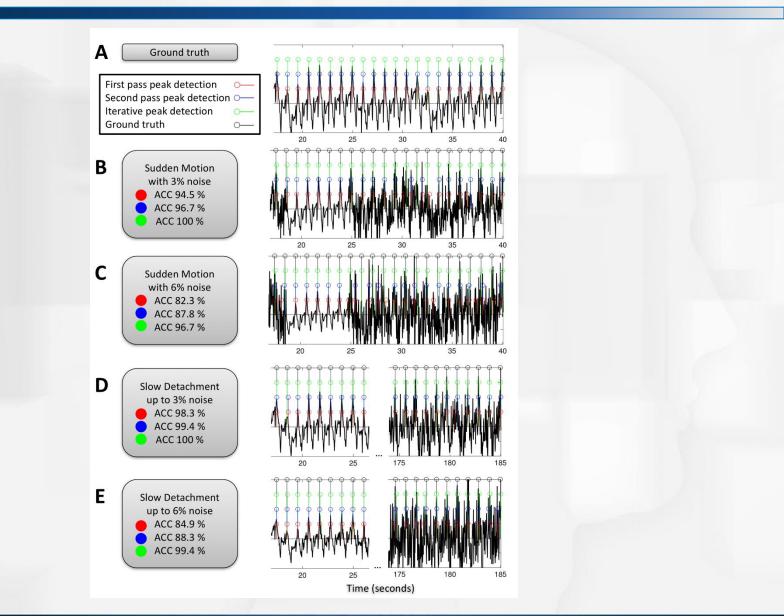


#### **Preprocessing: Peak Detection**



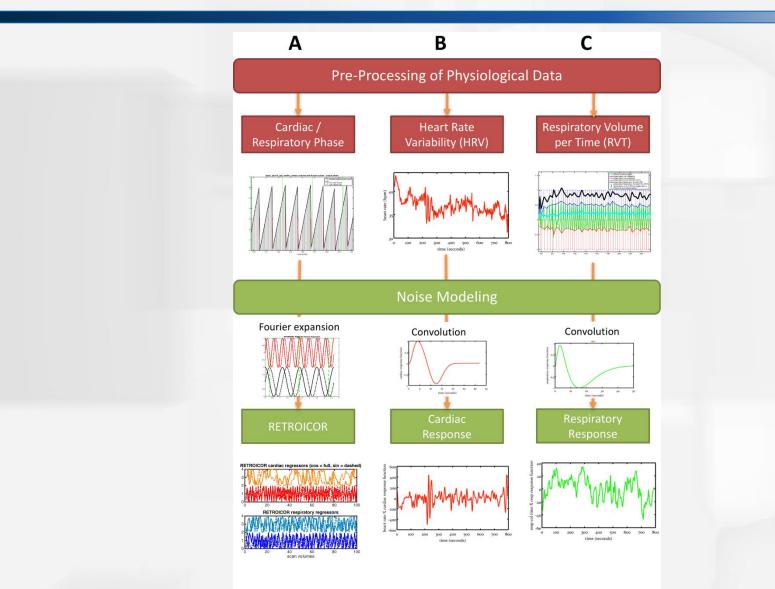
#### **Peak Detection: Robustness**





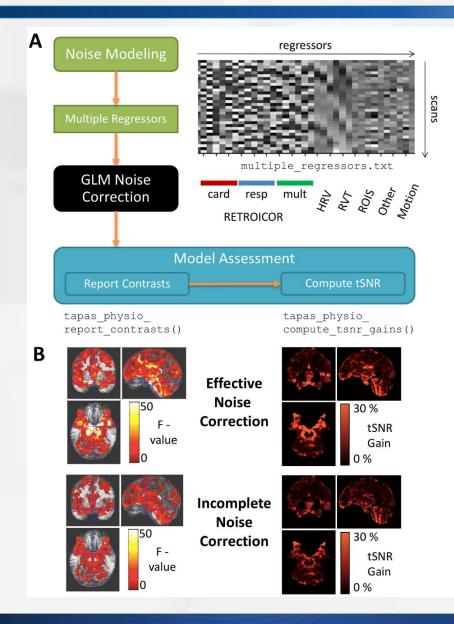
#### **Noise Modeling**





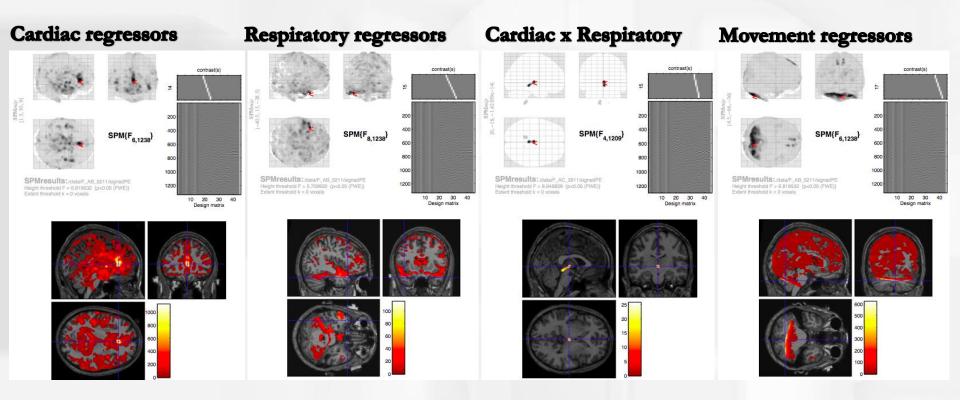
#### **Diagnostics: Model Assessment**





#### **Model Check: SPM F-contrasts**





Check Influence of Physiological Noise (Correction) on Data

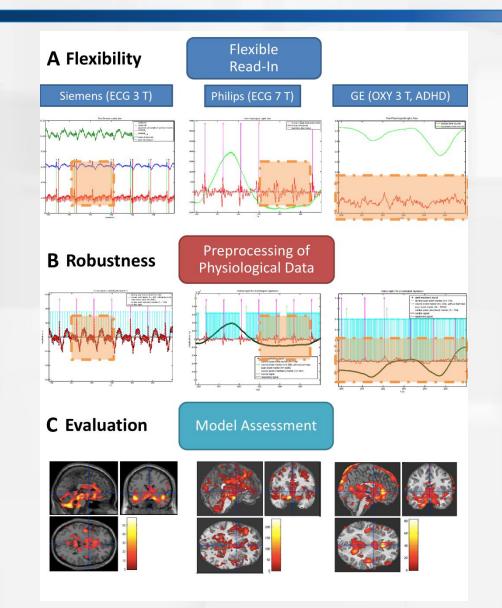
• SPM

Finally:

F-contrast on 1st and second level

#### Flexibility: Scanner vendors





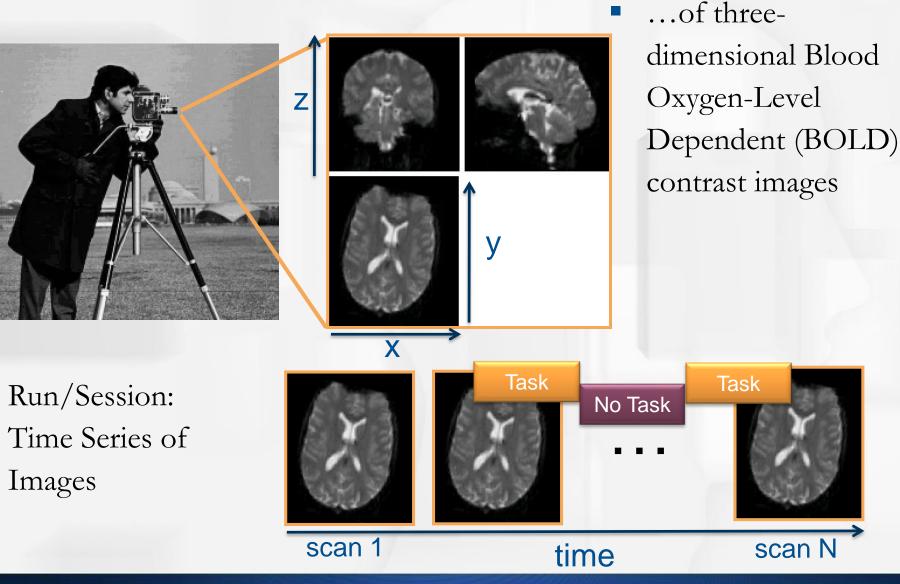
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## fMRI = Acquiring Movies





Matthias Müller-Schrader: fMRI Noise Models & Correction

#### Help for the exercises



#### Old version of an batch. OR: physio = tapas\_physio\_new();

