

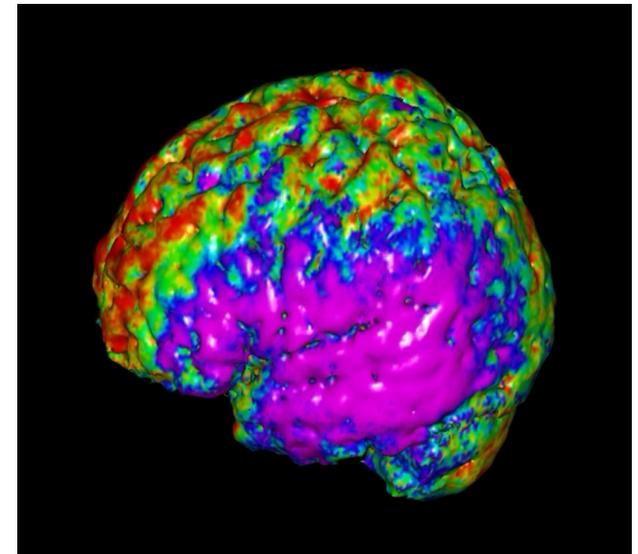
# Smoothing Images from Population Studies

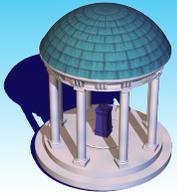
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**Hongtu Zhu, Ph.D.**

**Department of Biostatistics and Biomedical  
Research Imaging Center**

**University of North Carolina at Chapel Hill**





# Outline

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**Motivation**

**Smoothing Strategies**

**Multiscale Adaptive Smoothing Models**

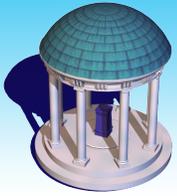
**Simulation Studies**

**Real Data Analysis**



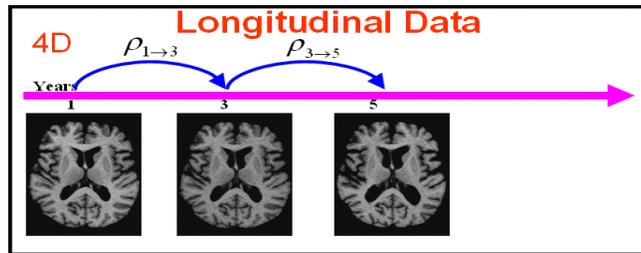
**UNC**  
SCHOOL OF  
PUBLIC HEALTH



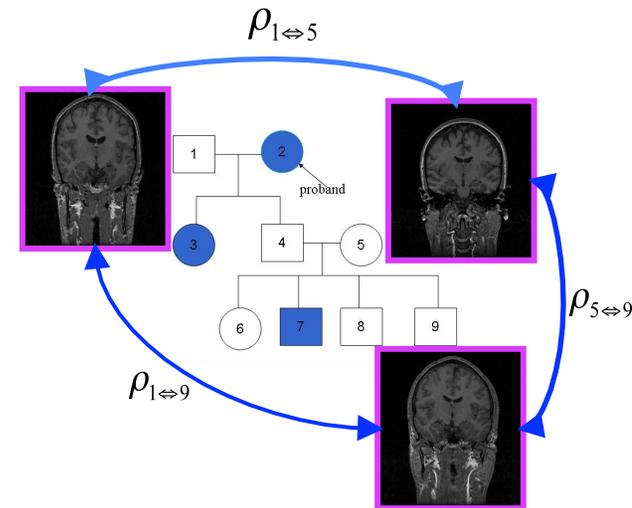
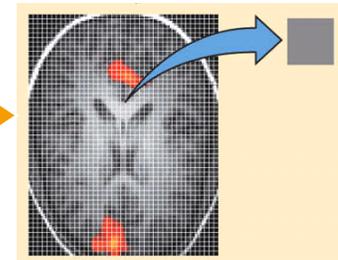
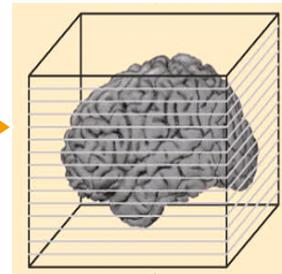


# Motivation

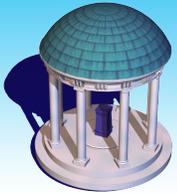
## Study Design



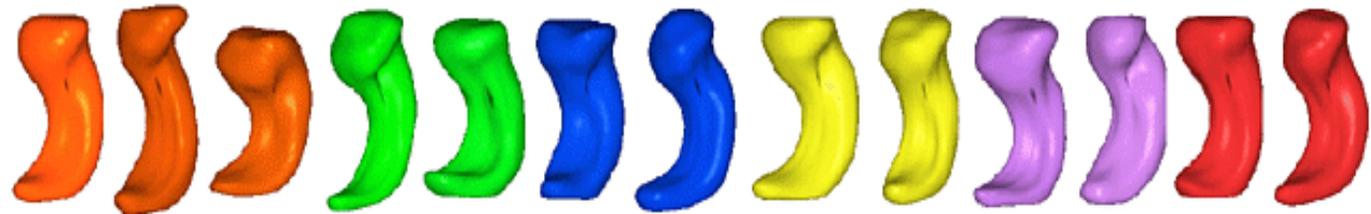
## Neuroimaging Data



[www.guysandstthomas.nhs.uk/.../T/Twins400.jpg](http://www.guysandstthomas.nhs.uk/.../T/Twins400.jpg)



## Motivation



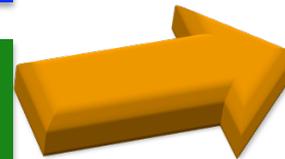
Raw Images



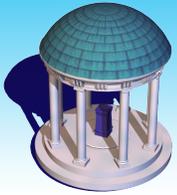
Image  
Reconstruction

Image  
Registration

Imaging  
Smoothing



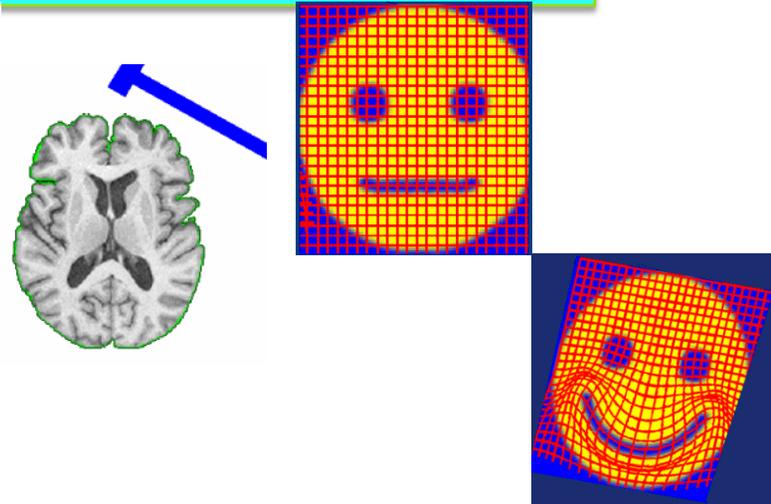
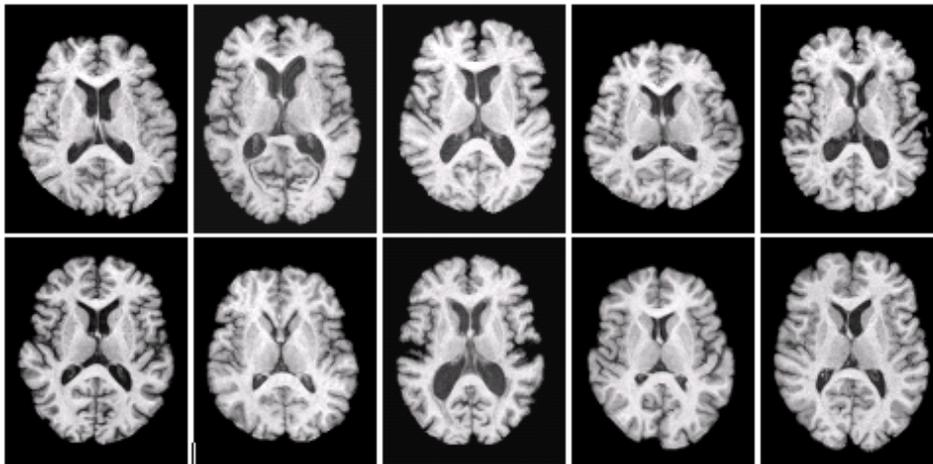
**Statistical  
Analysis**

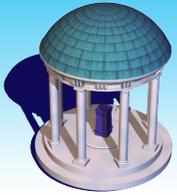


# Motivation

## Image Registration

Image registration is the process of **transforming different sets of data into one coordinate system**. Data may be multiple photographs, data from different sensors, from different times, or from different viewpoints.



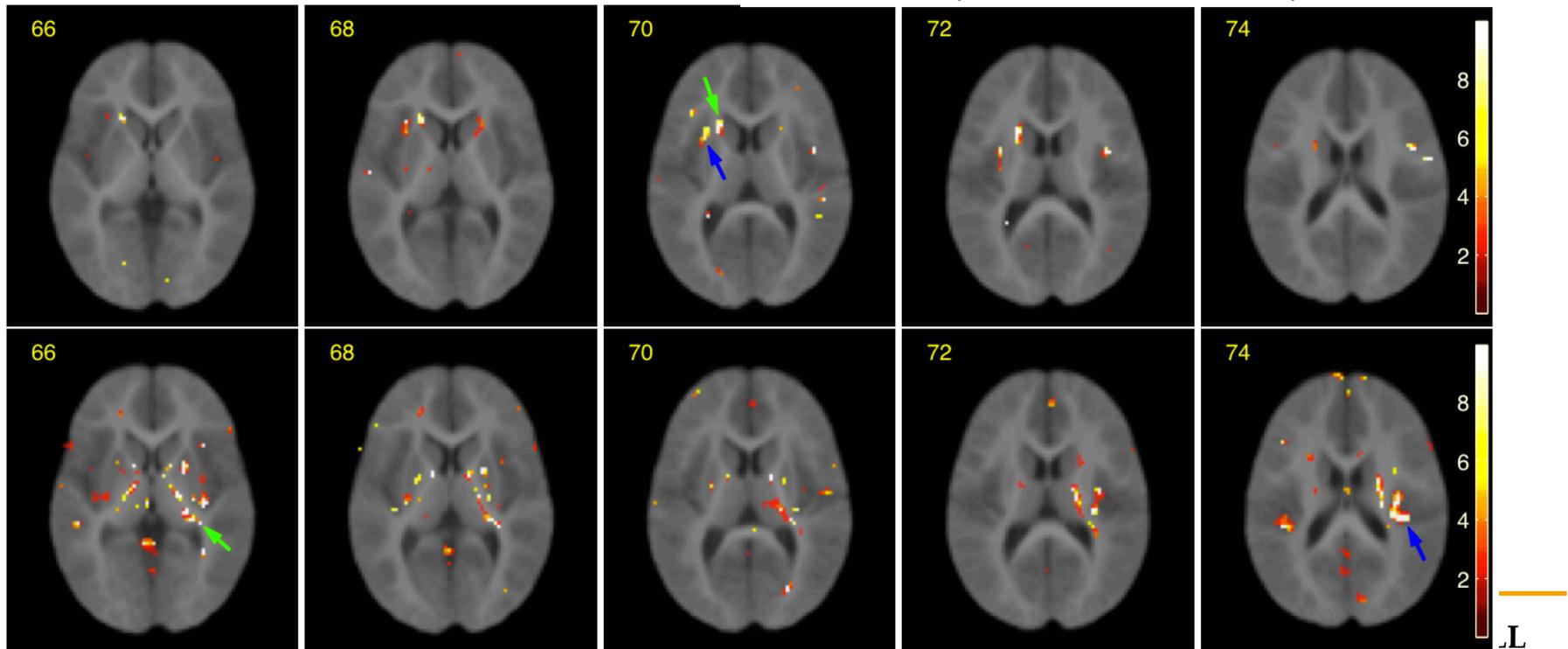
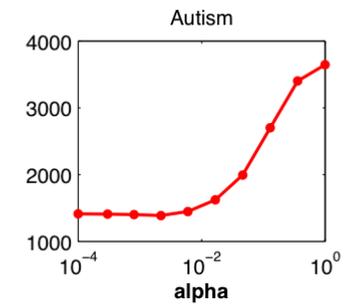
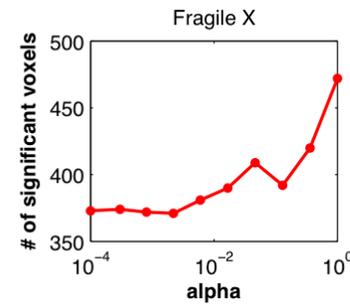


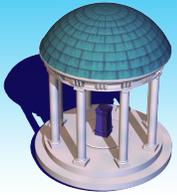
# Motivation

25 normal control (NC), 19 fragile X syndrome (FX), and 47 autism (AU)

FX vs NC

AU vs NC

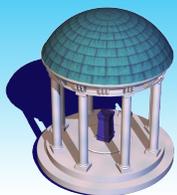




# Motivation

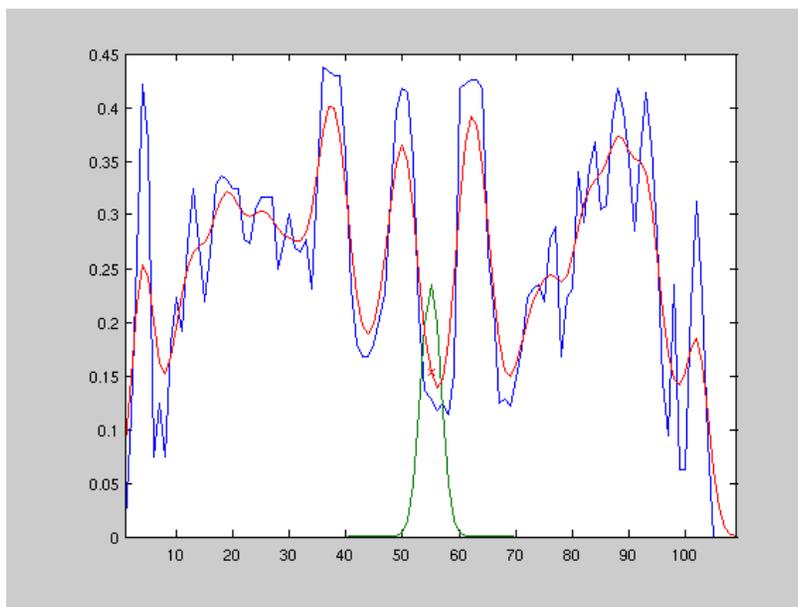
## Image Smoothing

- ◆ Registration
- ◆ Signal-to-noise Ratio
- ◆ Gaussian
- How is it implemented?
  - ◆ Convolution with a 3D Gaussian kernel, of specified Full-width at half-maximum (FWHM) in mm



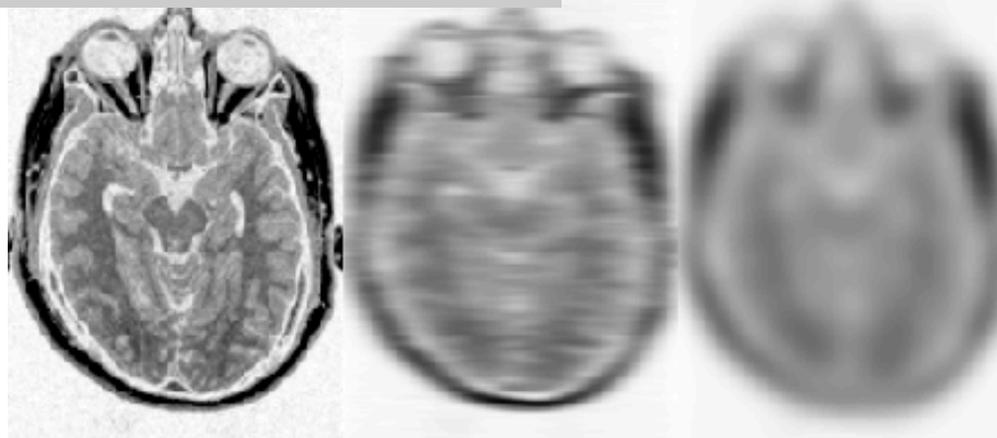
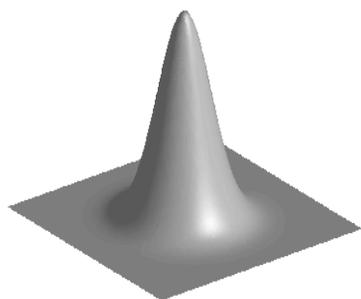
# Motivation

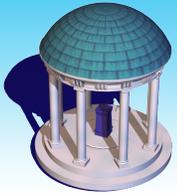
## Example of Gaussian smoothing in one-dimension



The Gaussian kernel is **separable** we can smooth 2D data with two 1D convolutions.

SPM training course

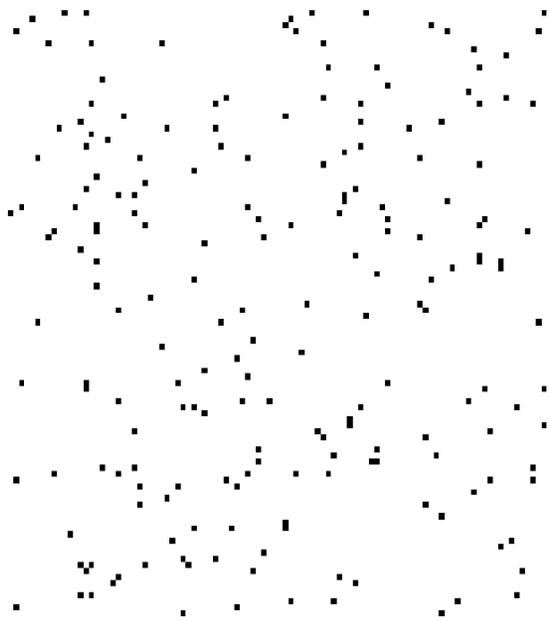




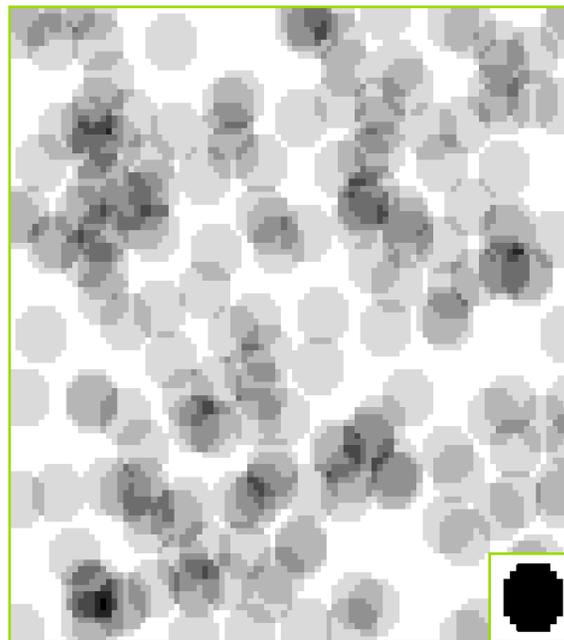
# Motivation

Each voxel after smoothing effectively represents a weighted average over its local region of interest (ROI)

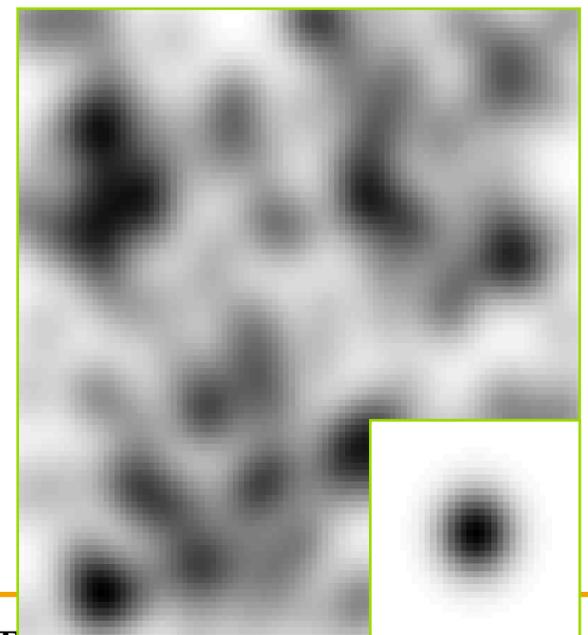
Before convolution

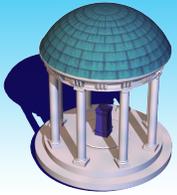


Convolved with a circle



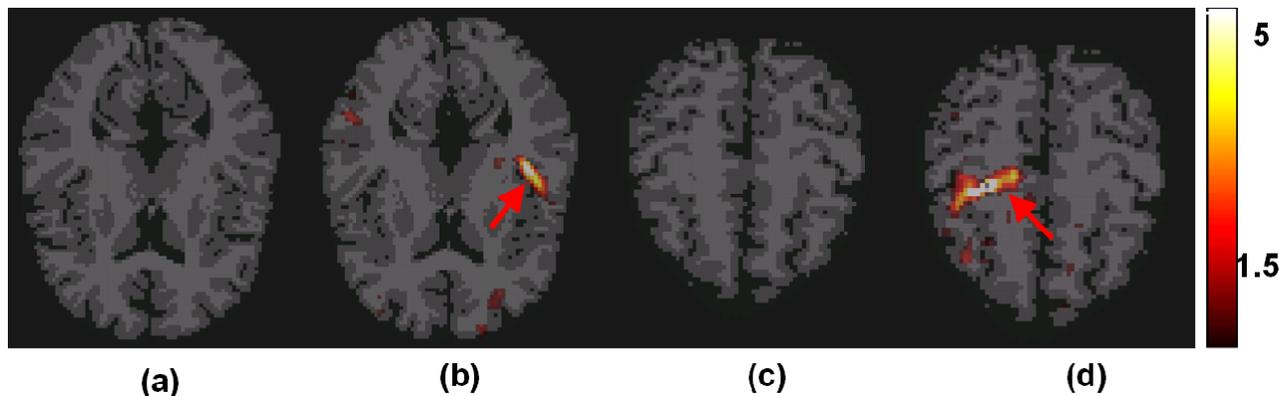
Gaussian convolution



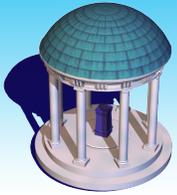


# Motivation

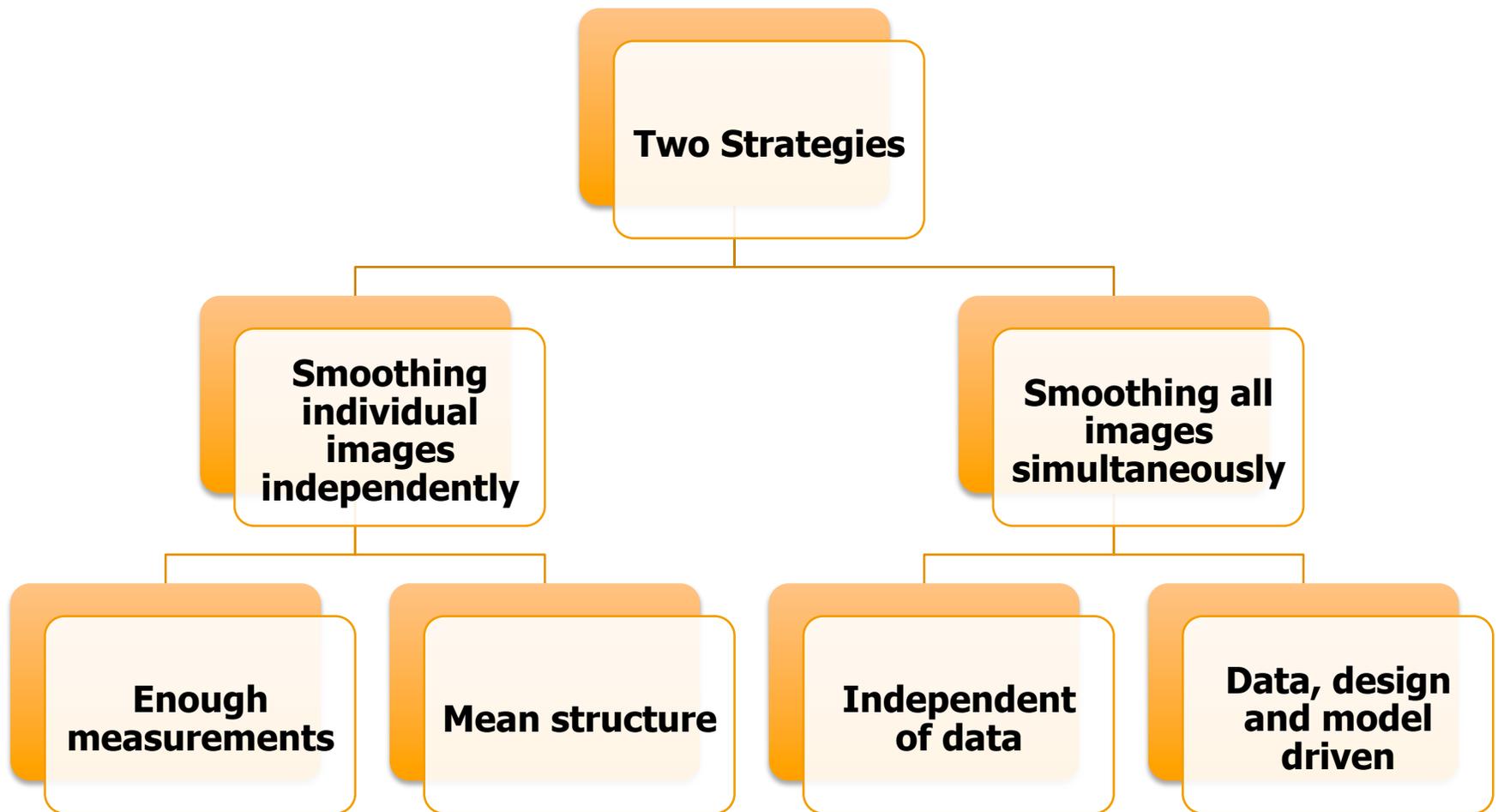
- Smoothing method is independent of **data**
- Degree of smoothness is **arbitrary**
- Effect of smoothness is **profound**
- The relationship between smoothing method and study design is **unknown**

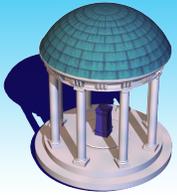


Jones et al. (2006),  
Yue et al. (2010)



# Smoothing Methods





# Smoothing Methods

## What is image?

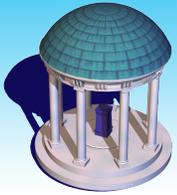
*an exact **replica** of the contents of a storage device*



*an optically formed **duplicate** or other reproduction of an object*

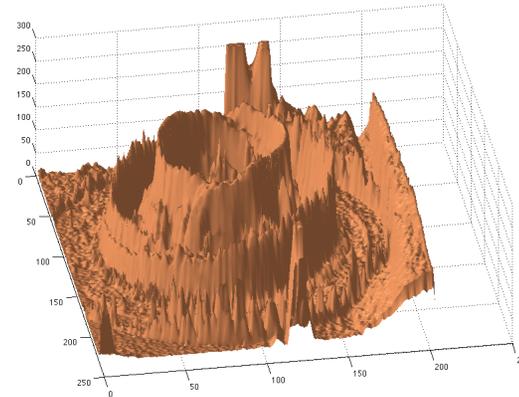


Google/wiki



# Smoothing Methods

**Mathematics.**

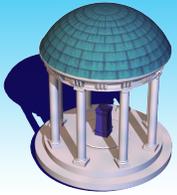


*Image is the point or set of points in the range corresponding to a designated point in the domain of a given function.*

▲  $\Omega$  is a compact set.  $\tilde{x} \in \Omega \subseteq \mathbb{R}^k$

➔  $f(\tilde{x}) \in M \subseteq \mathbb{R}^m$        $f : \Omega \rightarrow M \subseteq \mathbb{R}^m$

★  $\int_{\Omega} \|f(\tilde{x})\|^k d\tilde{x} < \infty$  for some  $k > 0$



# Smoothing Methods

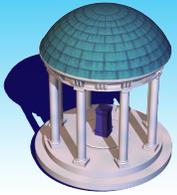
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## Digitized Images

$$f : \Omega_0 \rightarrow \{0, 1, \dots, M_0\}$$

- **Sampling (grid points)**  $\Omega_0 \in \Omega$
- **Sampling Rate**
- **Quantization**

$0, 1, 2, \dots, 2^m$  for  $m = 5 \sim 12$ , that is  $M_0 = 2^m$

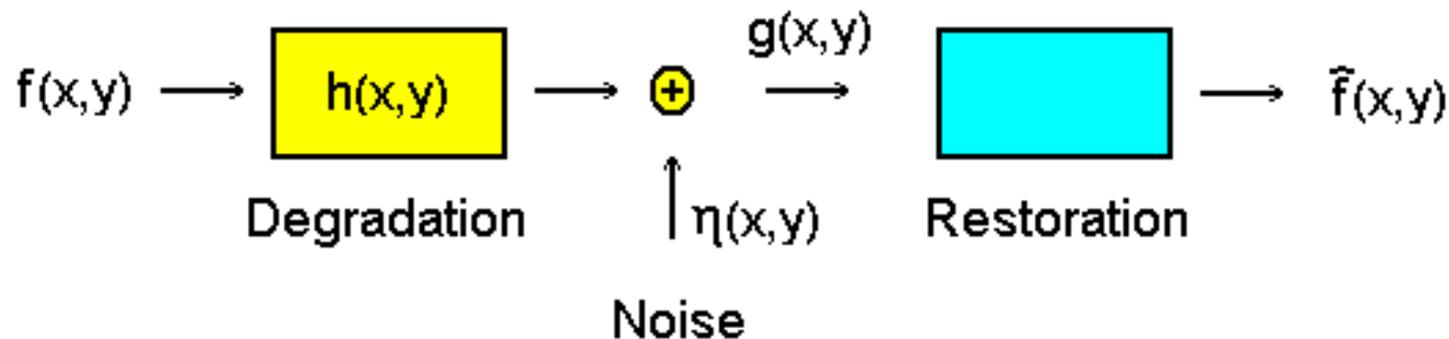


# Smoothing Methods

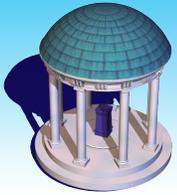
## Image Degradation/Restoration Process

The goal of image restoration is to improve a degraded image in some predefined sense. Schematically this process can be visualized as

$$g(x, y) = h(f(x, y)) + \eta(x, y)$$



where  $f$  is the original image,  $g$  is a degraded/noisy version of the original image and  $\tilde{f}$  is a restored version.



# Smoothing Methods

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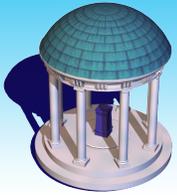
## Methods

- **Filters**  
Lowpass filtering, Wiener filters, Median filtering
- **Wavelet Shrinkage Denoising**  
Soft and Hard thresholding
- **Variational Denoising based on Bounded Variation Models**
- **Nonlinear Diffusion and Scale-space theory**
- **Bayesian Models**  
Markov random field

## Issues:

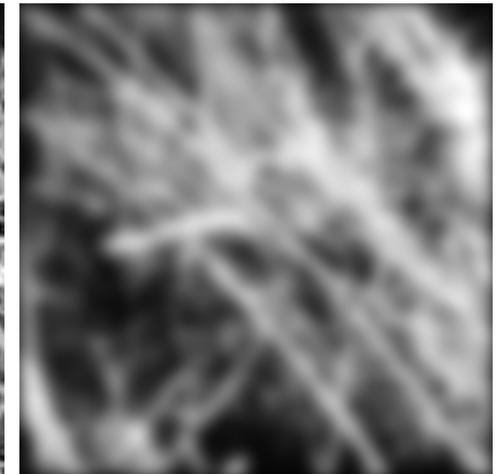
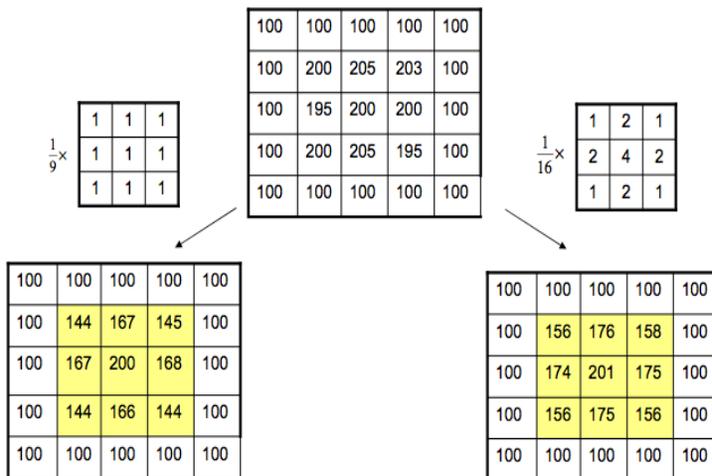
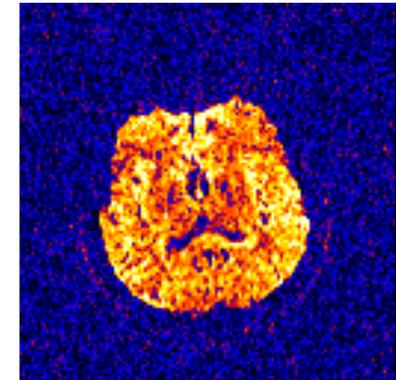
- **Noise Distribution**
- **Window Size**
- **Localization**
- **Tuning Parameters**

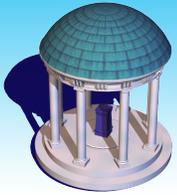
[Chen and Shen \(2005\)](#)



# Smoothing Methods

- An image may be 'dirty' with dots, speckles, stains
- Noise removal
  - Dots can be modeled as impulses (salt-and-pepper or speckle) or continuously varying (Gaussian noise)
  - Low-pass filtering
- Problem with low-pass filtering
  - May blur edges
  - Adaptive, edge preserving





# Smoothing Methods



original image



1px median filter



3px median filter

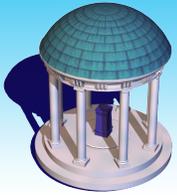


10px median filter

## Bandwidth Selection

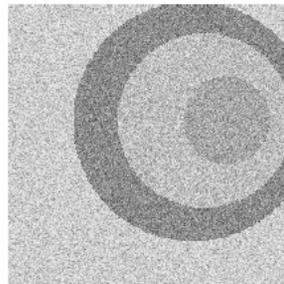


## Different Smoothing Methods

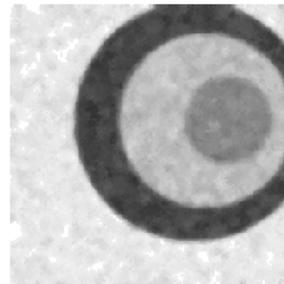


# Smoothing Methods

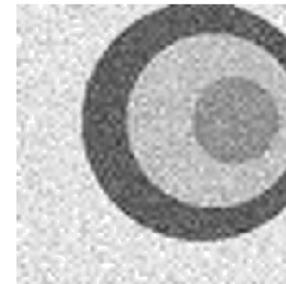
## Location Adaptation



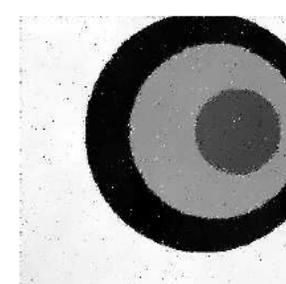
(a)



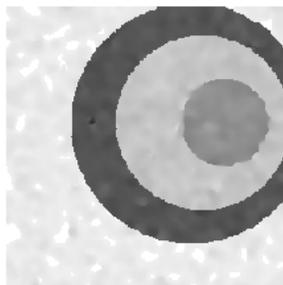
(b)



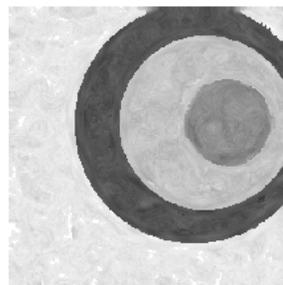
(c)



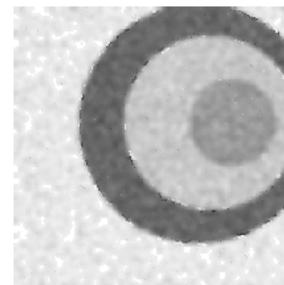
(d)



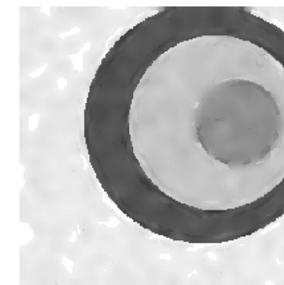
(e)



(f)



(g)

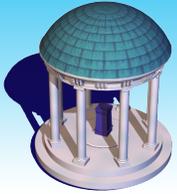


(h)

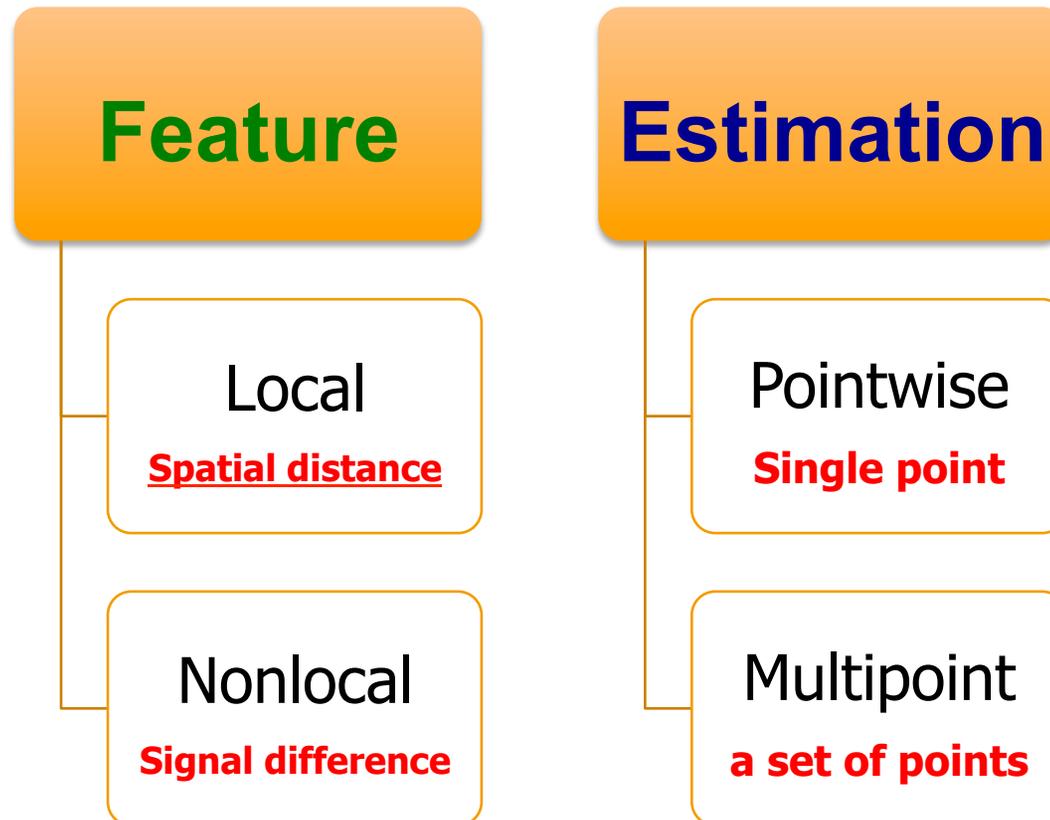
## Different Smoothing Methods

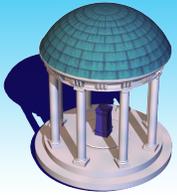
Figure Plot (a): The noisy test image. Plots (b)-(h): the reconstructed images by the local median smoothing procedure, the DWT procedure, the MRF procedure, the AWS procedure, and procedures (6)-(8), respectively.

**Qiu (2005)**



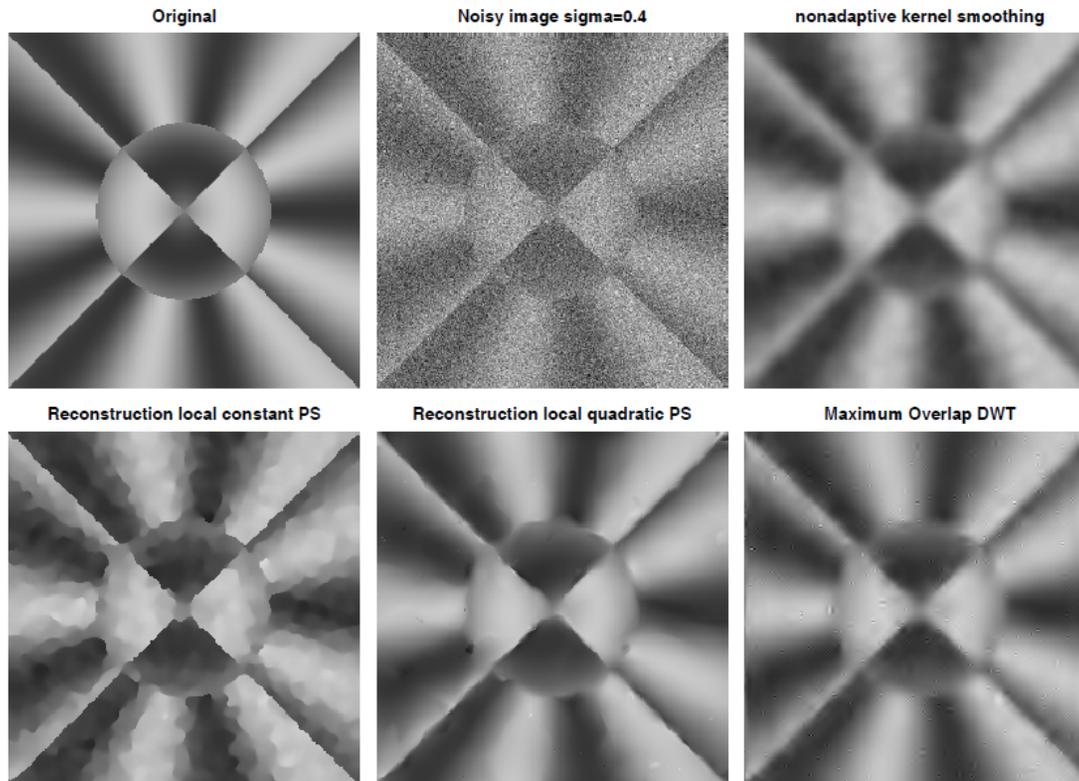
# Smoothing Methods





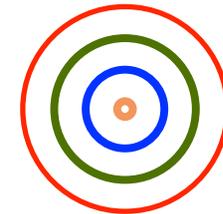
# Smoothing Methods

## Propagation-Separation Method

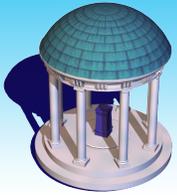


## Features

- Increasing Bandwidth



- Adaptive Weights
- Adaptive Estimates



# Smoothing Methods

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## Propogation-Seperation Theory

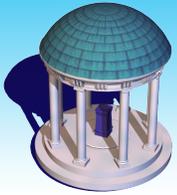
- **Exponential Family**  $Y(d) \sim EF(\theta(d))$

$$\hat{\theta}(d) = \operatorname{argmax}_{\theta(d)} \sum_{d'} w(d, d') \ell(Y(d'), \theta(d)) = \operatorname{argmax}_{\theta(d)} \ell\left(\sum_{d'} \tilde{w}(d, d') Y(d'), \theta(d)\right)$$

Smoothing Imaging Intensities

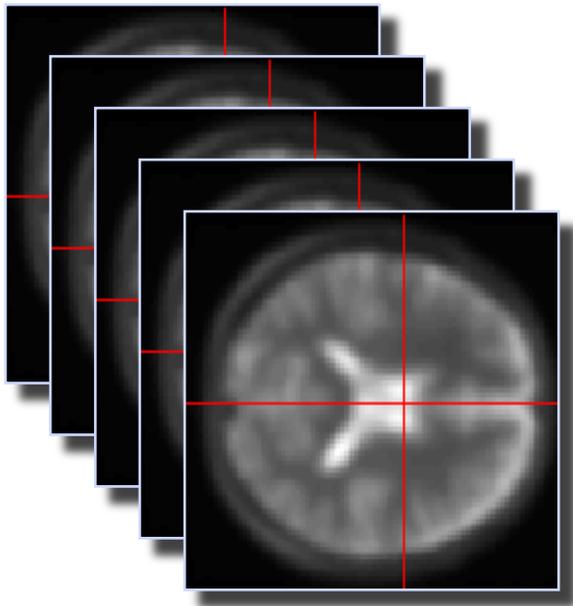
- **Under strong conditions,** **Katkovnik, V and Spokoiny, V. (2008)**  
**J. Polzehl and V. Spokoiny, (2005)**

$$P(K(\hat{\theta}(d), \theta(d))^{1/2} > C(\log(N(D)) / N(D))^{1/(2+c)}) \rightarrow 0$$



# Smoothing Methods

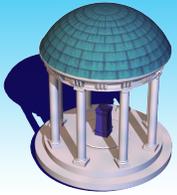
Images from Multiple Subjects  
Multiple Images from a single subject



Tabelow et al. (2006, 2008a, 2008b),  
Polzehl, et al. (2010)

- Denoising fMRI, DTI
- SPM

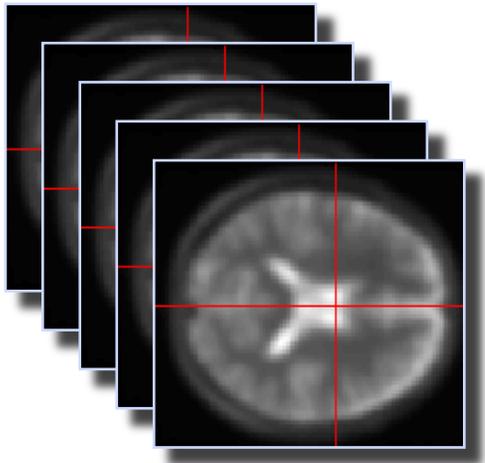
$$Y_i(d) = x_i^T \beta(d) + \varepsilon_i(d) \sigma(d)_i$$



# Multiscale Adaptive Smoothing Models

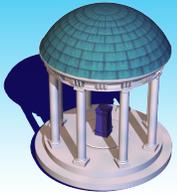
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## Images from Multiple Subjects



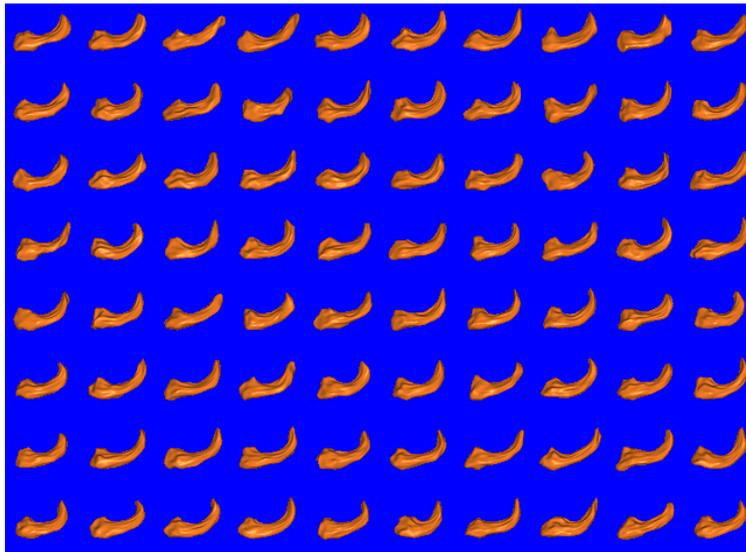
## Questions of Interest

- **Complex design**  
**Longitudinal, Twin, and family studies**
- **Models with parametric and/or nonparametric components**
- **Consistency results**
- **Standard deviation images**
- **Testing theory**



# Multiscale Adaptive Smoothing Models

My goal is to develop a class of MASMs with necessary statistical properties for imaging data collected from cross-sectional, longitudinal, twin, and familial studies.



$M$

$$= g(x, \theta(d), f(d)) \oplus \varepsilon$$

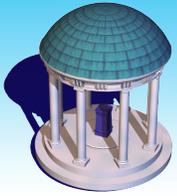
$$x \in R^k, \theta(d) \in \Theta \subset R^p, f(d) \in F$$

$$g: R^k \times R^p \times F \rightarrow M$$

**Problems of interest:**

$$\{(\theta(d), f(d)) : d \in D\}$$

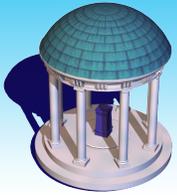
$$\{\varepsilon(d) : d \in D\}$$



## **MARM**

# **Multiscale Adaptive Regression Models**

- **Integrate Parametric Models with PS**
- **Standard Deviation Image**
- **Consistency and Asymptotic Distribution**



# Multiscale Adaptive Regression Model

$D$ : 3D volume or 2D surface.

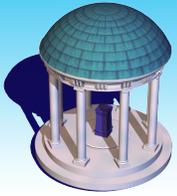
$N_D$ : the number of points on  $D$

$d$ : a voxel in  $D$

$\{Y_{i,D} : i = 1, \dots, n\}$  : data on  $D$  for the  $i^{\text{th}}$  subject

$X_i$  : a  $k \times 1$  covariates

$\theta(d)$  : unknown parameter



# Multiscale Adaptive Regression Model

## Voxel-wise Approach

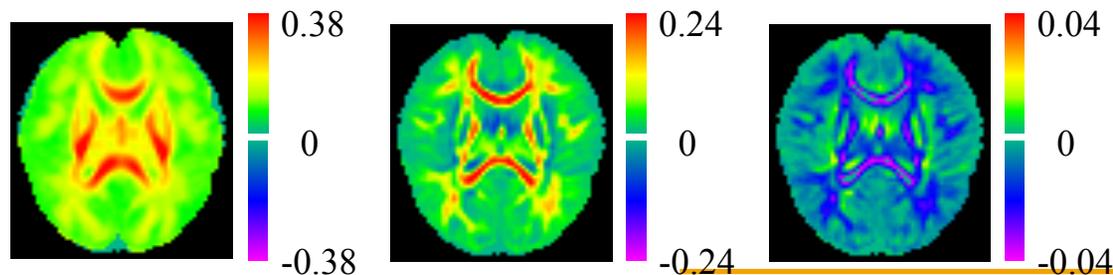
$$p(\mathbf{Y}_{i,\mathcal{D}}|\mathbf{X}_i) = \prod_{d \in \mathcal{D}} p(Y_i(d)|\mathbf{x}_i, \boldsymbol{\theta}(d)),$$

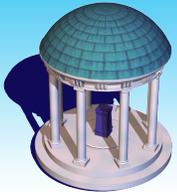
**MARM**

**Being Spatial**

$$p(\mathbf{Y}_{i,\mathcal{D}}|\mathbf{X}_i) \approx \prod_{D_k} p(\{Y_i(d') : d' \in D_k\}|\mathbf{x}_i)$$

$D_k$  denotes the set of all voxels in a homogeneous region

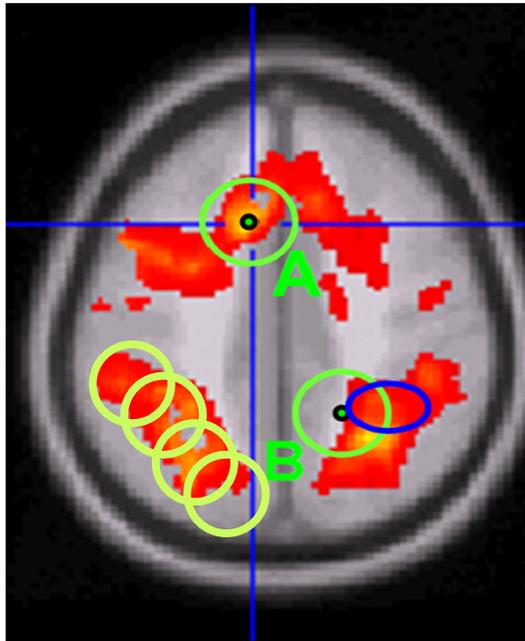




# Multiscale Adaptive Regression Model

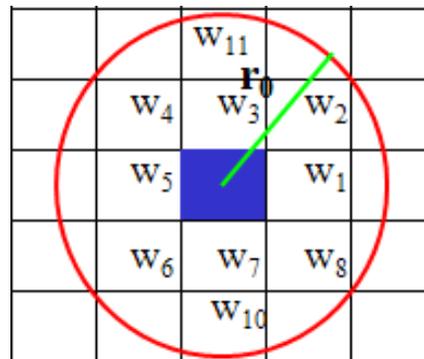
Identifying homogeneous regions

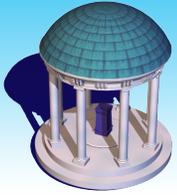
$$D_k$$



Drawing a sphere with radius  $r_0$  at each voxel

Calculating the similarities between the current voxel and its neighboring voxels.





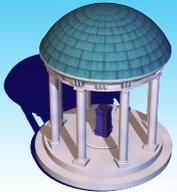
# Multiscale Adaptive Regression Model

## Model Specification

$$p(Y_i | x_i, \theta) = \int_{d \in \mathcal{D}} p(Y_i | x_i, \theta(d)) \omega(d, d'; r_0) \theta(d)$$

$$\ell(\{Y_i(d') : d' \in B(d, r_0)\} | x_i) = \sum_{d' \in B(d, r_0)} \omega(d, d'; r_0) \ell(Y_i(d') | x_i, \theta(d'))$$

$\omega(d, d'; r_0)$  is a **weight** function for characterizing the **similarity** between the data in voxels  $d$  and  $d'$ .



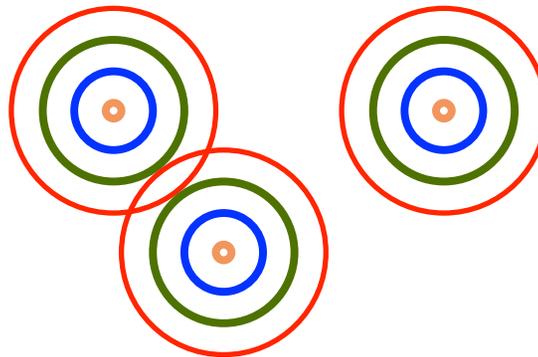
# Multiscale Adaptive Regression Model

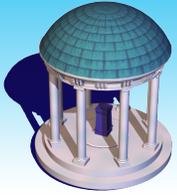
Being Hierarchical

Drawing nested spheres with increasing radiuses at each voxel



$$h_0 = 0 < h_1 < \dots < h_S = r_0$$

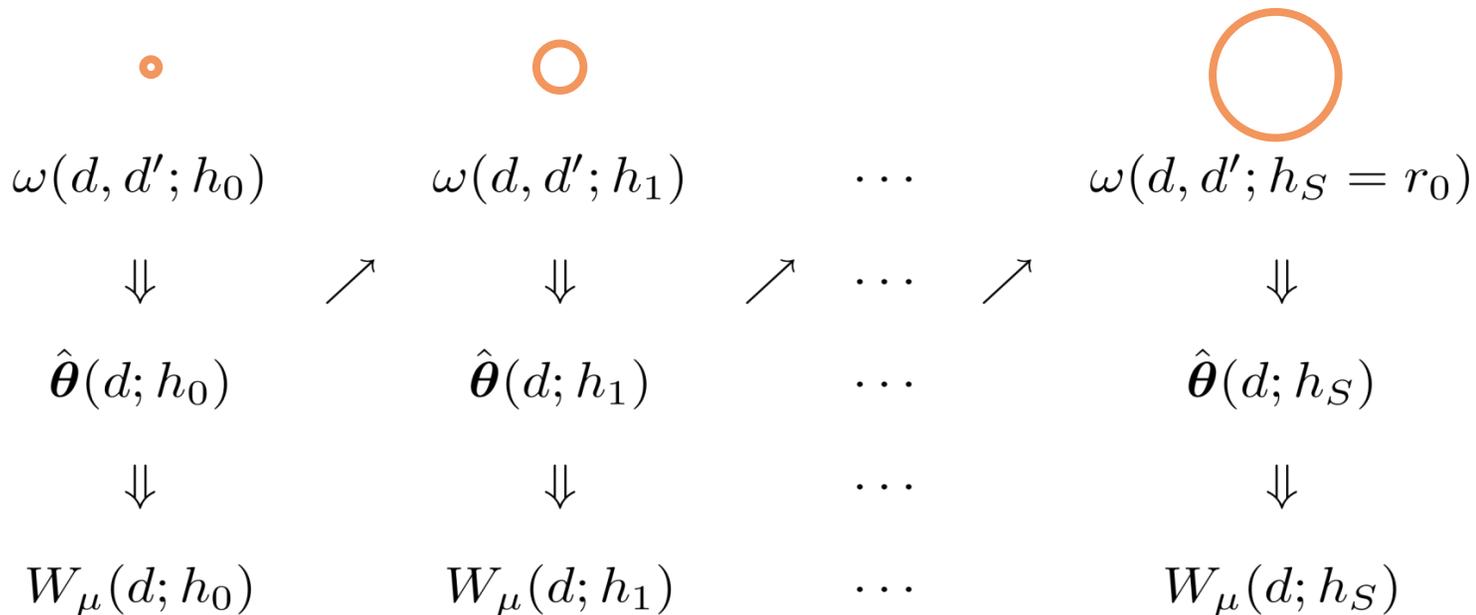


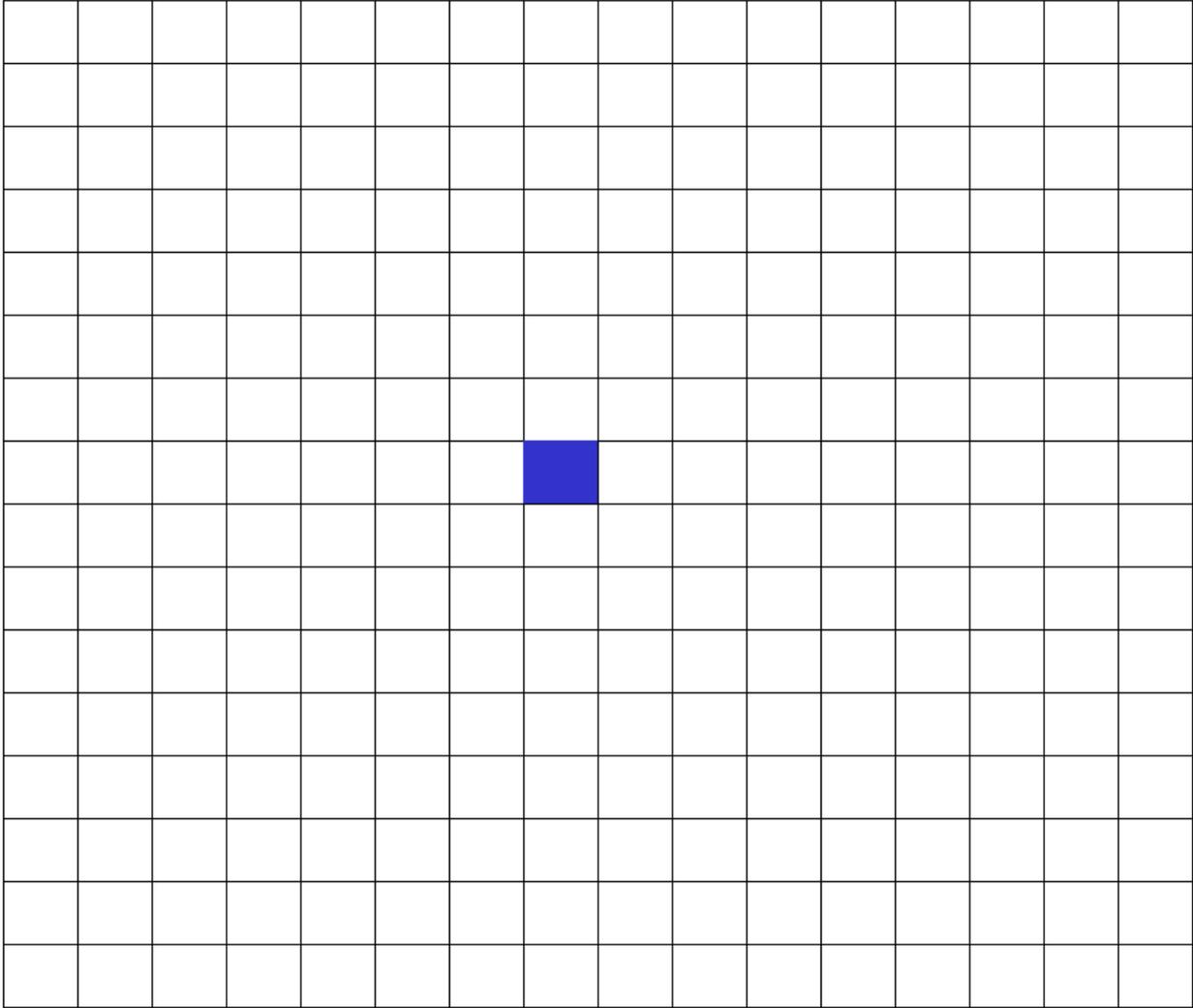


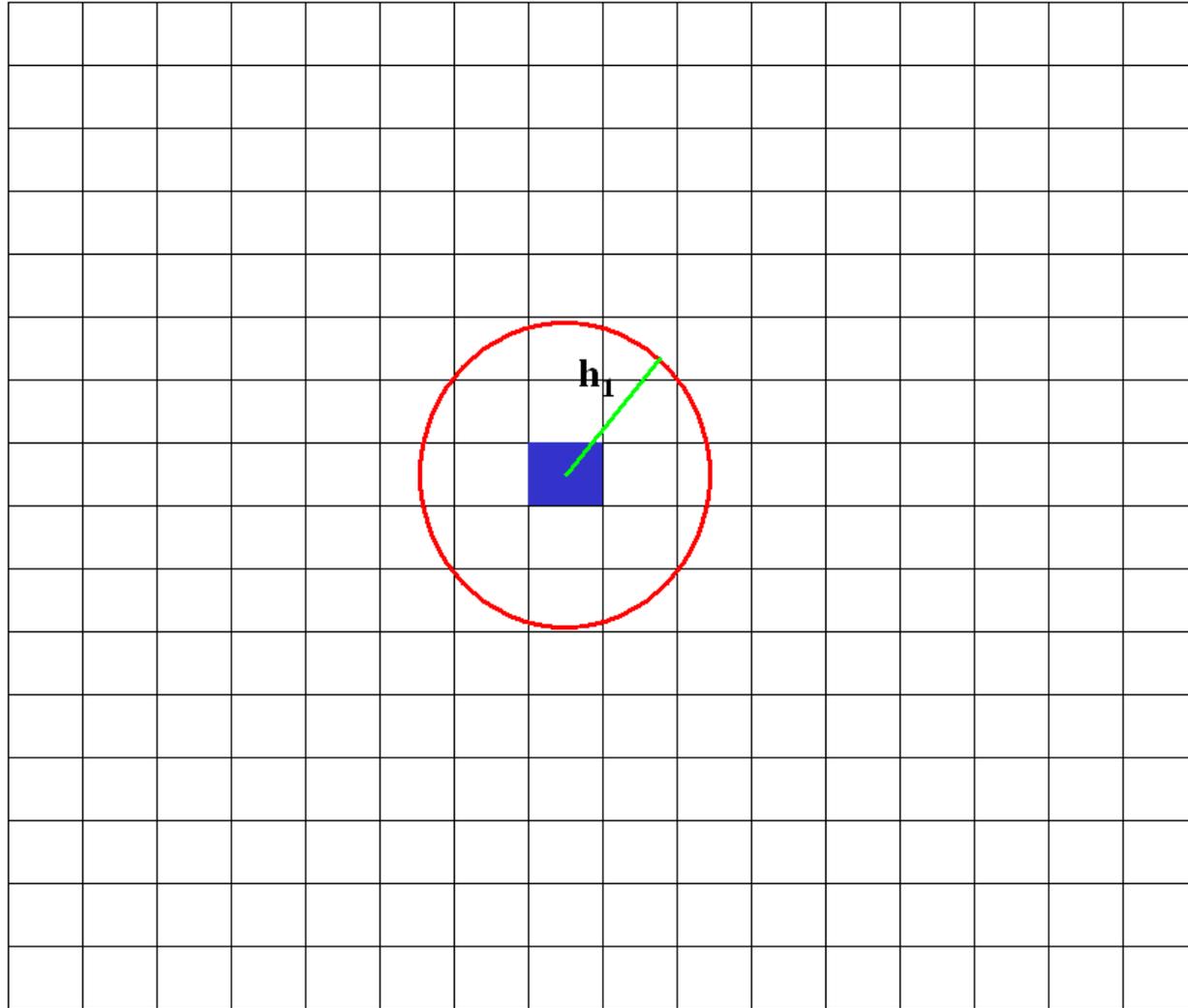
# Multiscale Adaptive Regression Model

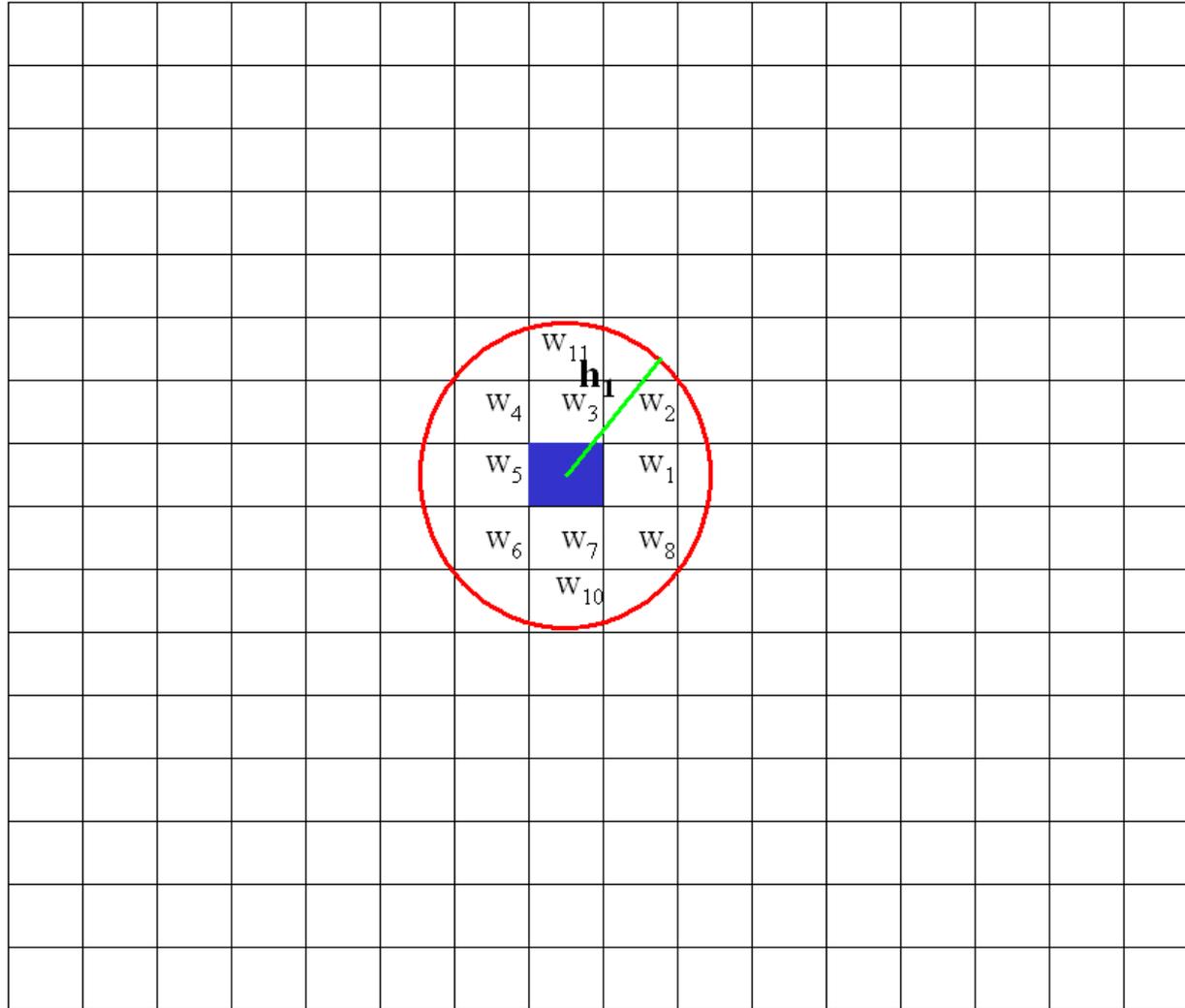
## Being Adaptive

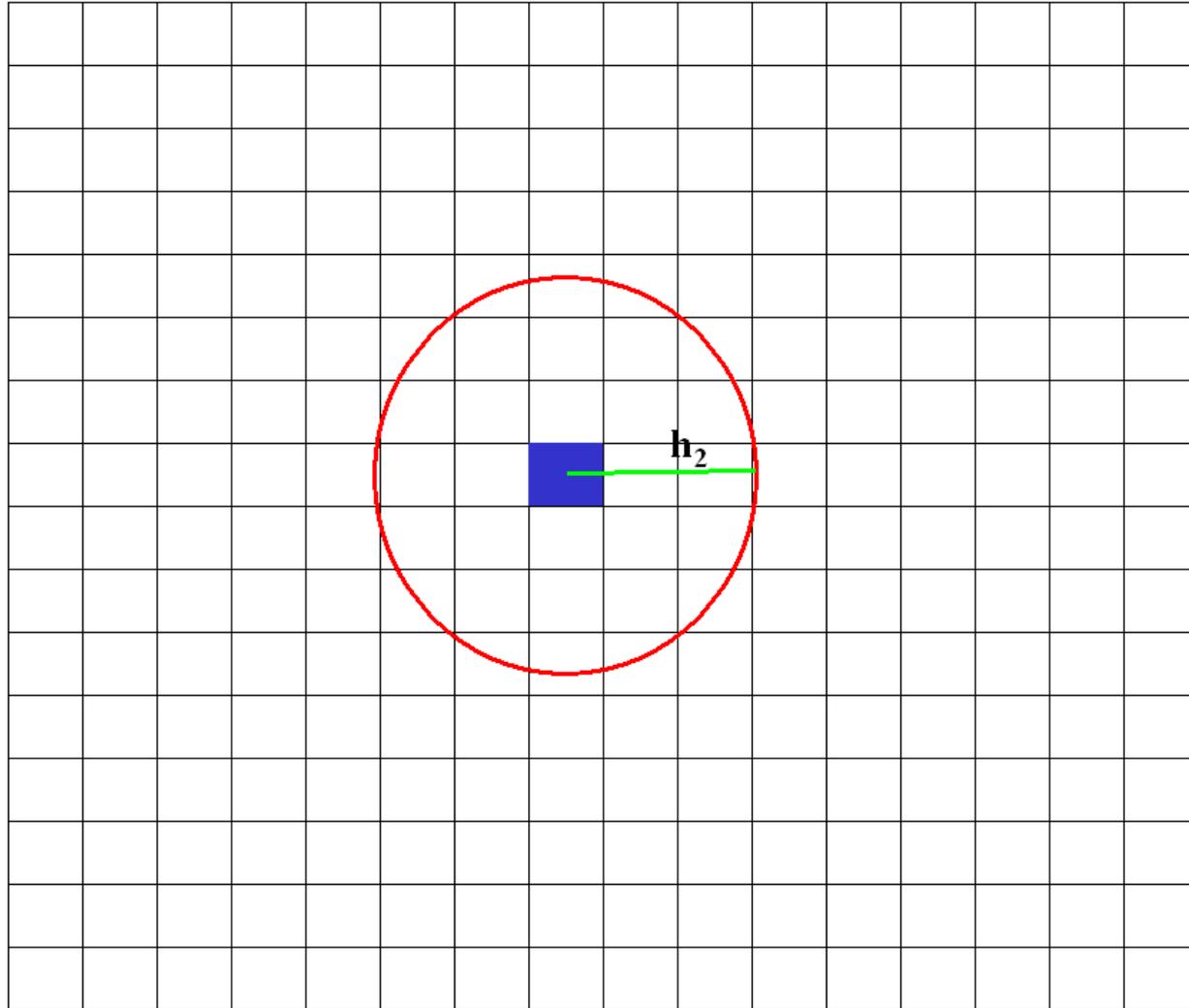
Sequentially determine  $\omega(d, d'; h)$  and adaptively update  $\hat{\theta}(d, h)$



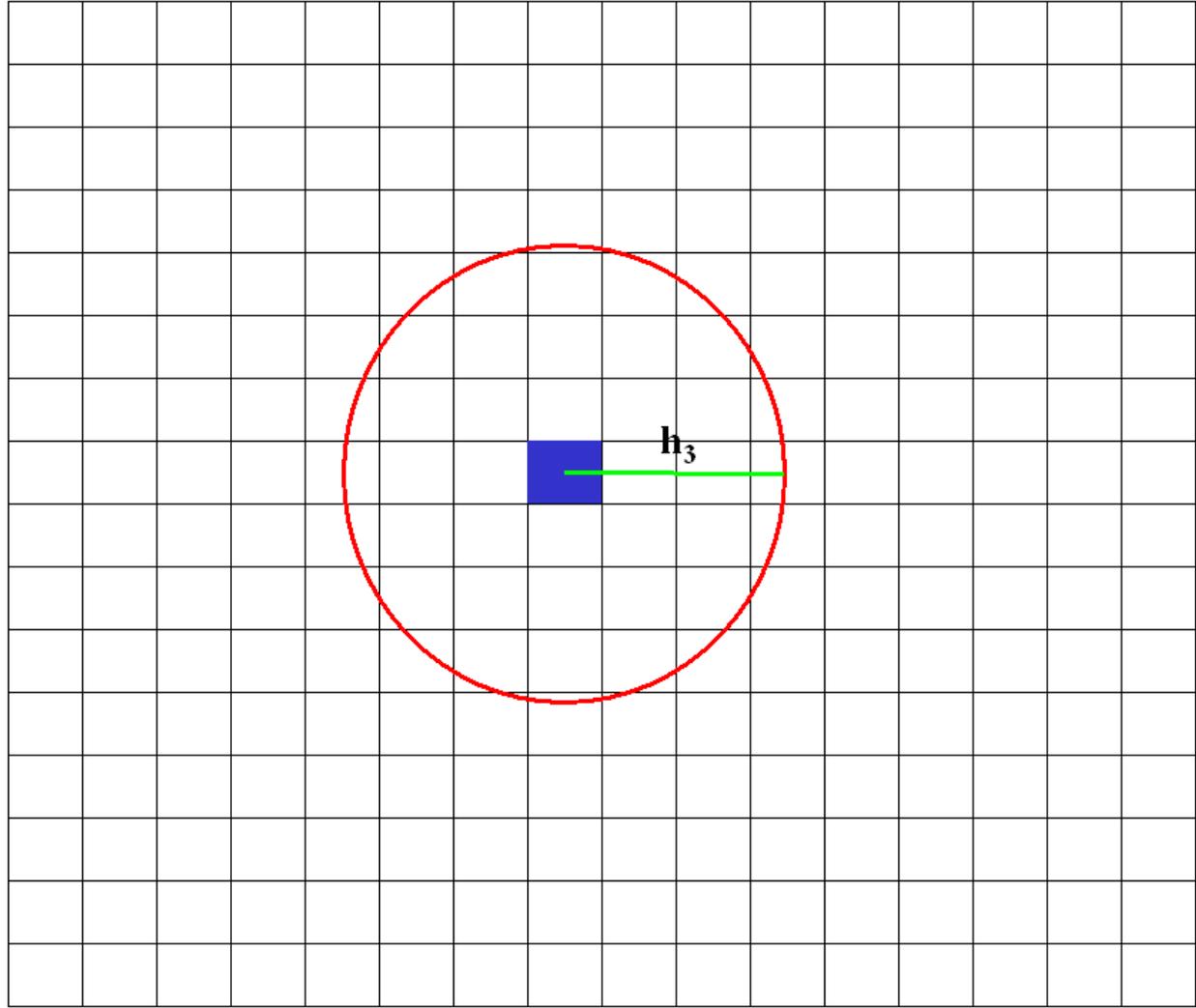


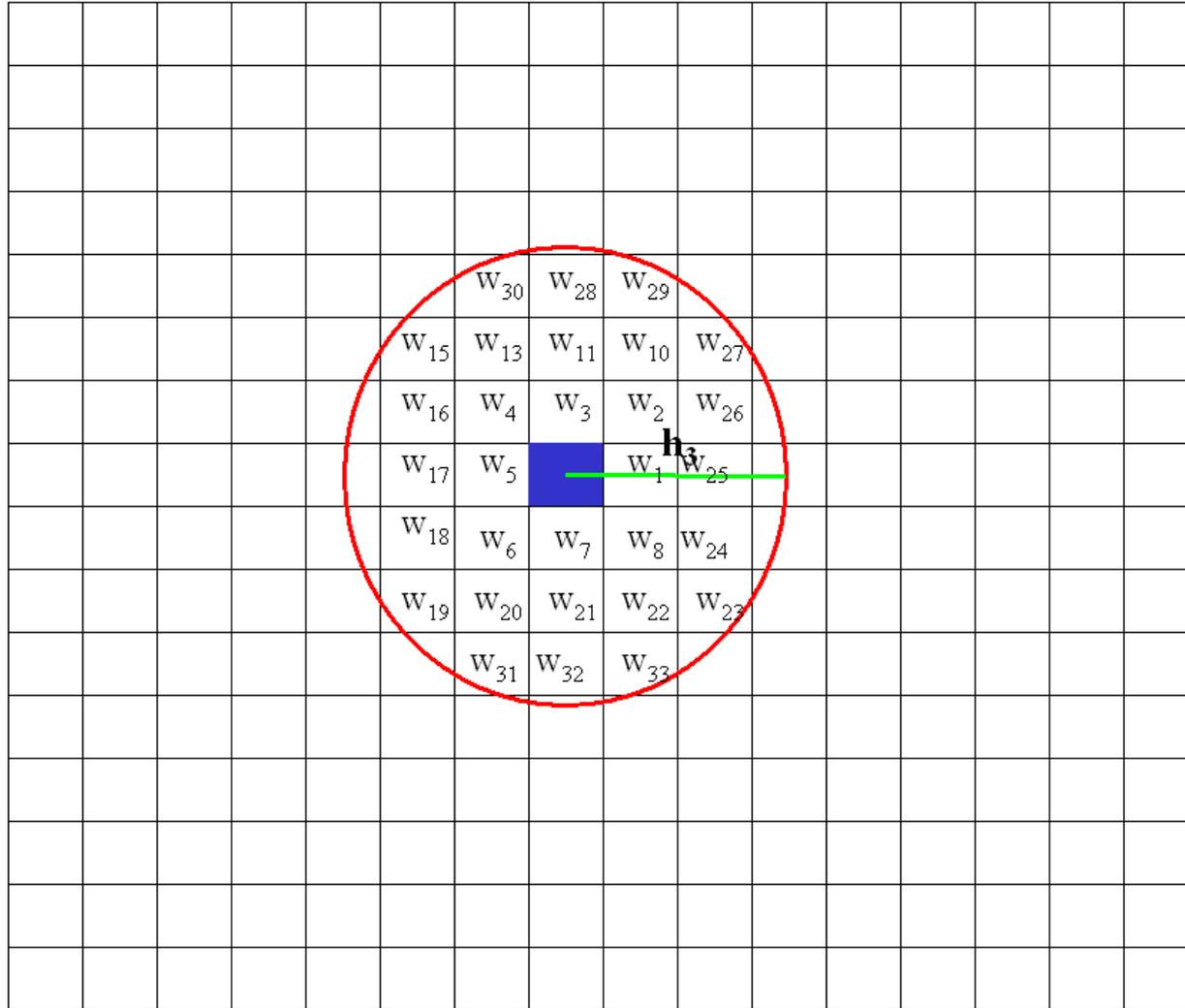


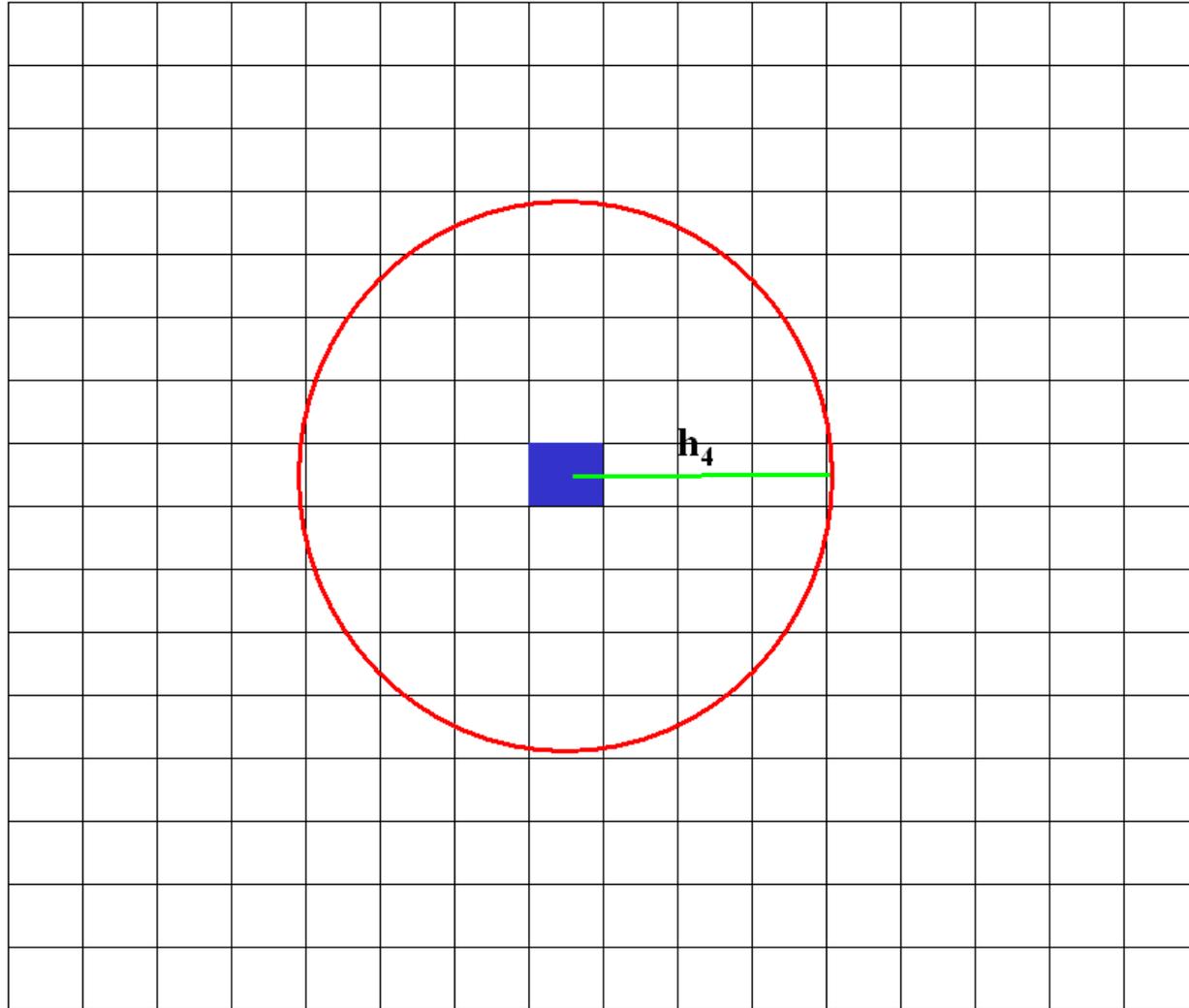


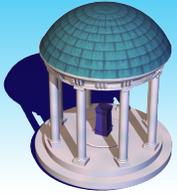












# Multiscale Adaptive Regression Model

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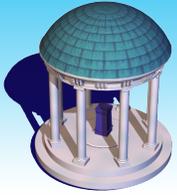
**MARM/PS**

**Learning Voxel Feature**

**Local Feature Adaptation**

**Adaptive Estimation and Testing**

**Automatic Stop**



# Multiscale Adaptive Regression Model

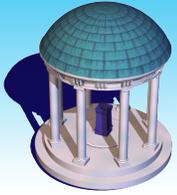
## Local Feature Adaptation

- For any radius  $h_s > h_0$ , define

$$\omega(d, d'; h_s) = K_{loc}(\|d - d'\|_2 / h_s) K_{st}(D_{\theta}(d, d'; h_{s-1}) / C_n)$$

- $K_{loc}(u)$  and  $K_{st}(u)$  are two decreasing kernel functions
- Smoothing kernel:  $K_{loc}(u) = (1 - u^2)_+$
- Similarity kernel:  $K_{st}(u) = \exp(-u) \mathbf{1}\left(u \leq \frac{s+2}{s(\log s+2)}\right)$
- Dissimilarity measure:

$$D_{\theta}(d, d'; h_{s-1}) = [\hat{\theta}(d; h_{s-1}) - \hat{\theta}(d'; h_{s-1})]^T \hat{\Sigma}(\hat{\theta}(d; h_{s-1}))^{-1} [\hat{\theta}(d; h_{s-1}) - \hat{\theta}(d'; h_{s-1})].$$



# Multiscale Adaptive Regression Model

## Adaptive Estimation and Testing

### Weighted quasi-likelihood

$$\ell_n(\boldsymbol{\theta}(d); h, \tilde{\omega}) = \sum_{i=1}^n \sum_{d' \in B(d, h)} \tilde{\omega}(d, d'; h) \log p(Y_i(d') | \mathbf{x}_i, \boldsymbol{\theta}(d))$$

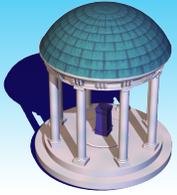
### MWQLE

$$\hat{\boldsymbol{\theta}}(d, h) = \operatorname{argmax}_{\boldsymbol{\theta}(d)} n^{-1} \ell_n(\boldsymbol{\theta}(d); h, \tilde{\omega})$$

### Newton-Raphson Algorithm

$$\hat{\boldsymbol{\theta}}(d, h)^{(t+1)} = \hat{\boldsymbol{\theta}}(d, h)^{(t)} + \{-\partial_{\boldsymbol{\theta}(d)}^2 \ell_n(\hat{\boldsymbol{\theta}}(d, h)^{(t)}; h, \tilde{\omega})\}^{-1} \partial_{\boldsymbol{\theta}(d)} \ell_n(\hat{\boldsymbol{\theta}}(d, h)^{(t)}; h, \tilde{\omega})$$

### Expectation-Maximization Algorithm



# Multiscale Adaptive Regression Model

## Adaptive Estimation and Testing

### Sandwich Estimator

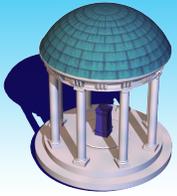
$$\text{Cov}[\hat{\boldsymbol{\theta}}(d, h)] \approx \Sigma_n(\hat{\boldsymbol{\theta}}(d, h)) = [\Sigma_{n,1}(\hat{\boldsymbol{\theta}}(d, h))]^{-1} \Sigma_{n,2}(\hat{\boldsymbol{\theta}}(d, h)) [\Sigma_{n,1}(\hat{\boldsymbol{\theta}}(d, h))]^{-1}$$

$$\Sigma_{n,1}(\boldsymbol{\theta}(d)) = -\partial_{\boldsymbol{\theta}(d)}^2 \ell_n(\boldsymbol{\theta}(d); h, \tilde{\boldsymbol{\omega}}) \text{ and}$$

$$\Sigma_{n,2}(\boldsymbol{\theta}(d)) = \sum_{i=1}^n \left[ \sum_{d' \in B(d, h)} \tilde{\omega}(d, d'; h) \partial_{\boldsymbol{\theta}(d)} \log p(Y_i(d') | \mathbf{x}_i, \boldsymbol{\theta}(d)) \right]^{\otimes 2}$$

### Wald Test Statistic

$$[R(\hat{\boldsymbol{\theta}}(d; h)) - \mathbf{b}_0]^T [\partial_{\boldsymbol{\theta}(d)} R(\hat{\boldsymbol{\theta}}(d; h)) \hat{\Sigma}_n(\hat{\boldsymbol{\theta}}(d; h)) \partial_{\boldsymbol{\theta}(d)} R(\hat{\boldsymbol{\theta}}(d; h))^T]^{-1} [R(\hat{\boldsymbol{\theta}}(d; h)) - \mathbf{b}_0]$$



# Multiscale Adaptive Regression Model

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$\log(\text{Voxel size}) \ll Cn \ll \text{sample size}$

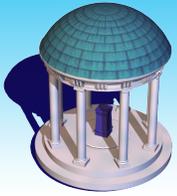
Kernel functions

Conditions for M-estimators hold uniformly

Weak Consistency

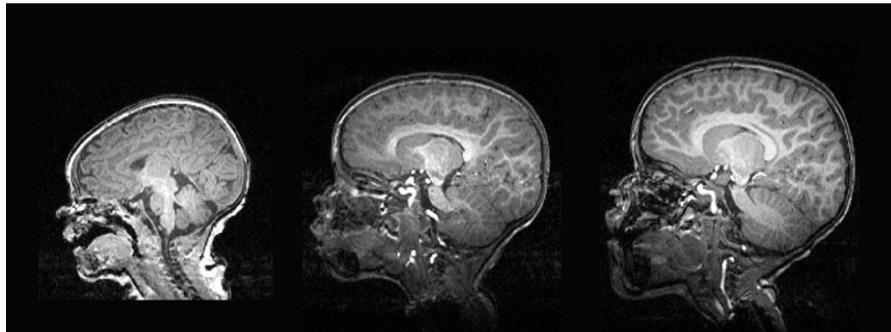
Asymptotical Normality

Asymptotically Chi-squared distribution



## Real Data

- **Early Brain Development Project**
- **Objective:** We want to assess the brain structural connectivity change in the early brain development.
- **Subject:** 250 infants.
- **PI:** John Gilmore
- **MRIs:** DWI, resting fMRI, and T1 MRI were acquired for each subject at 2 weeks, 1, 2, 3, 4 years old.

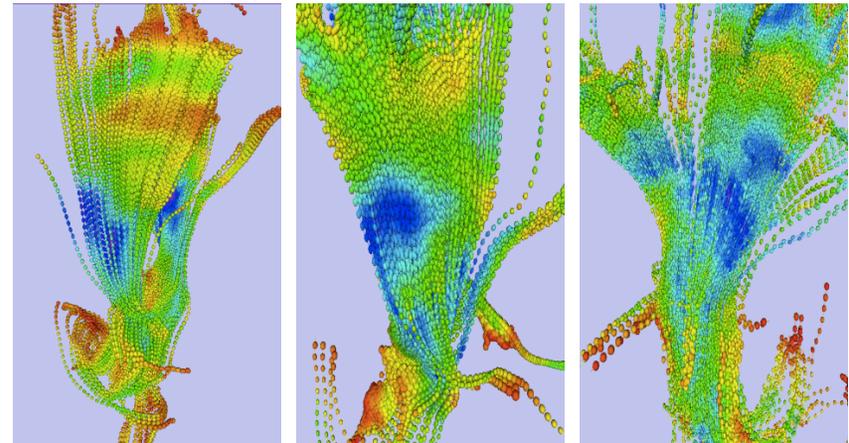


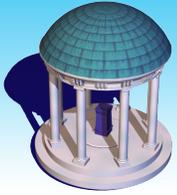
2 weeks

1 year

2 year

Knickmeyer RC, et al. (2008) *J Neurosci* 28: 12176-12182.

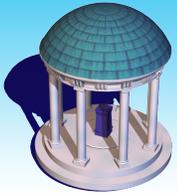




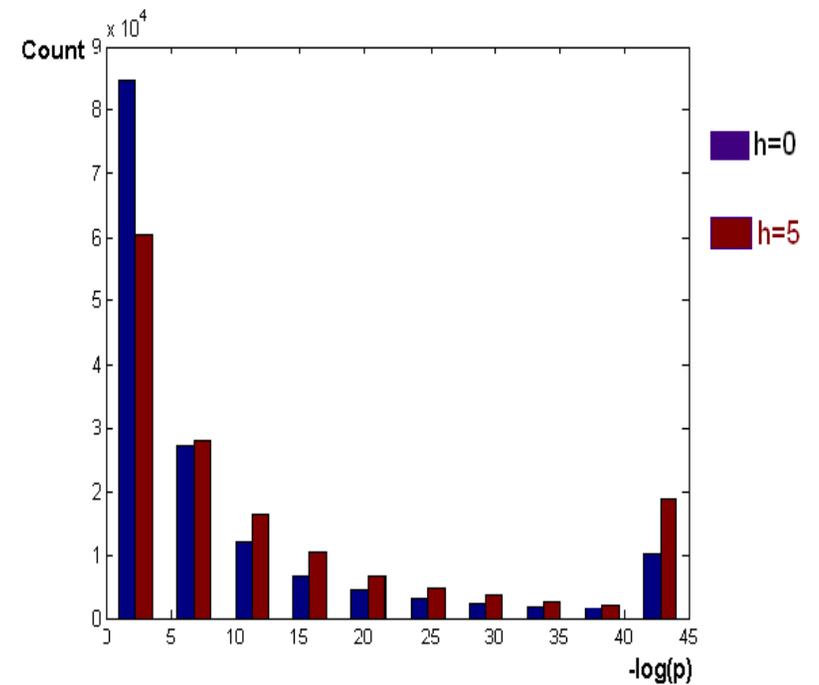
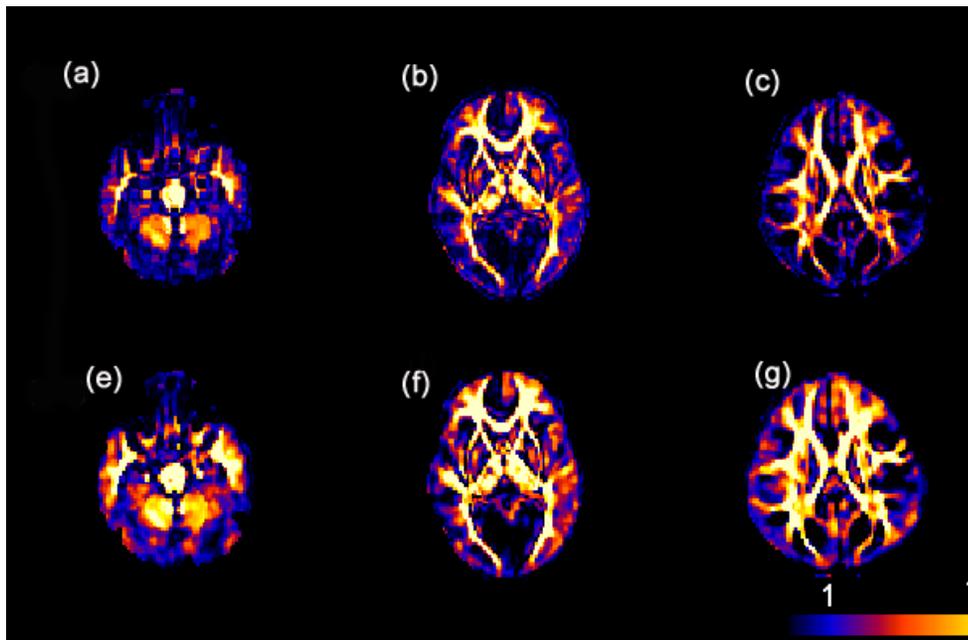
# Infant Brain Development Data

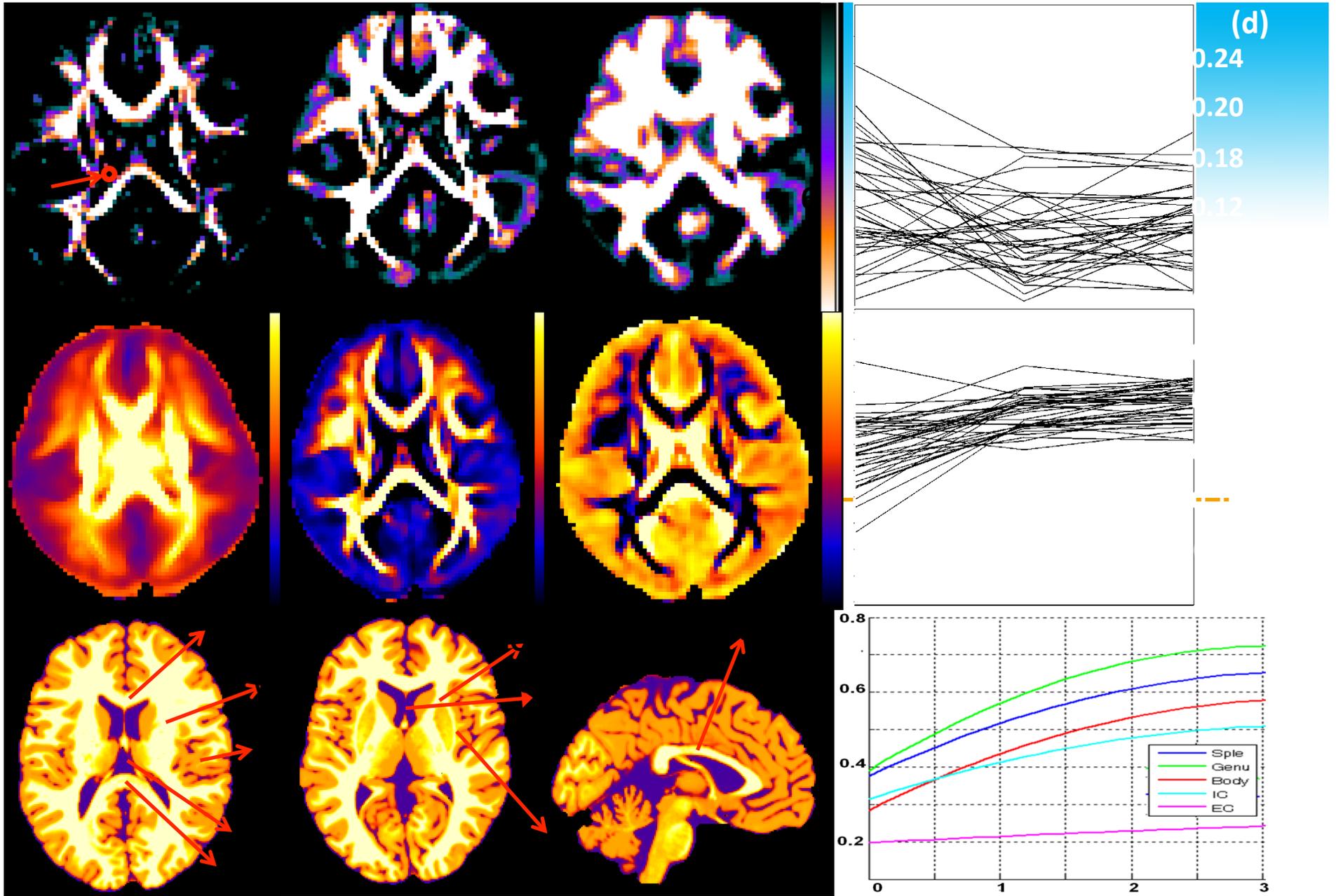
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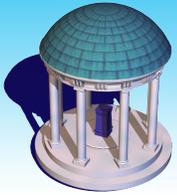
- **Objective:** We want to assess the brain structure change in the early brain development.
- **Subject:** 38 infants.
- **Image:** Diffusion-weighted images and T1 weighted images were acquired for each subject at 2 weeks, 1 and 2 years old.
- **Method:** Voxel-wise imaging analysis and MARM.



# Time Effect and Comparison

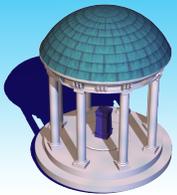




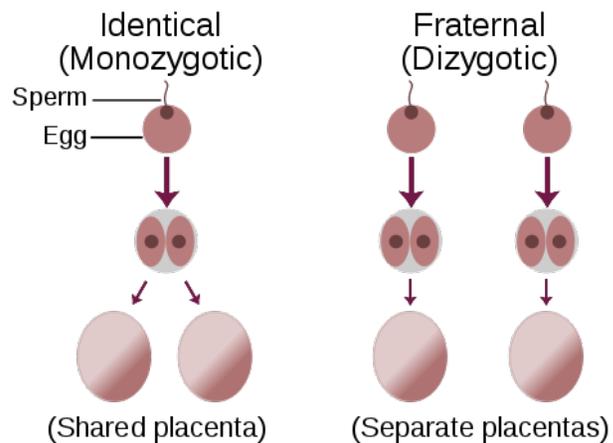
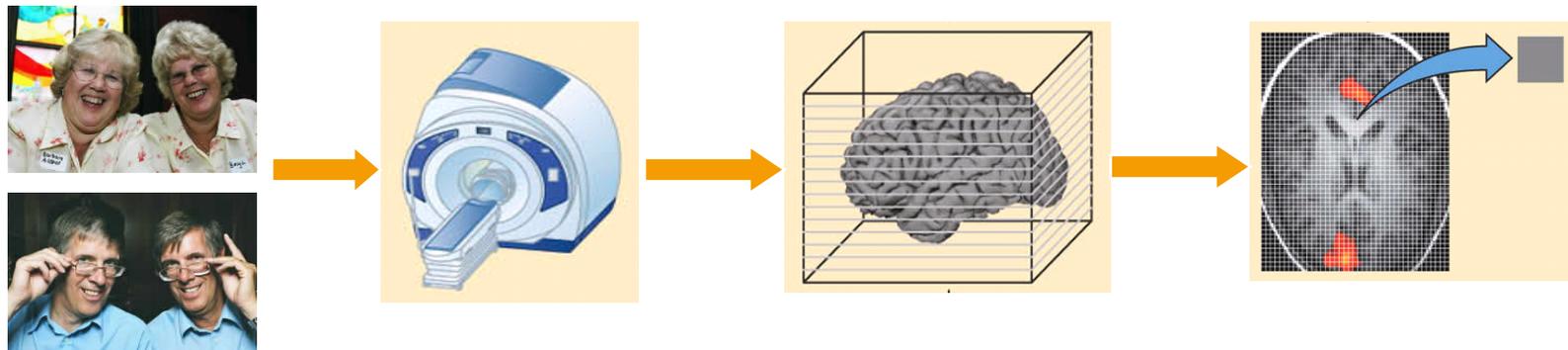


## Twin-MARM

- **ACE/ADE Models**
- **Two-stage MARM**
- **Consistency and Asymptotic Distribution**

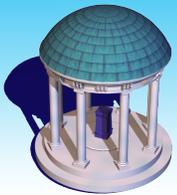


# Twin-MARM



MZ twins share same genetic material

DZ twins share average 50% of their genes

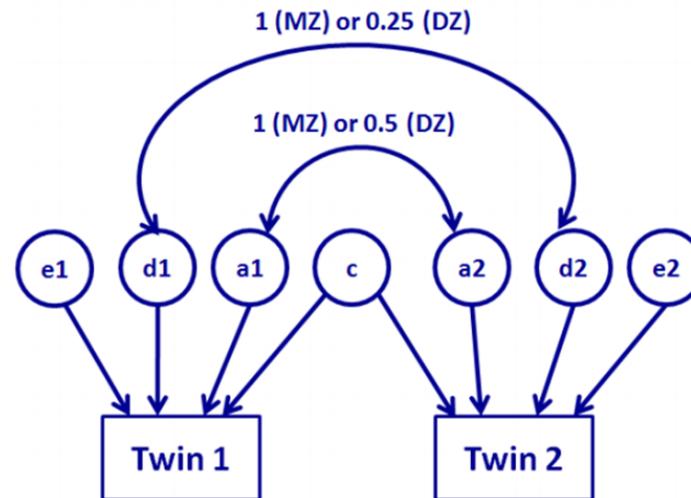


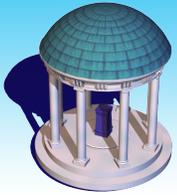
# Twin-MARM

At specific voxel  $v$ , we consider the structural equation model:

$$y_{ij}(v) = x_{ij}^T \beta(v) + a_{ij}(v) + d_{ij}(v) + c_i(v) + e_{ij}(v)$$

$a_{ij}(v)$ ,  $d_{ij}(v)$ ,  $c_i(v)$  and  $e_{ij}(v)$  : the additive genetic, dominance genetic, common environmental and residual effects on  $i$ -th twin pair. We assume they are independently normally distributed with mean 0 and variance  $\sigma_a(v)^2$ ,  $\sigma_d(v)^2$ ,  $\sigma_c(v)^2$  and  $\sigma_e(v)^2$ .





## Twin-MARM

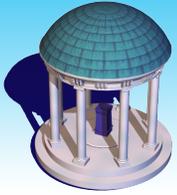
There are two sets of parameters:  
mean structure  
**variance structure**

$$\omega(d, d'; h_s) = K_{loc}(\|d - d'\|_2 / h_s) K_{st}(D_{\theta}(d, d'; h_{s-1}) / C_n)$$

$$D_{\theta}(d, d'; h_{s-1}) = [\hat{\theta}(d; h_{s-1}) - \hat{\theta}(d'; h_{s-1})]^T \hat{\Sigma}(\hat{\theta}(d; h_{s-1}))^{-1} [\hat{\theta}(d; h_{s-1}) - \hat{\theta}(d'; h_{s-1})].$$

**Question of interest:**

**Mean and variance images may have different patterns.**



# Twin-MARM

## Two-stage Approach

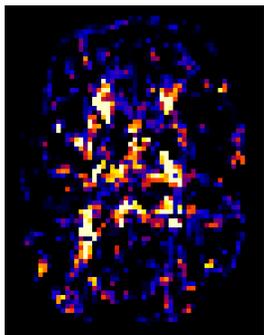
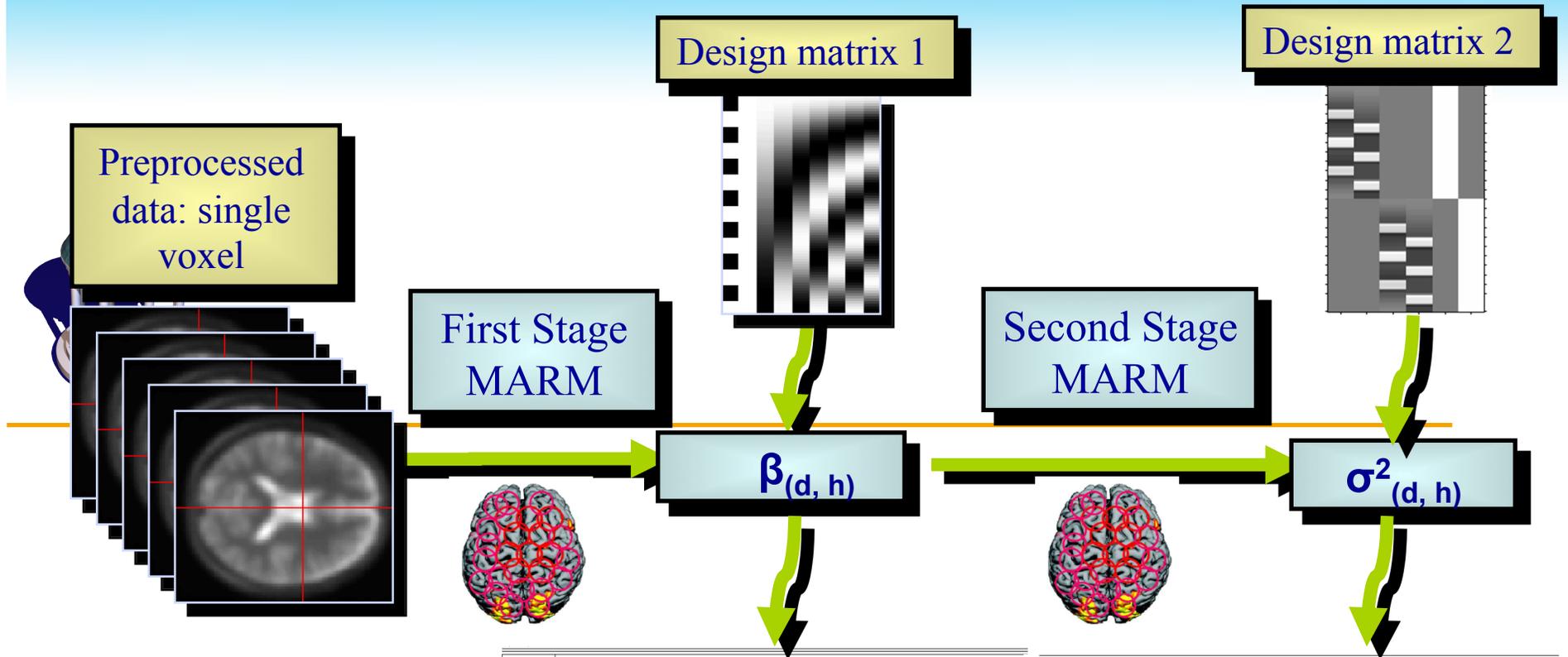
- **Mean structure**

$$Y_{ij}(d) = x_{ij}^T \beta(d) + \varepsilon_{ij}(d) \Rightarrow \{\hat{\beta}(d; h) : d \in D\}$$

- **Variance structure**

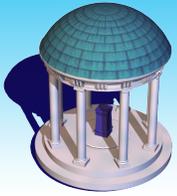
$$\{Y_{ij}(d) - x_{ij}^T \hat{\beta}(d; h)\}^2 = z_{ij}^T \rho(d) + \delta_{ij}(d) \Rightarrow \{\hat{\rho}(d; h) : d \in D\}$$

# TwinMARM



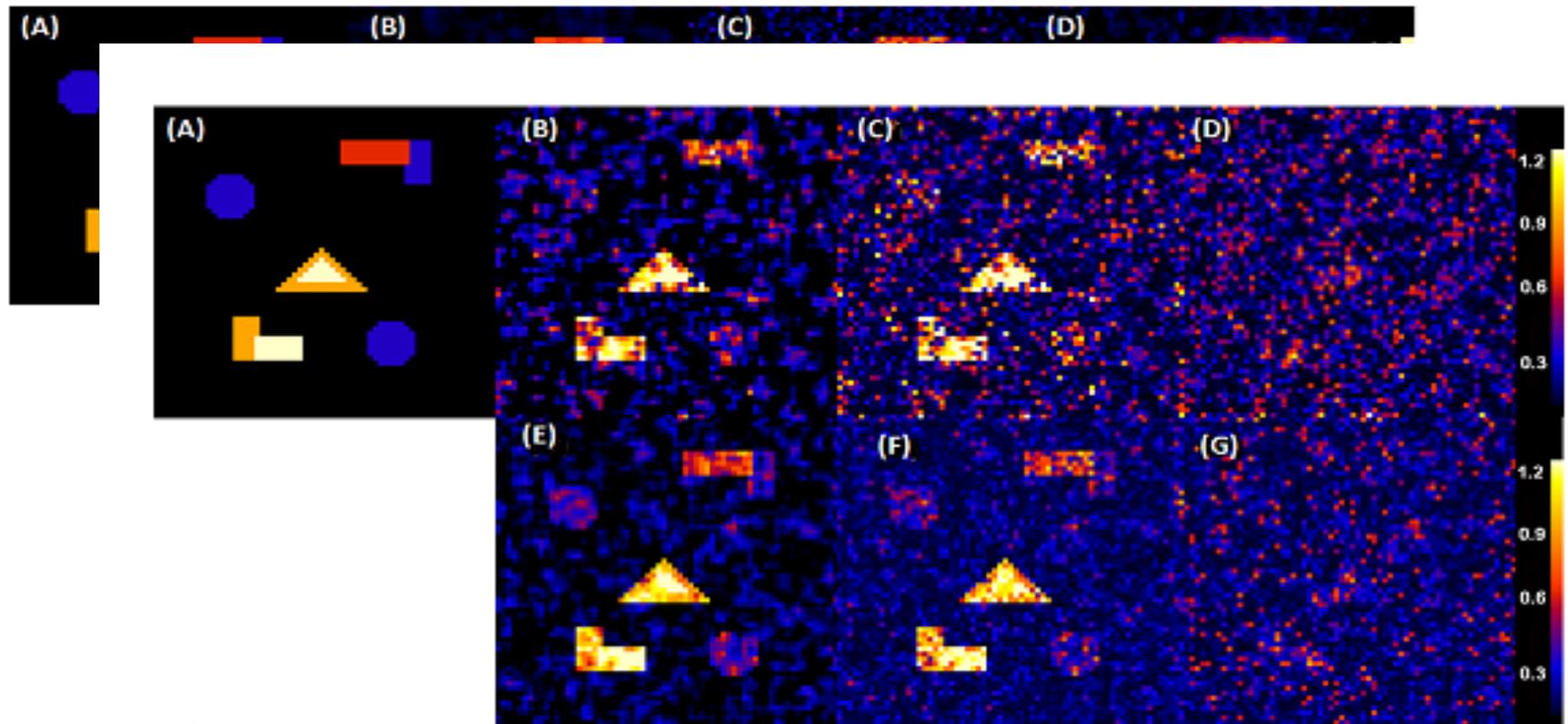
$\beta_1(d)$		$n = 100$			$n = 200$			$n = 400$		
		$h_0$	$h_5$	$h_{10}$	$h_0$	$h_5$	$h_{10}$	$h_0$	$h_5$	$h_{10}$
0	BIAS	0.00	0.00	0.00	0.00	0.00	0.00	0.00	0.00	0.00
	RMS	0.21	0.12	0.08	0.15	0.09	0.05	0.11	0.06	0.04
	SD	0.21	0.12	0.07	0.15	0.08	0.05	0.11	0.06	0.04
	RE	1.02	1.06	1.06	1.01	1.03	1.03	1.00	1.01	1.01
0.5	BIAS	0.00	-0.01	0.00	0.00	0.00	0.00	0.00	0.00	0.00
	RMS	0.22	0.14	0.09	0.16	0.10	0.06	0.12	0.07	0.04
	SD	0.22	0.13	0.08	0.16	0.09	0.06	0.12	0.07	0.04
	RE	1.01	1.09	1.11	1.01	1.04	1.03	1.00	1.00	0.98

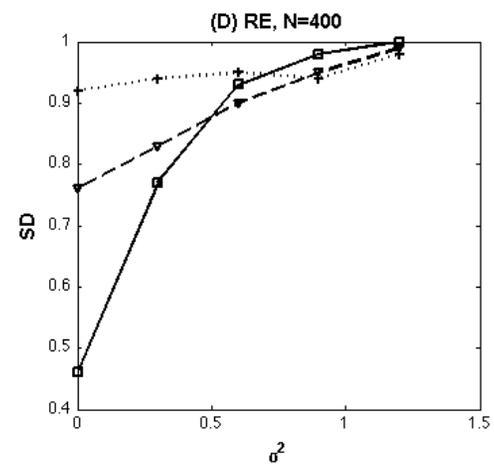
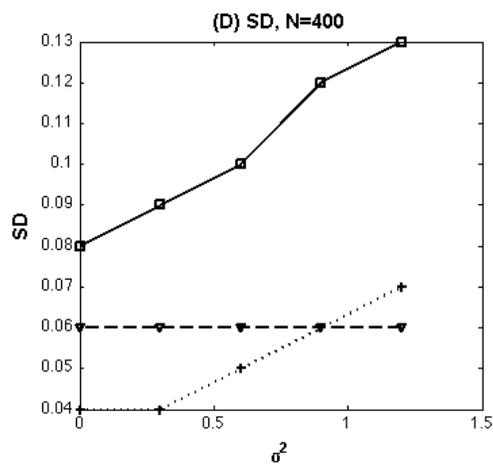
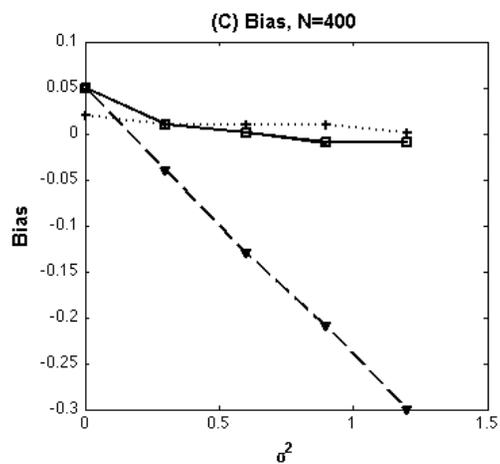
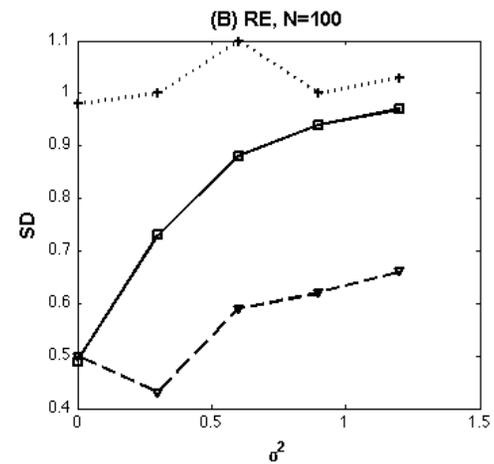
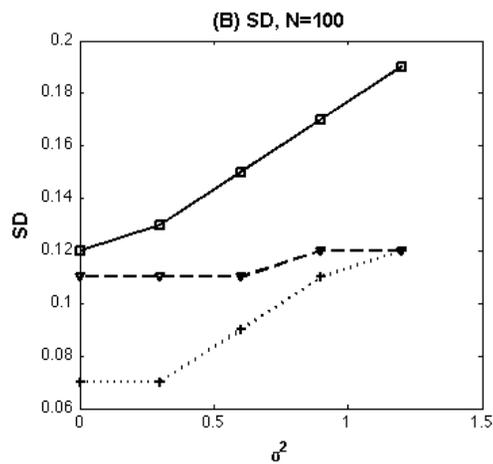
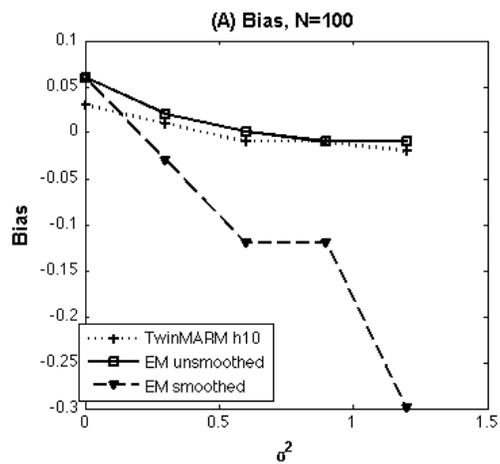
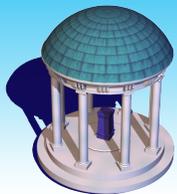
$\sigma_a^2(d)$		$n = 100$			$n = 200$			$n = 400$		
		$h_0$	$h_5$	$h_{10}$	$h_0$	$h_5$	$h_{10}$	$h_0$	$h_5$	$h_{10}$
0	BIAS	0.03	0.03	0.03	0.02	0.02	0.02	0.02	0.02	0.02
	RMS	0.13	0.08	0.07	0.08	0.06	0.05	0.06	0.04	0.04
	SD	0.13	0.08	0.07	0.09	0.06	0.05	0.06	0.04	0.04
	RE	0.99	0.99	0.98	0.99	0.98	0.98	0.96	0.93	0.92
0.3	BIAS	0.03	0.01	0.01	0.02	0.00	0.00	0.02	0.02	0.01
	RMS	0.14	0.09	0.07	0.09	0.06	0.05	0.07	0.04	0.04
	SD	0.14	0.09	0.07	0.10	0.06	0.05	0.07	0.05	0.04
	RE	1.00	1.00	1.00	0.99	0.97	0.97	0.97	0.95	0.94

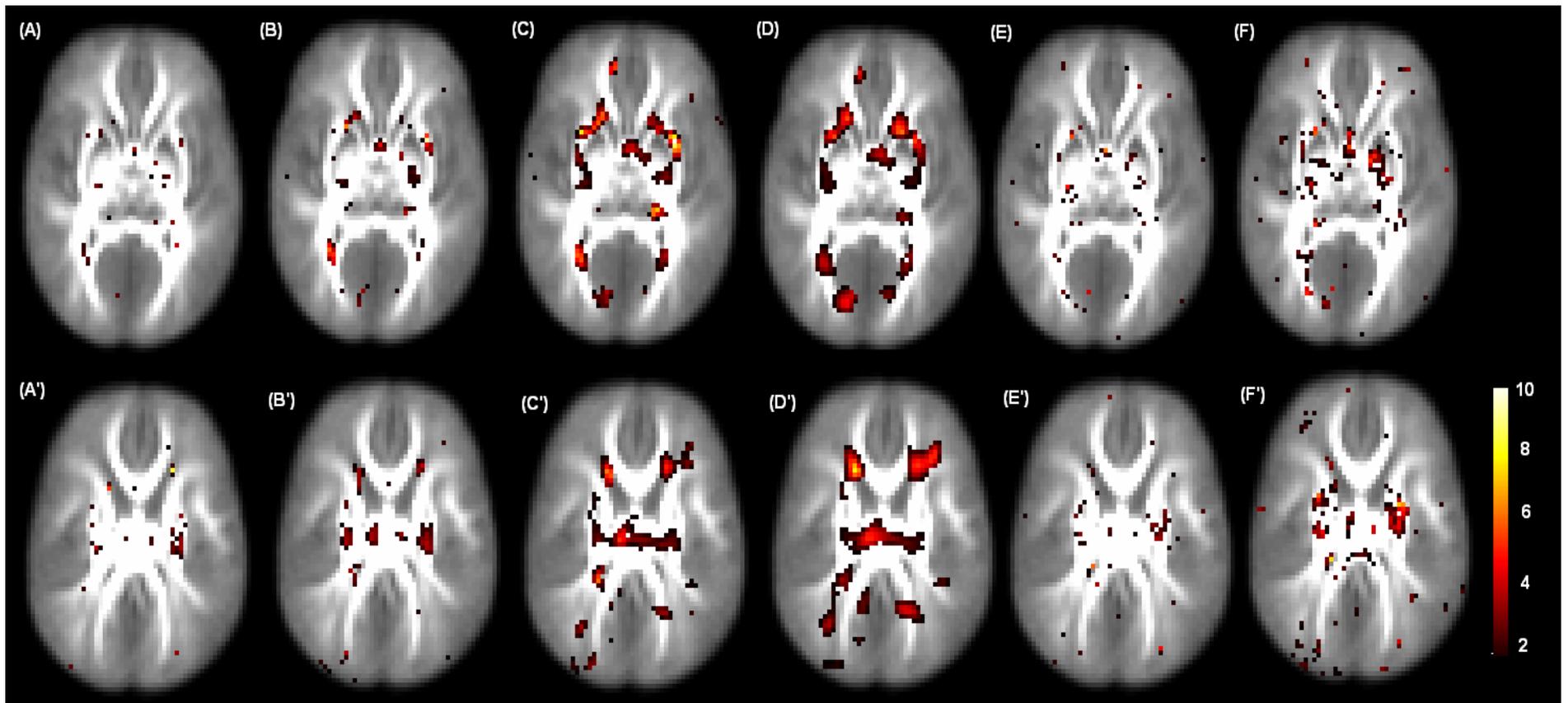
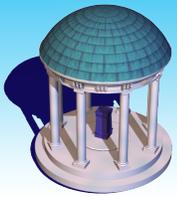


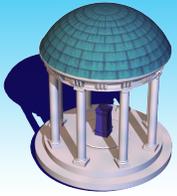
## Twin-MARM

It is dangerous to use Gaussian-kernel to smooth imaging data and then carry out twin analysis.



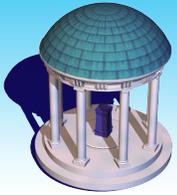






## Multiscale Adaptive Smoothing Models for HRF in fMRI

- Convolution Models in Frequency Domain
- Back Fitting Methods
- Multi-stage MARM



# Multiscale Adaptive Smoothing Model

**$D$** : 3D volume

**$N_D$** : the number of points on  $D$

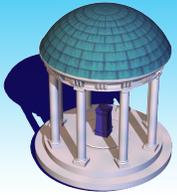
**$d$** : a voxel in  $D$

$\{Y(t, d) : t = 1 \times t_{TR}, \dots, T \times t_{TR}, d \in D\}$ : spatial-temporal process

$\{X(t) : t \in [0, T \times t_{TR}]\}$ : external stimulus process

$\{H(t, d) : t \in [0, T \times t_{TR}], d \in D\}$ : spatial-temporal HRF process

$\{\varepsilon(t, d) : t \in [0, T \times t_{TR}], d \in D\}$ : error process



## Voxel-wise Approach

$$Y(t, d) = H(\bullet, d) \otimes X(t) + \varepsilon(t, d) = \int H(t - u, d) X(u) du + \varepsilon(t, d)$$

**time-domain**

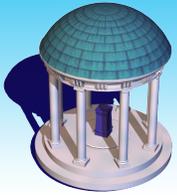
$$H(t, d) = \sum_{k=1}^K \beta_k(d) f_k(t)$$

$$Y(t, d) = \sum_{k=1}^K \beta_k(d) \int f_k(t - u, d) X(u) du + \varepsilon(t, d) = \sum_{k=1}^K \beta_k(d) x_k(t) + \varepsilon(t, d)$$

**frequency-domain**

$$F_Y(f, d) = F_H(f, d) F_X(f) + F_\varepsilon(f, d) \text{ for } f \in [0, T \times t_{TR}]$$

$$F_Y(f, d) = \int_0^{T \times t_{TR}} Y(t, d) \exp(-2\pi i f t / (T \times t_{TR})) dt$$



## Continuous

$$F_Y(f, d) = F_H(f, d)F_X(f) + F_\varepsilon(t, d) \text{ for } f \in [0, T \times t_{TR}]$$

## Discrete

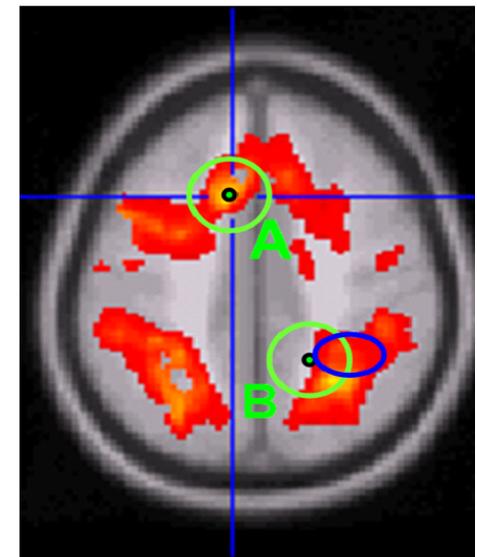
$$\phi_Y(f, d) = \phi_H(f, d)\phi_X(f) + \phi_\varepsilon(f, d) \text{ for } f \in [0, T \times t_{TR}]$$

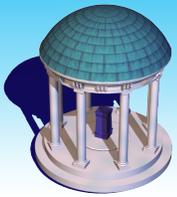
$$\phi_Y(f, d) = \sum_{t=0}^{T \times t_{TR}} Y(t, d) \exp(-2\pi i f t / (T \times t_{TR}))$$

## Key Assumptions:

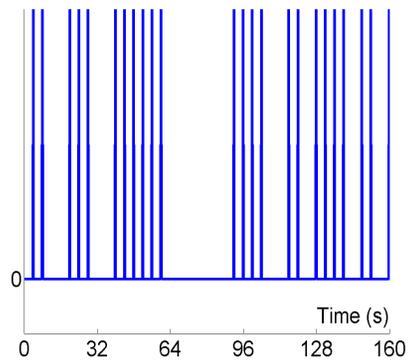
$$\phi_\varepsilon(f, d) \sim (0, 1(f = f')\sigma(f, f'; d, d'))$$

$\phi_H(f, d)$  is piecewisely smooth for  $(f, d) \in \mathcal{N}(f, d)$

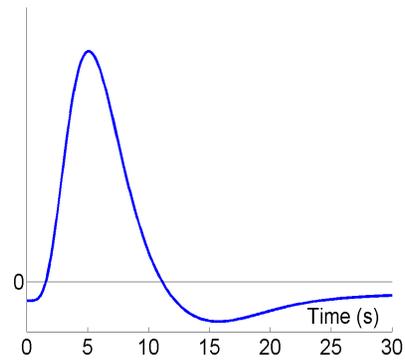




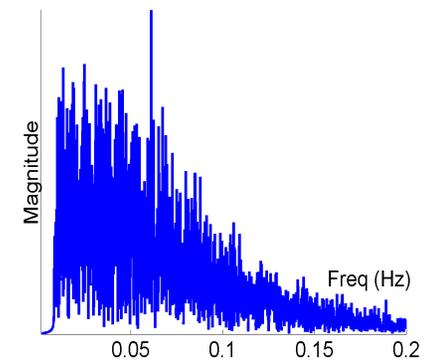
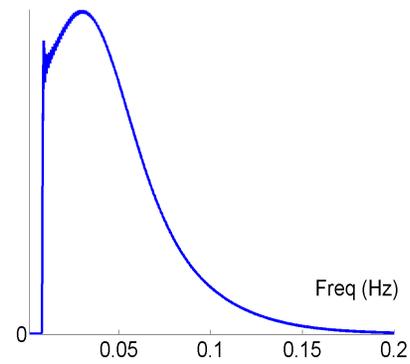
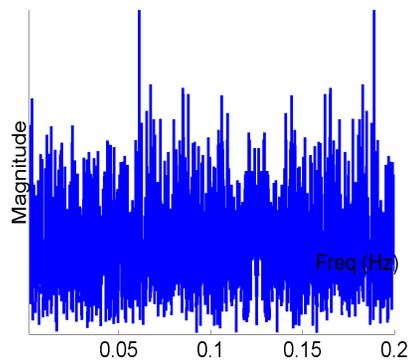
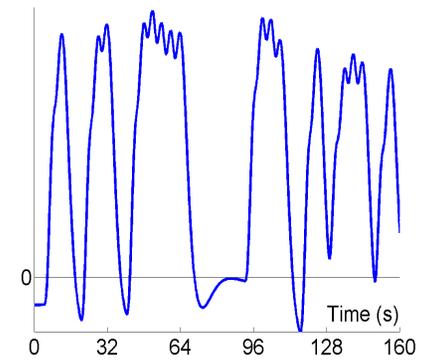
Stimulus ("Neural")

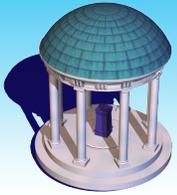


HRF



Predicted Data





$$\phi_Y(f, d) = \phi_H(f, d)\phi_X(f) + \phi_\varepsilon(f, d) \text{ for } (f, d) \in \mathcal{N}(f, d)$$

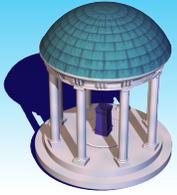
**Approximation**

**Unknown**

$$\begin{aligned}\phi_Y(f_k, d') &= \phi_H(f_k, d')\phi_X(f_k) + \phi_\varepsilon(f_k, d') \\ &\approx \phi_H(f, d)\phi_X(f_k) + \phi_\varepsilon(f_k, d')\end{aligned}$$

$$(f_k, d') \in B((f, d); \varepsilon, r) = (f - \varepsilon, f + \varepsilon) \times B(d', r)$$

**Multiple Events: Backfitting Methods**



## Weighted LSE

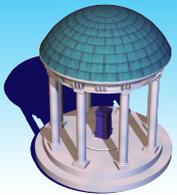
$$L(\phi_H(f, d); B((f, d); r, h)) = \sum_{(f_k, d')} [\phi_Y(f_k, d') - \phi_H(f, d)\phi_X(f_k)]^2 w(f, d, f_k, d'; \varepsilon, r)$$

$$\hat{\phi}_H(f, d) = \sum_{(f_k, d')} \phi_Y(f_k, d') \bar{\phi}_X(f_k) w(f, d, f_k, d'; \varepsilon, r) / \sum_{(f_k, d')} \phi_X(f_k) \bar{\phi}_X(f_k) w(f, d, f_k, d'; \varepsilon, r)$$

$$\text{Var}(\hat{\phi}_H(f, d))$$

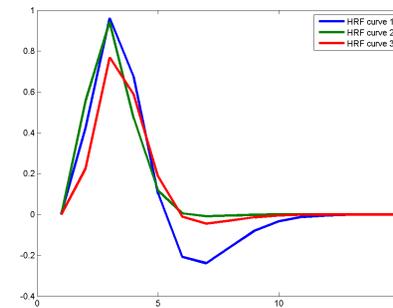
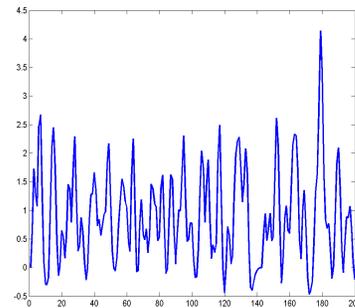
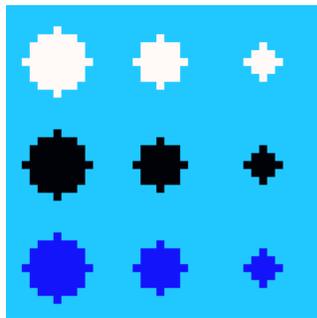
## Estimated HRF

$$\hat{H}(t, d) = \sum_{k=0}^{T-1} \hat{\phi}_H(f, d) \exp(i2\pi t f_k) [1 - \cos(2\pi t / T)] / (2\pi^2 t^2 / T)$$



## Simulation II: Multivariate Case

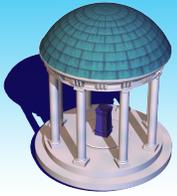
The background image and the simulated one with their related curves.  
In this simulation the smallest SNR is between 0.5 and 0.7.



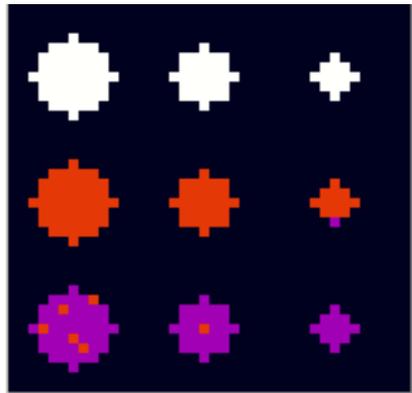
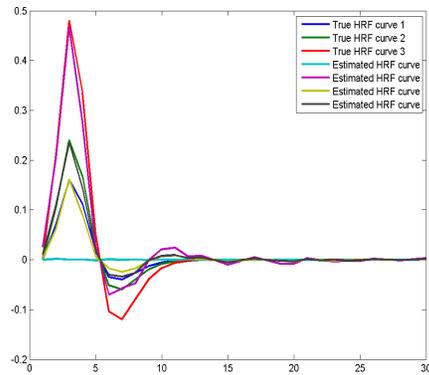
Three activated regions for each sequence of events correspond with three different HRFs:  $h_j(t)/2$ ,  $h_j(t)/4$ ,  $h_j(t)/6$ ,  $j=1,2,3$

$$h_j(t) = A_j \cdot (t/d_{j1})^{a_{j1}} \exp(-(t - d_{j1})/b_{j1}) - c(t/d_{j2})^{a_{j2}} \exp(-(t - d_{j2})/b_{j2}) \quad j = 1, 2, 3$$

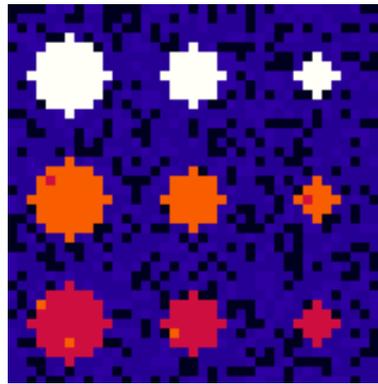
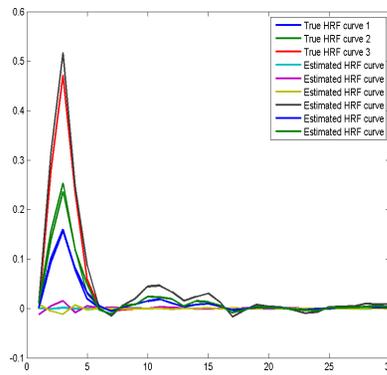
$$\epsilon(t) \sim N(0, \sigma^2) \quad X_j(t) \sim B(1, 0.15)$$



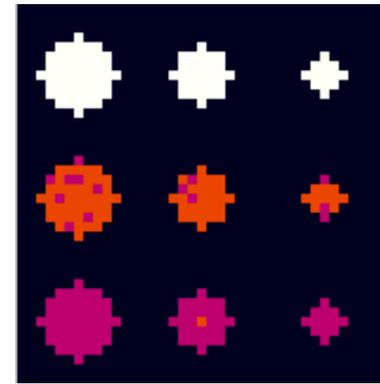
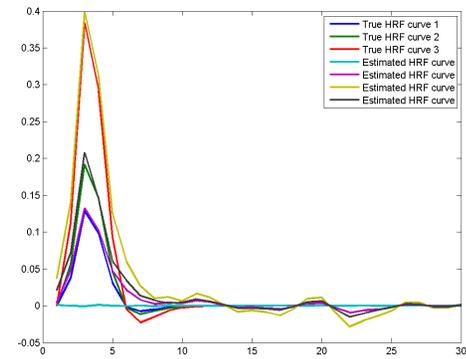
The estimates of the HRFs, from the left to right, are the 1st, 2nd and 3rd sequences of events.



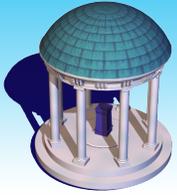
H1(t)



H2(t)



H3(t)

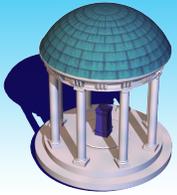


## Comparison with Lindquist *et al* (2009)

$D = \frac{1}{n} \frac{1}{m} \sum_{i=1}^n \sum_{j=1}^m (|\hat{x}_{ij} - x_0| - |\hat{y}_{ij} - x_0|)$  n is the sample size, m is the number of voxels in each active region.

Param	sFIR			IR			GAM					
	A1	A2	A3		A1	A2	A3		A1	A2	A3	
H	C1	-0.02 (0.045)	-0.04 (0.045)	-0.09 (0.060)	C1	-0.11 (0.0819)	-0.16 (0.115)	-0.32 (0.218)	C1	-0.13 (0.039)	-0.20 (0.044)	-0.42 (0.059)
	C2	-0.02 (0.041)	-0.04 (0.043)	-0.10 (0.057)	C2	-0.09 (0.0699)	-0.14 (0.096)	-0.27 (0.175)	C2	-0.11 (0.041)	-0.17 (0.045)	-0.36 (0.057)
	C3	-0.01 (0.046)	-0.02 (0.047)	-0.07 (0.064)	C3	-0.07 (0.0673)	-0.10 (0.089)	-0.21 (0.161)	C3	-0.07 (0.038)	-0.11 (0.045)	-0.25 (0.066)
T	C1	-0.08 (0.722)	0.05 (0.326)	0.01 (0.073)	C1	-3.74 (3.313)	-3.49 (3.309)	-3.19 (3.292)	C1	-2.50 (0.425)	-2.66 (0.298)	-2.84 (0.070)
	C2	-0.05 (0.685)	0.07 (0.292)	0.01 (0.069)	C2	-3.50 (3.440)	-3.34 (3.475)	-2.88 (3.419)	C2	-2.57 (0.452)	-2.75 (0.287)	-2.91 (0.069)
	C3	-0.55 (1.159)	-0.10 (0.513)	-0.10 (0.513)	C3	-3.54 (3.404)	-3.26 (3.430)	-3.03 (3.415)	C3	-2.46 (0.577)	-2.66 (0.427)	-2.87 (0.254)
W	C1	-0.20 (0.671)	-0.28 (0.596)	-0.42 (0.518)	C1	-1.70 (2.122)	-1.73 (2.127)	-1.66 (2.094)	C1	-3.33 (0.623)	-3.41 (0.576)	-3.46 (0.515)
	C2	-0.38 (0.760)	-0.41 (0.597)	-0.49 (0.513)	C2	-1.78 (2.143)	-1.80 (2.099)	-1.79 (2.018)	C2	-3.35 (0.634)	-3.41 (0.575)	-3.49 (0.512)
	C3	-0.32 (0.870)	-0.33 (0.741)	-0.48 (0.658)	C3	-1.79 (2.179)	-1.85 (2.123)	-2.08 (2.221)	C3	-3.30 (0.713)	-3.42 (0.677)	-3.63 (0.550)

Comparisons of the differences of the absolute errors between our method with smooth finite impulse response (sFIR), inverse logit (IL) and SPM canonical HRF (GAM), respectively. C1, C2 and C3 denotes the 1st, 2nd and 3rd sequences of events, respectively. A1, A2 and A3 denotes the 1st, 2nd and 3rd active regions. Values in the blanket are the standard deviations. H=Height, W=Width, T=Time-to-Peak.

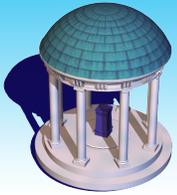


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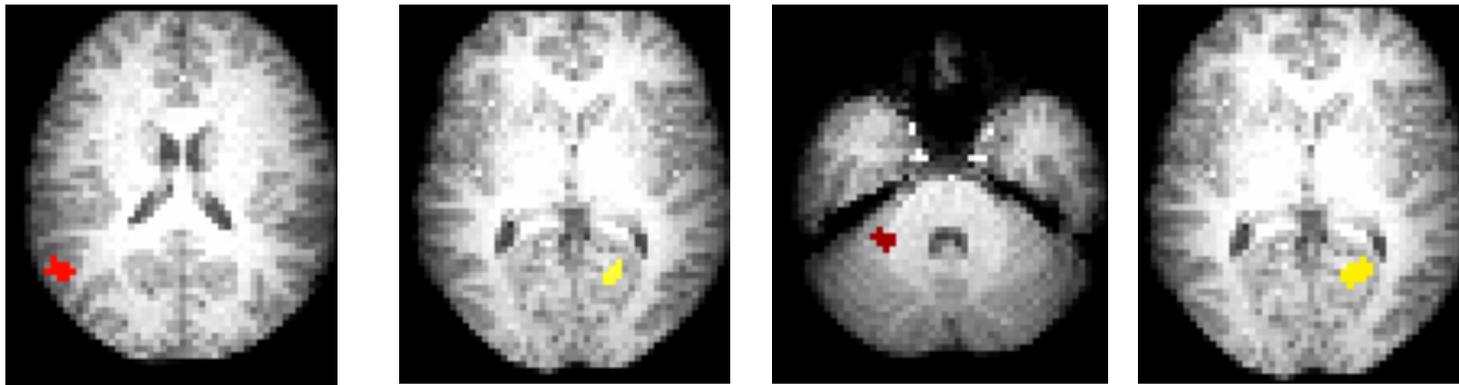
**This data set is from a memory related experiment to compare the neural correlates of relational memory during implicit (non-strategic) versus explicit (conscious, strategic) retrieval.**

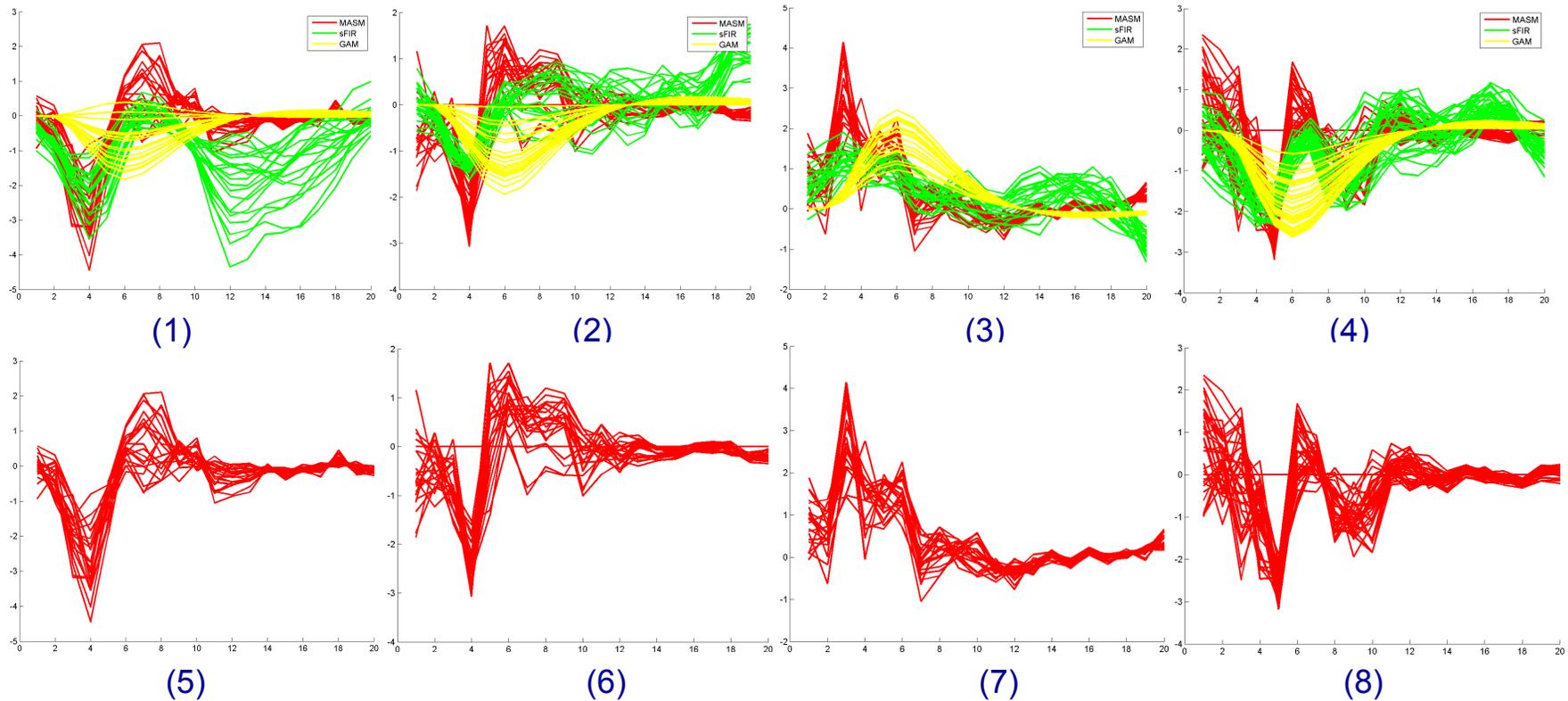
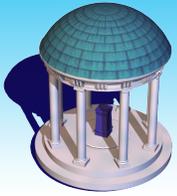
**There are four different sequences of stimuli.**

**We use SPM8 to preprocess the images including the realignment, timing slicing, segmentation, coregistration, normalization and spatial smoothing.**

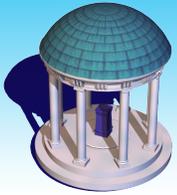


**We focus on some significant regions of interest (ROI) detected by SPM to study the HRFs of the voxels by our method. The results are verified by sFIR and GAM.**





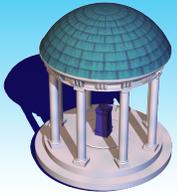
(1)-(4) Estimated HRFs at the significant ROIs corresponding each condition from MASM (red), sFIR(green) and GAM(yellow); (5)-(8) Estimated HRFs from only MASM in the each ROI.



## References

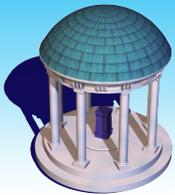
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- Li, YM, Zhu, HT, Shen, DG, Lin WL., Gilmore, J, Ibrahim, JG. (2010). Multiscale Adaptive Regression Models for Neuroimaging Data. *JRSSB*, in press.
- Li, YM, Ja-an Lin, M. Styner, John Gilmore, Zhu HT. Two-stage spatial adaptive analysis of twin neuroimaging data. *Submitted*.
- Li, YM, M. Styner, John Gilmore, Zhu HT. Spatial adaptive generalized estimating equations for longitudinal neuroimaging data. *Submitted*.
- Skup, M, Zhu, HT, Zhang HP. Spatial adaptive generalized Moment Estimation for longitudinal neuroimaging data. *Submitted*.
- Wang, J. Zhu, H.T., Fan, J.Q., Giovanello, K., and Lin, W. (2011). Multiscale Adaptive Smoothing Models for the Hemodynamic Response Function in fMRI. *Submitted*.



# Acknowledgements





## SI and SINS

### Establish ASA Section: **Statistics in Imaging (SI)**

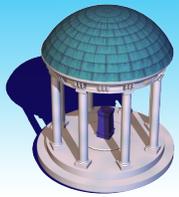
- Advisory board committee members for SI:  
Martin Lindquist, Daniel Rowe, Brian Caffo,  
Hernando Ombao, F. Dubois Bowman, Thomas Nichols,  
Ranjan Maitra, Hongtu Zhu, Kary Myers.



### Society of Imaging **Neuroscience** Statisticians (SINS)

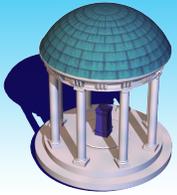
<http://www.mscs.mu.edu/~dbrowe/sins.html>

- Roundtables in ENAR 2011 and **JSM 2011.**



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# Simulation Studies

$$y_i(d) = x_i^T \beta(d) + \varepsilon_i(d)$$

$$\varepsilon_i(d) \sim N(0,1) \quad \varepsilon_i(d) \sim \chi^2(3) - 3$$

$$n = 60 \quad \text{or} \quad n = 80$$

$$x_i = (1, x_{i2}, x_{i3})^T$$

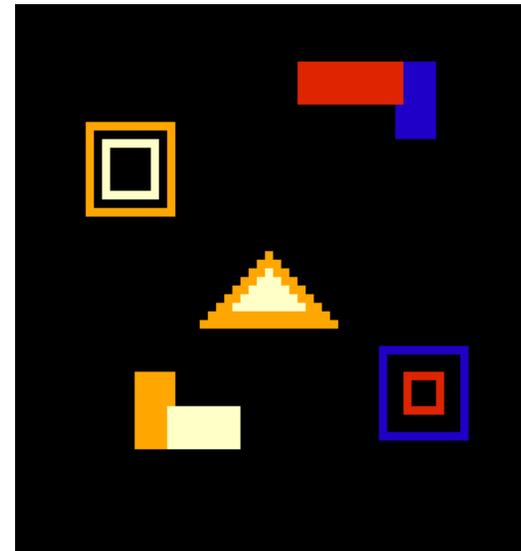
$$x_{i2} \sim \text{Bernoulli}(0.5)$$

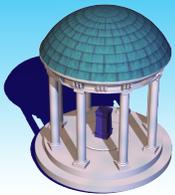
$$x_{i3} \sim \text{Uniform}[1,2]$$

$$\beta(d) = (\beta_1(d), \beta_2(d), \beta_3(d))^T$$

$$\beta_1(d) = \beta_2(d) = 0$$

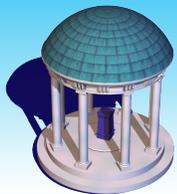
ROI	<i>black</i>	<i>blue</i>	<i>red</i>	<i>yellow</i>	<i>white</i>
$\beta_3(d)$	0.0	0.2	0.4	0.6	0.8





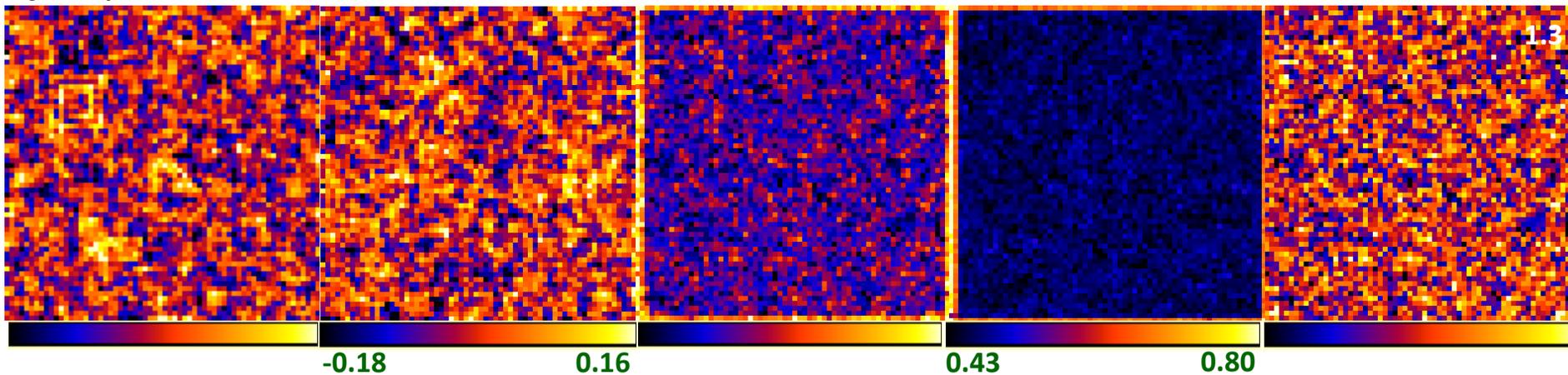
# Simulation Studies

$\beta_2(d)$		$\chi^2(3) - 3$						$N(0, 1)$					
		$n = 60$			$n = 80$			$n = 60$			$n = 80$		
		$h_0$	$h_5$	$h_{10}$	$h_0$	$h_5$	$h_{10}$	$h_0$	$h_5$	$h_{10}$	$h_0$	$h_5$	$h_{10}$
0.0	BIAS	0.00	0.00	0.01	0.00	0.00	0.00	0.00	0.00	0.00	0.00	0.00	0.00
	RMS	0.48	0.35	0.26	0.41	0.31	0.22	0.20	0.15	0.11	0.17	0.13	0.09
	SD	0.47	0.34	0.24	0.41	0.30	0.21	0.19	0.14	0.10	0.17	0.12	0.09
	RE	1.03	1.05	1.06	1.02	1.03	1.04	1.03	1.05	1.06	1.02	1.03	1.04
0.2	BIAS	0.00	-0.03	-0.07	0.01	-0.02	-0.06	0.00	-0.03	-0.05	0.00	-0.02	-0.05
	RMS	0.46	0.34	0.24	0.39	0.29	0.21	0.19	0.14	0.11	0.16	0.12	0.09
	SD	0.46	0.33	0.24	0.40	0.29	0.21	0.19	0.14	0.10	0.16	0.12	0.09
	RE	1.01	1.01	1.01	0.99	1.00	1.01	1.02	1.04	1.06	1.02	1.02	1.03
0.4	BIAS	-0.01	-0.05	-0.09	0.01	-0.02	-0.06	0.00	0.00	-0.01	0.00	0.00	0.00
	RMS	0.46	0.34	0.25	0.40	0.30	0.22	0.19	0.15	0.12	0.16	0.13	0.10
	SD	0.46	0.33	0.24	0.40	0.29	0.21	0.19	0.14	0.11	0.16	0.12	0.09
	RE	1.01	1.02	1.03	1.01	1.02	1.03	1.03	1.05	1.07	1.00	1.01	1.02
0.6	BIAS	0.00	-0.05	-0.09	0.00	-0.04	-0.07	0.00	0.01	0.02	0.00	0.00	0.01
	RMS	0.46	0.35	0.26	0.40	0.30	0.23	0.19	0.15	0.12	0.16	0.13	0.10
	SD	0.46	0.34	0.25	0.40	0.30	0.22	0.19	0.14	0.11	0.16	0.13	0.10
	RE	1.01	1.03	1.04	1.01	1.02	1.03	1.02	1.04	1.06	1.01	1.03	1.04
0.8	BIAS	0.00	-0.04	-0.06	0.00	-0.02	-0.05	0.00	-0.01	-0.02	0.00	0.00	-0.01
	RMS	0.47	0.35	0.26	0.40	0.30	0.23	0.19	0.15	0.11	0.17	0.13	0.10
	SD	0.46	0.34	0.25	0.40	0.30	0.22	0.19	0.14	0.11	0.16	0.12	0.09
	RE	1.02	1.03	1.04	1.01	1.02	1.03	1.02	1.04	1.05	1.03	1.05	1.06

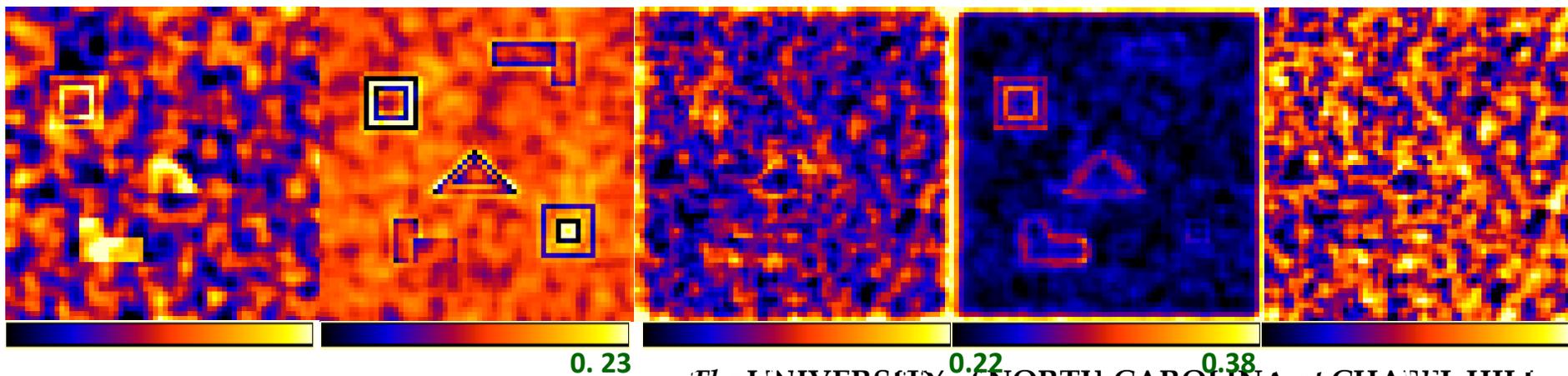


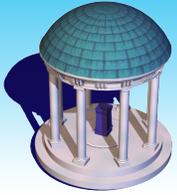
# Simulation Studies

$$\hat{\beta}_3(d, h_0)$$

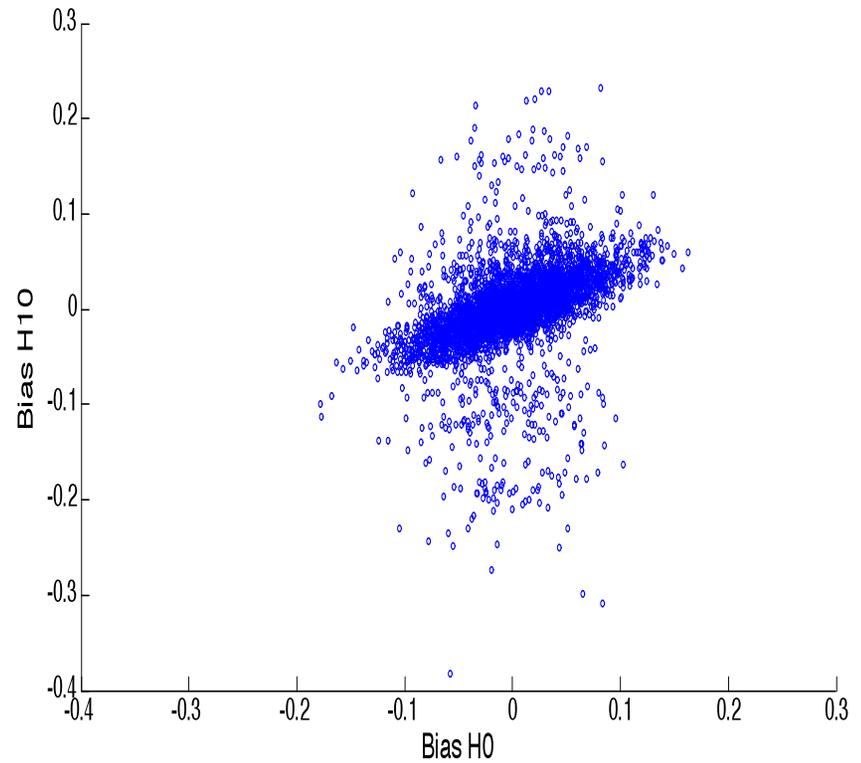
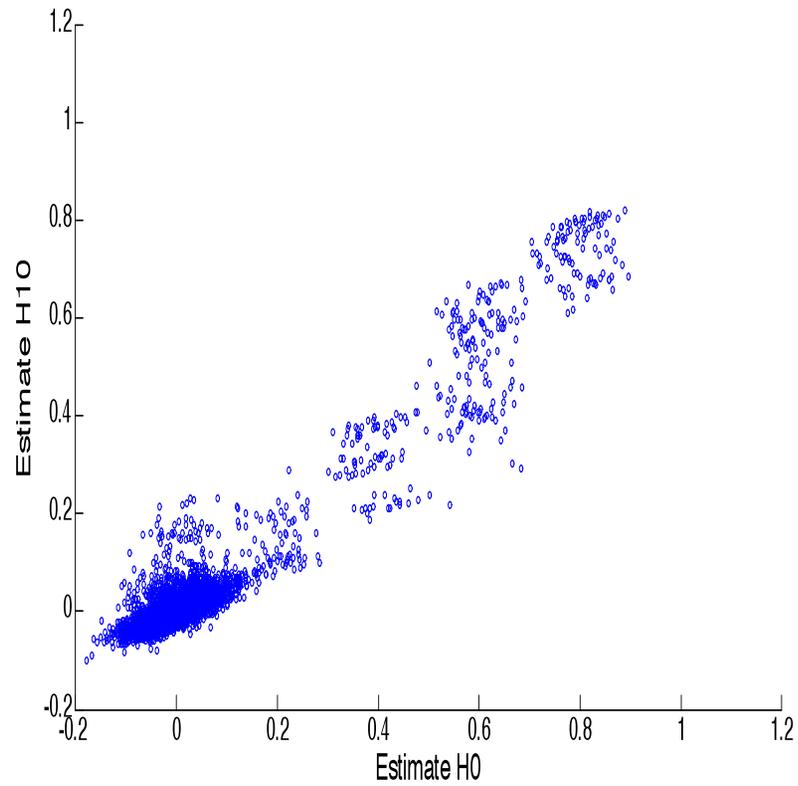


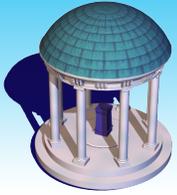
$$\hat{\beta}_3(d, h_{10})$$



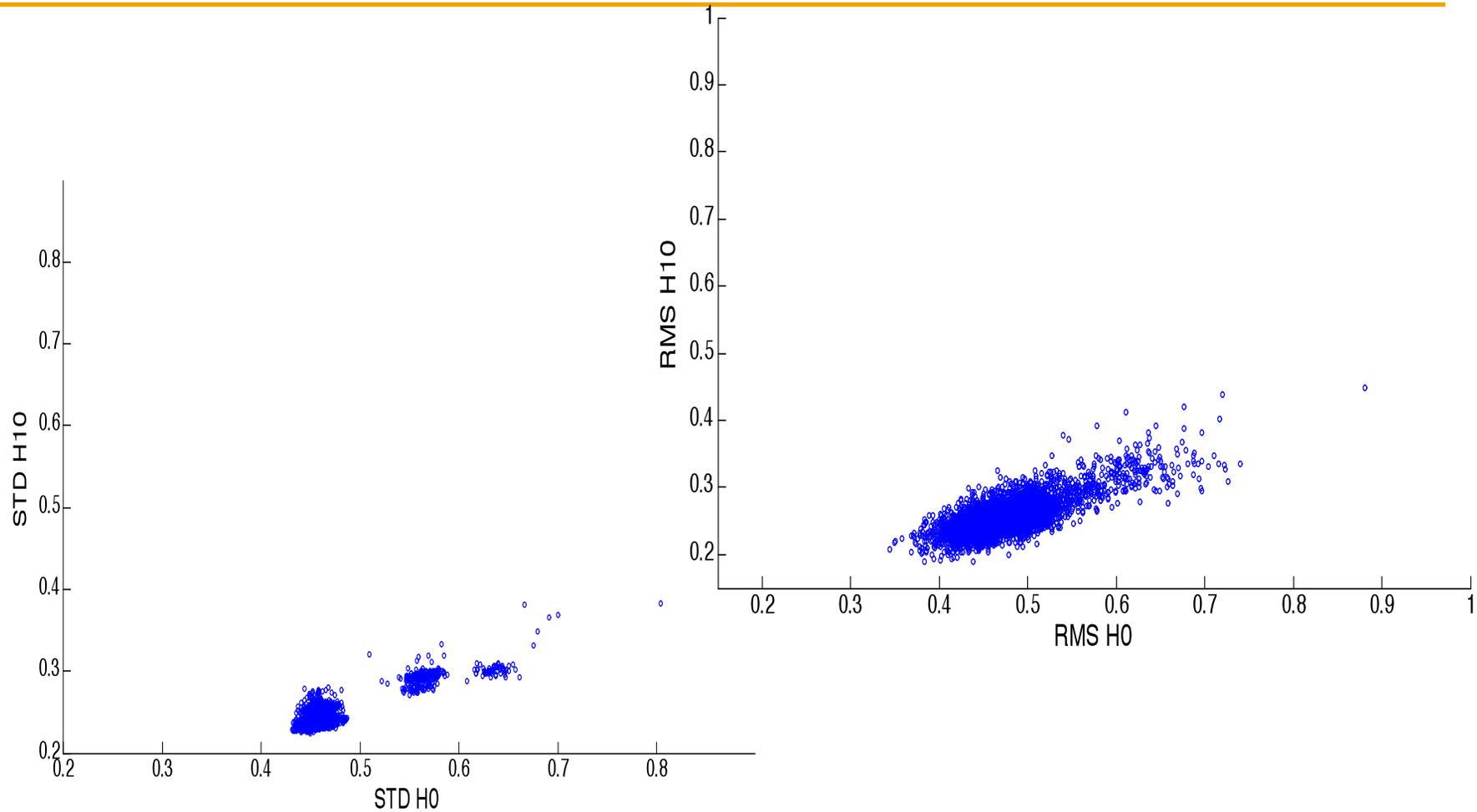


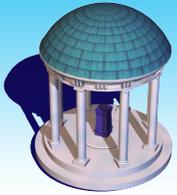
# Simulation Studies



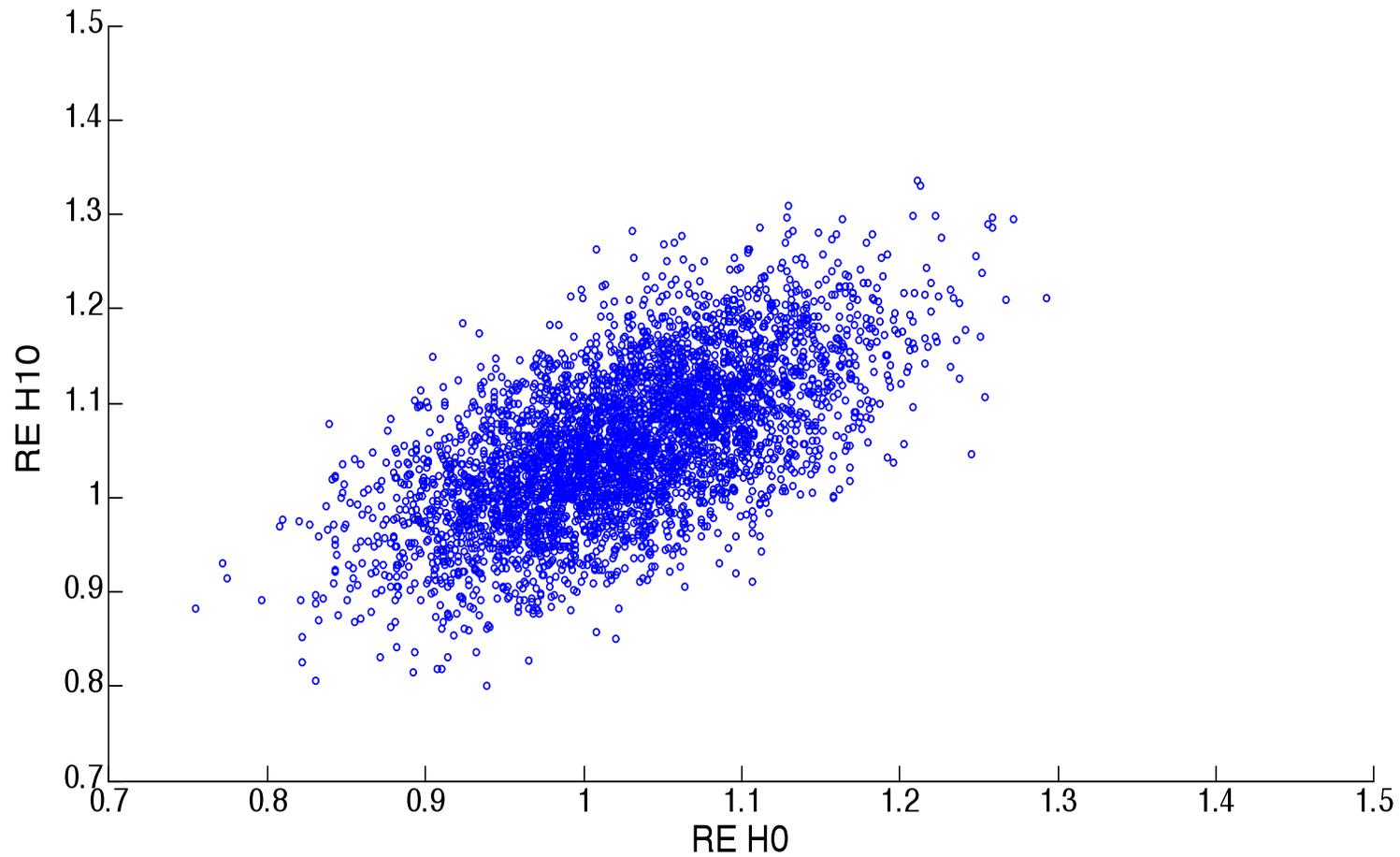


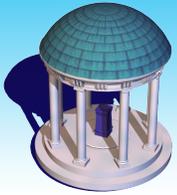
# Simulation Studies





# Simulation Studies

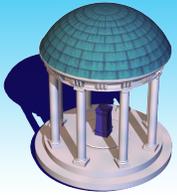




# Simulation Studies

**Table 2.** Simulation study for  $W_\mu(d, h)$ : estimates (ES) and standard errors (SE) of rejection rates for pixels inside the five ROIs were reported at 2 different scales ( $h_0, h_{10}$ ), 2 different distributions ( $N(0, 1)$  and  $\chi^2(3) - 3$ ), and 2 different sample sizes ( $n = 60, 80$ ) at  $\alpha = 5\%$ . For each case, 1,000 simulated data sets were used.

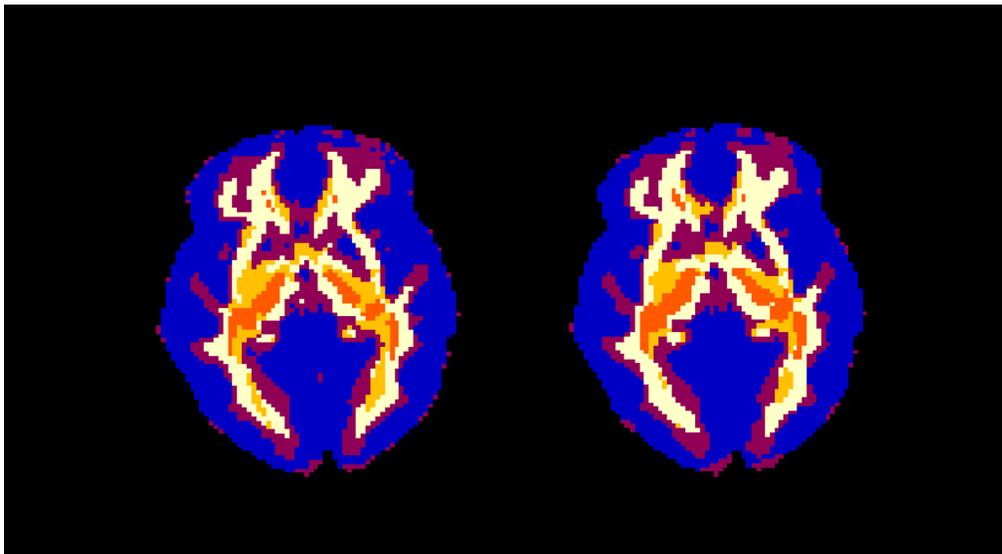
$\beta_2(d)$	$s$	$N(0, 1)$				$\chi^2(3) - 3$			
		$n = 60$		$n = 80$		$n = 60$		$n = 80$	
		ES	SE	ES	SE	ES	SE	ES	SE
0.2	$h_0$	0.20	0.066	0.24	0.070	0.08	0.038	0.08	0.037
	$h_{10}$	0.30	0.126	0.38	0.121	0.10	0.069	0.18	0.081
0.4	$h_0$	0.56	0.090	0.67	0.079	0.15	0.065	0.18	0.070
	$h_{10}$	0.93	0.051	0.98	0.030	0.26	0.129	0.35	0.159
0.6	$h_0$	0.88	0.039	0.95	0.024	0.27	0.057	0.33	0.050
	$h_{10}$	1.00	0.004	1.00	0.004	0.51	0.091	0.63	0.083
0.8	$h_0$	0.99	0.015	1.00	0.005	0.43	0.080	0.52	0.080
	$h_{10}$	0.99	0.010	0.99	0.011	0.78	0.099	0.90	0.006
0.0	$h_0$	0.07	0.006	0.07	0.006	0.06	0.007	0.07	0.006
	$h_{10}$	0.08	0.011	0.07	0.011	0.07	0.012	0.08	0.012



## Early Brain Development: Structural connectivity

Comparison:

**MAGEEC(0)** versus **MAGEEC(5)**



Model Selection:

