# Software tools motivated by analysis of fMRI data

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UNIVERSITY OF MINNESOTA

# Introduction

### Where I am

 Center for Magnetic Resonance Research, University of Minnesota

### What I work on

- Computational models of visual processing
- Object and form vision

### Approach

- Cognitive neuroscience (experiments, fMRI)
- Theoretical neuroscience (modeling)
- Data analysis (stats, programming)

### Resources

http://cvnlab.net





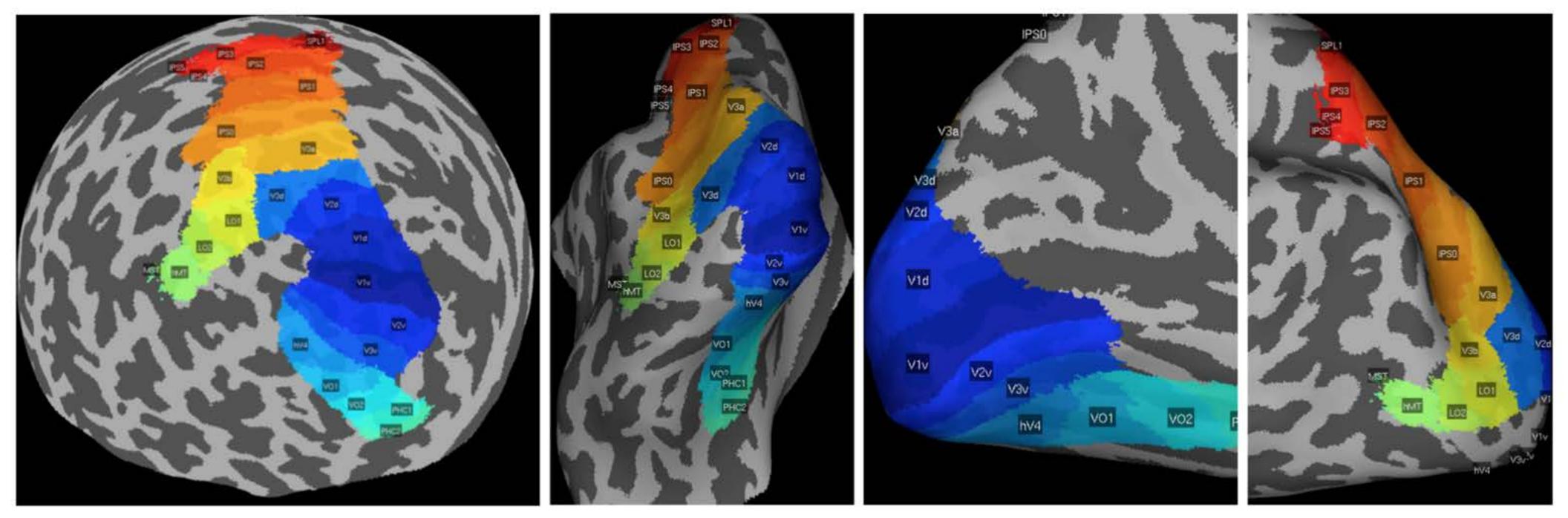
http://www.cmrr.umn.edu

# **Computing approach**

- Pull bits and pieces from:
  - FreeSurfer
  - SPM
  - FSL
- Integrate into MATLAB pipelines Some standalone MATLAB toolboxes (GLMdenoise, analyzePRF, etc.) Analysis is done mostly on a large workstation, using cluster computing for parallel analysis of
- individual voxels

http://github.com/kendrickkay/

- High-throughput (avoid GUI, automated)
- Customizable (colormap, overlays, etc.)
- FreeSurfer-oriented but could be generalized
- Support for high-resolution surfaces
- Method: map pixels to vertices using nearest-neighbor interpolation, use caching mechanism for speed



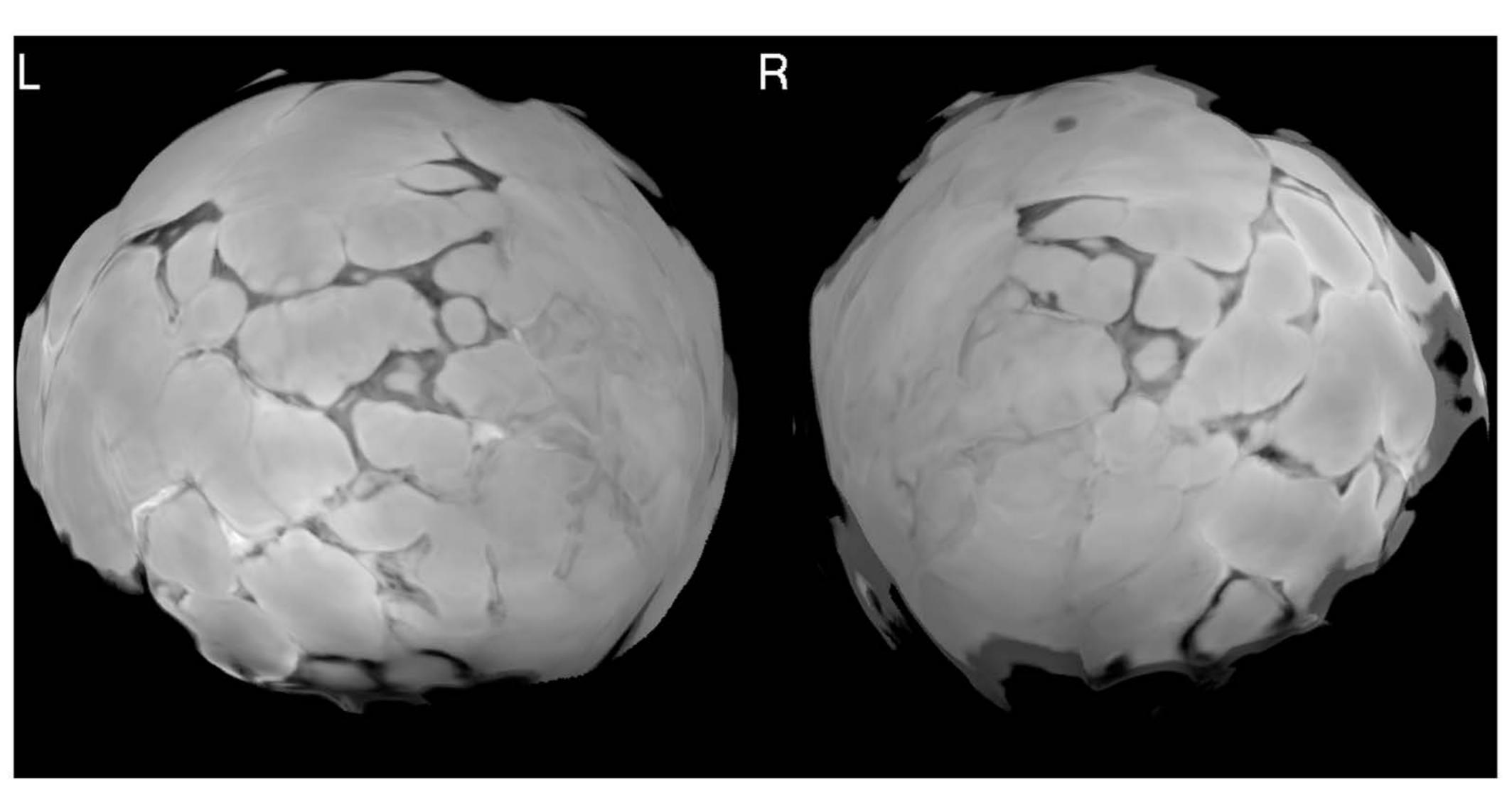
### sphere

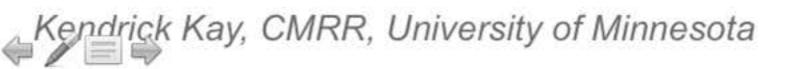
### occipital

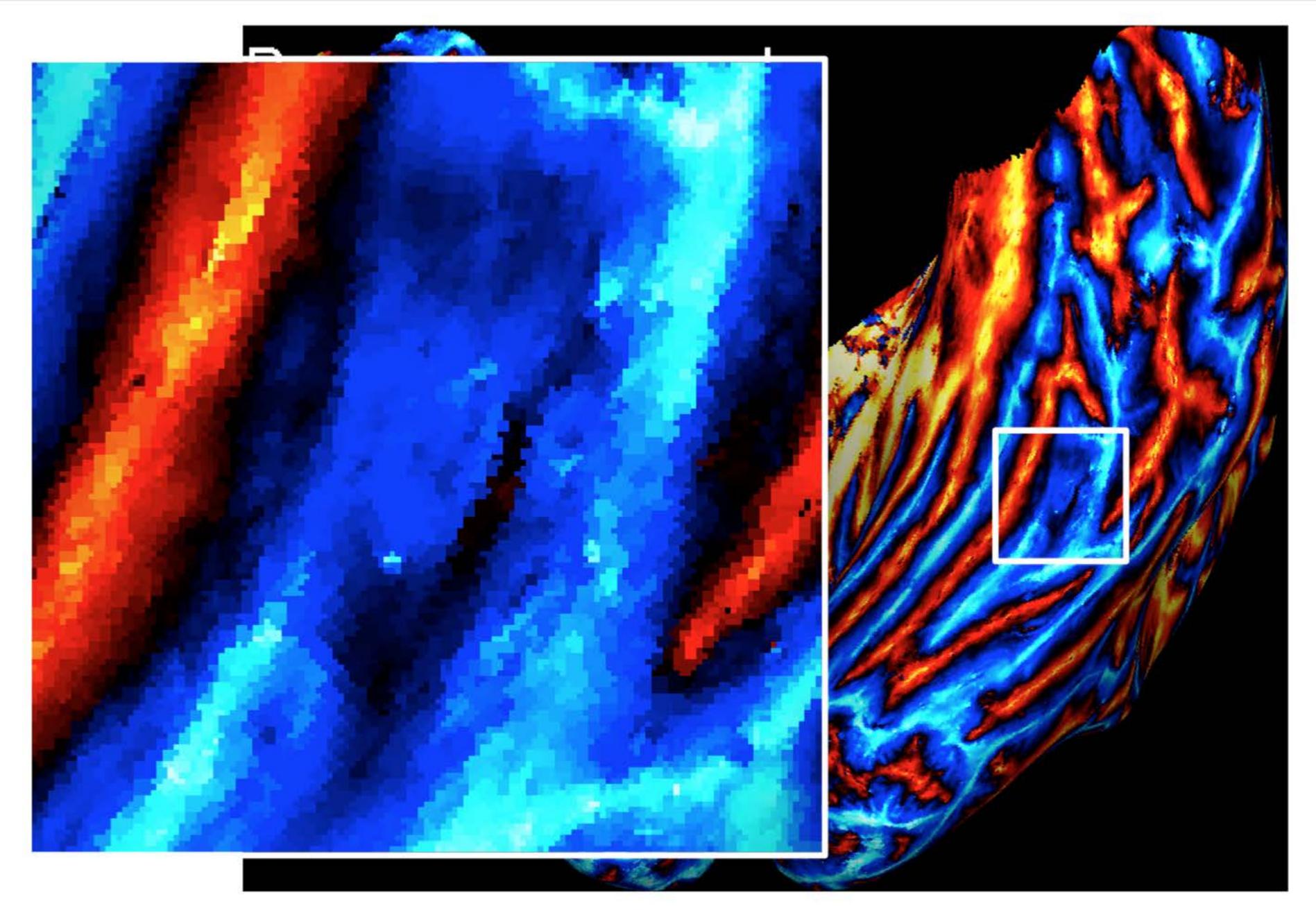
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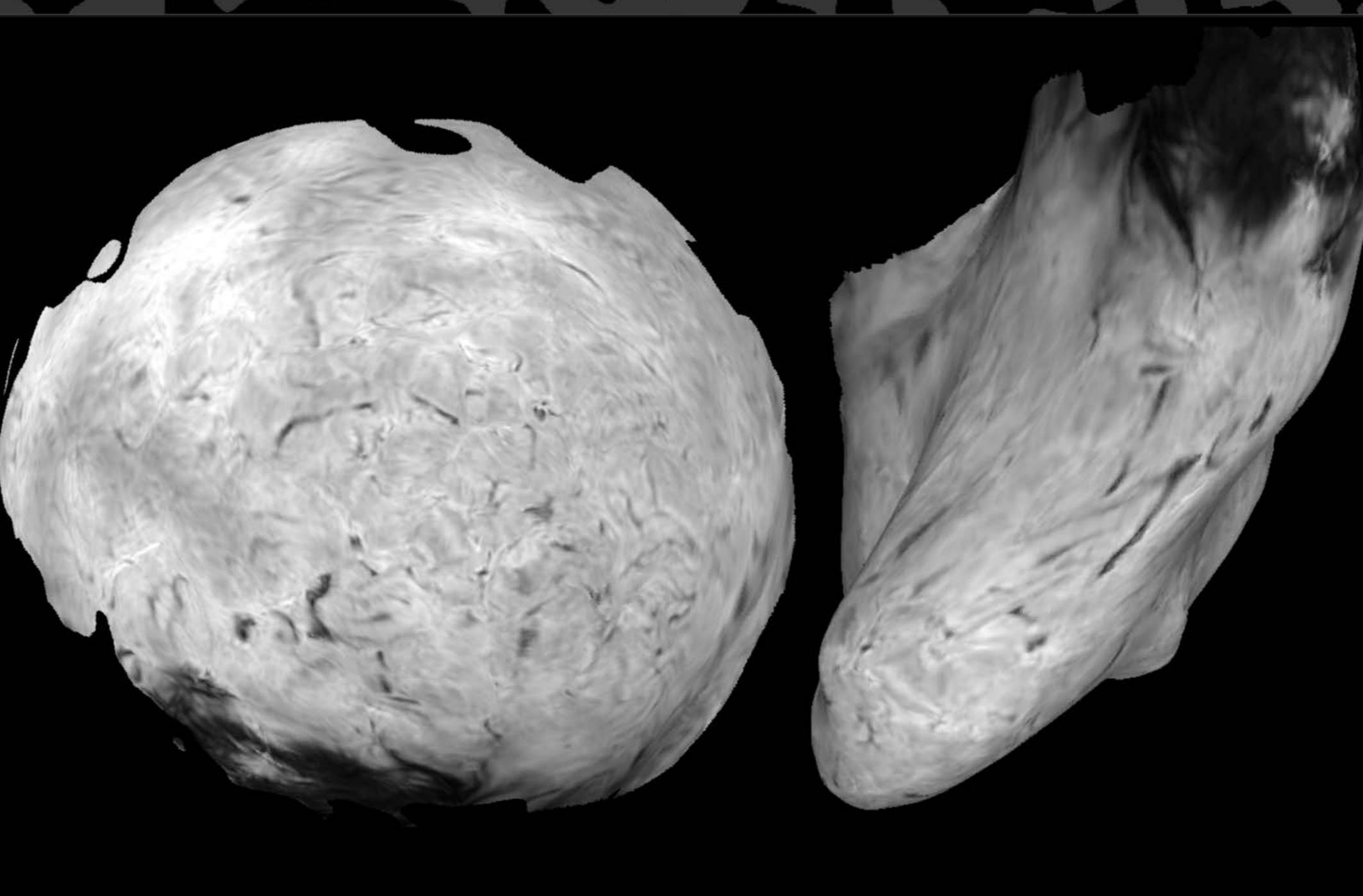
### medial

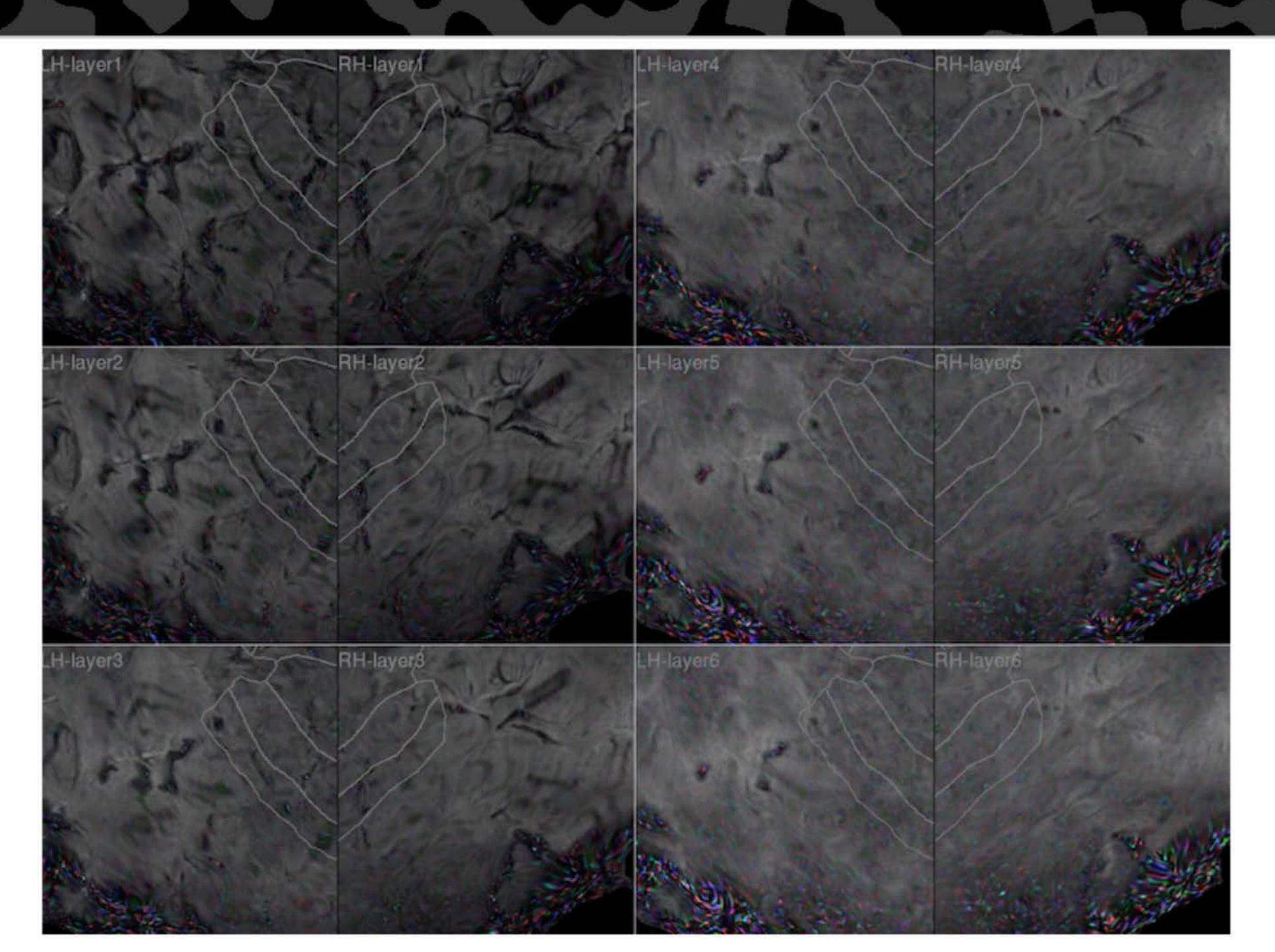
#### lateral









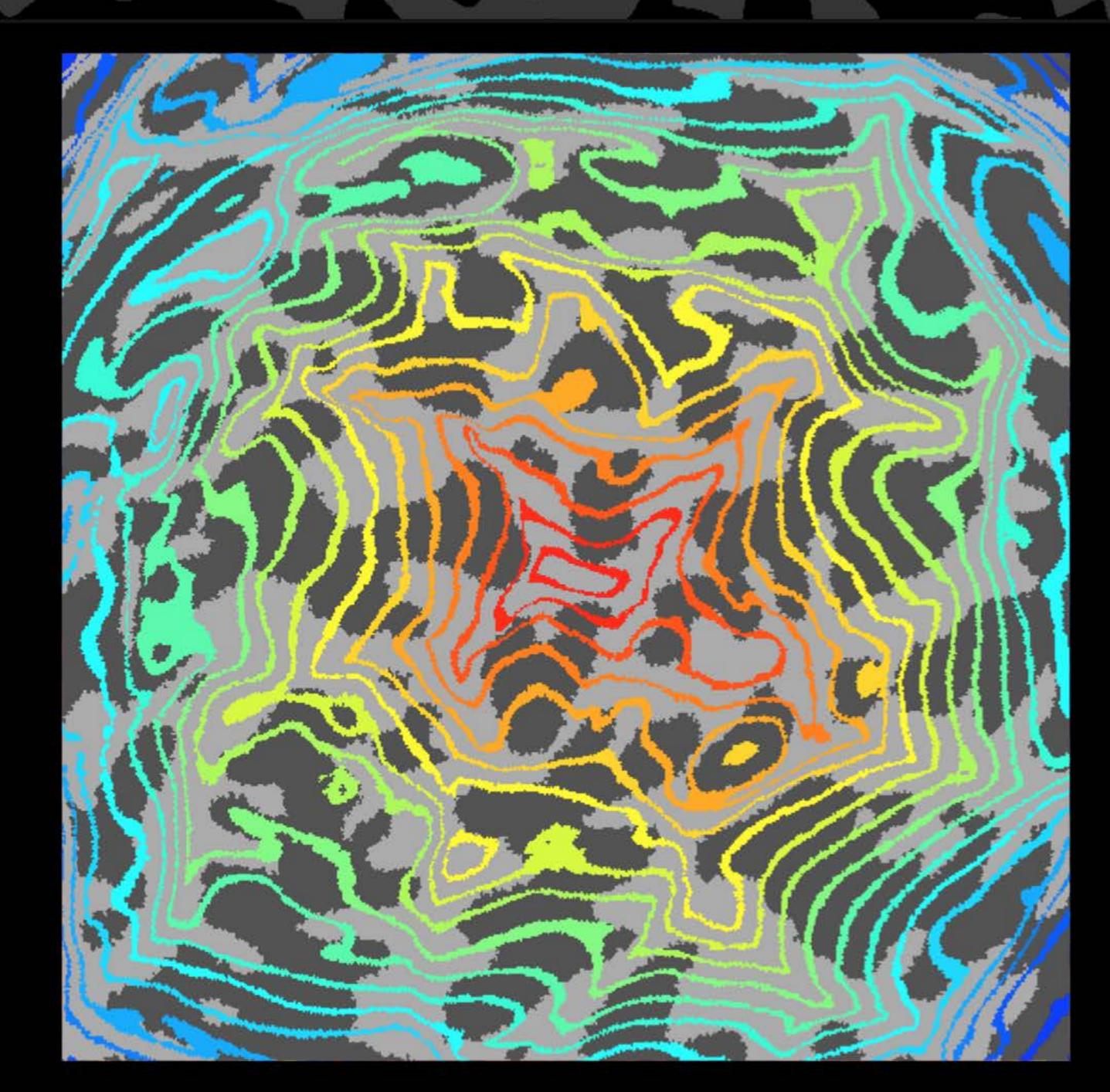


### Brain art (2/7)

Real data masquerading as art!



# Brain art (2/7)



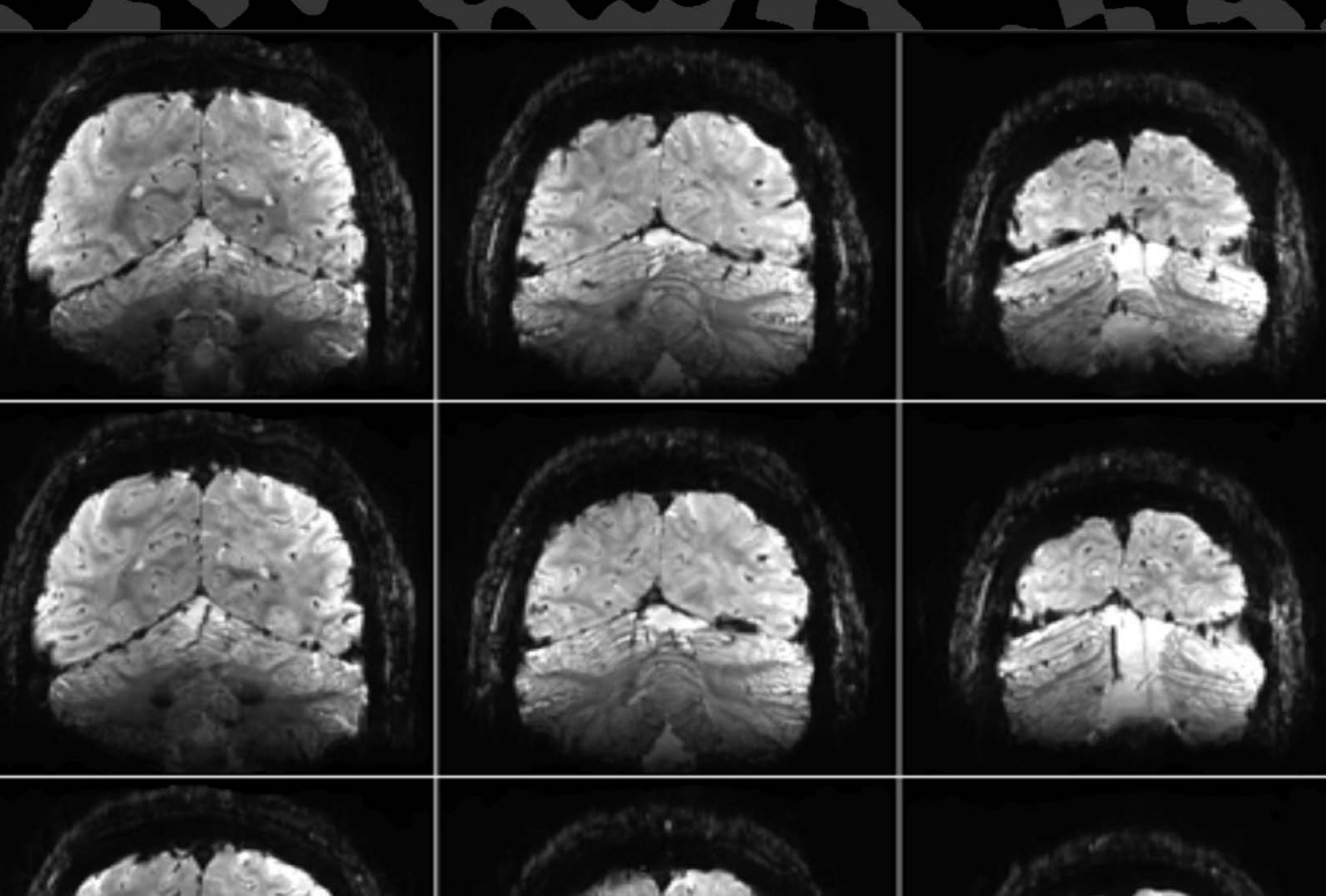
# Brain art (2/7)

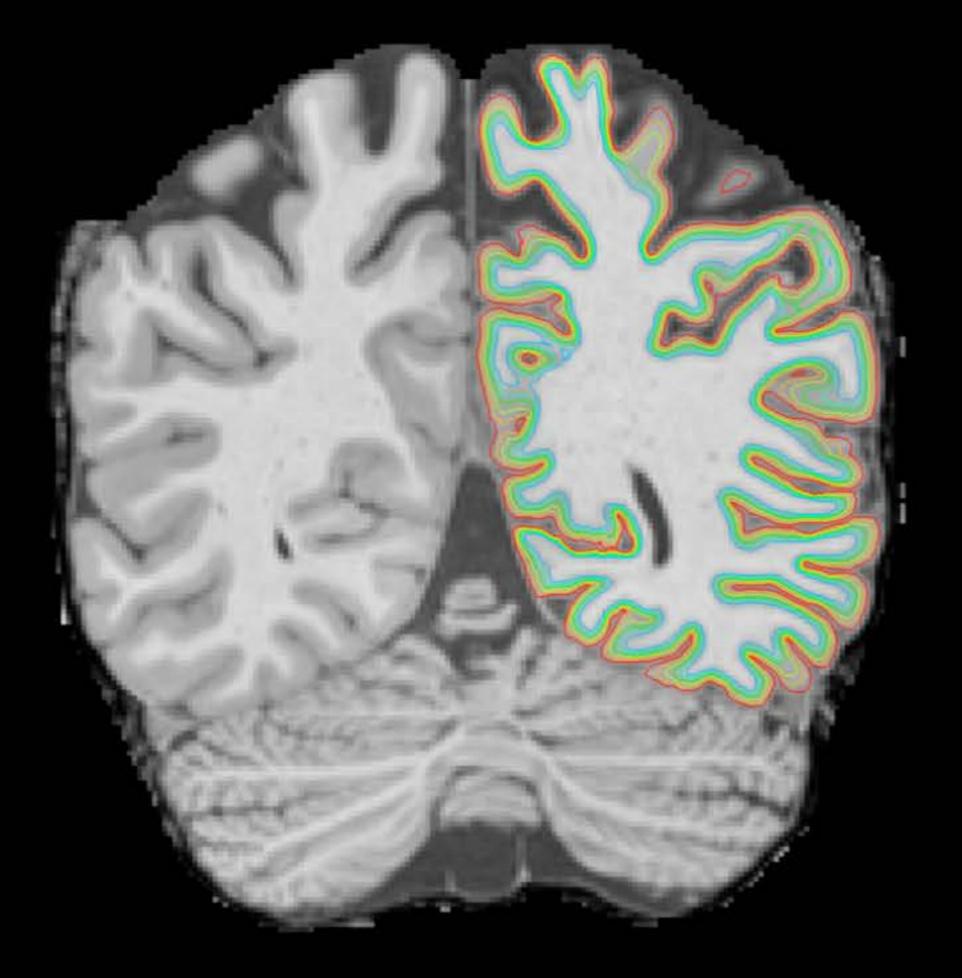


- Average T1s to improve SNR
- Co-registration: T1 to T1, T2 to T2, T2 to T1, EPI to T2
- Generate FreeSurfer cortical surfaces (dense, equidistant layers, truncated)
- Fieldmaps:
  - Regularize using local linear smoothing
  - Interpolate over time
- EPI:
  - Slice time correction
  - Motion correction
  - Fieldmap undistortion
  - Interpolation onto cortical surfaces
  - Total: 1 temporal resampling, 1 spatial resampling
- Homogenization of EPI intensities (polynomial basis functions, surface-based)

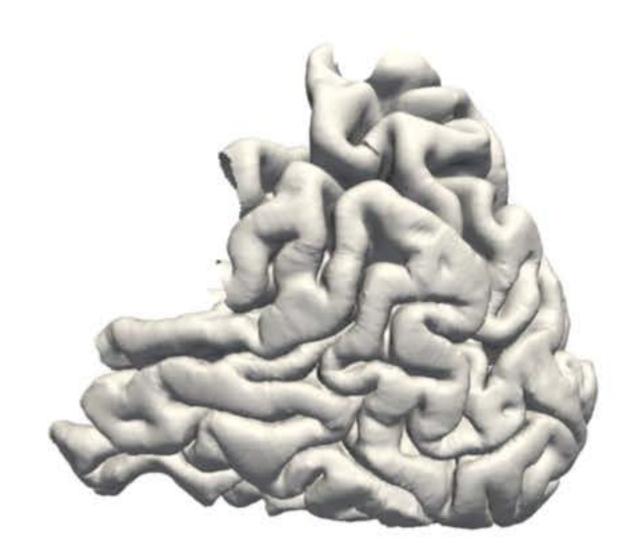
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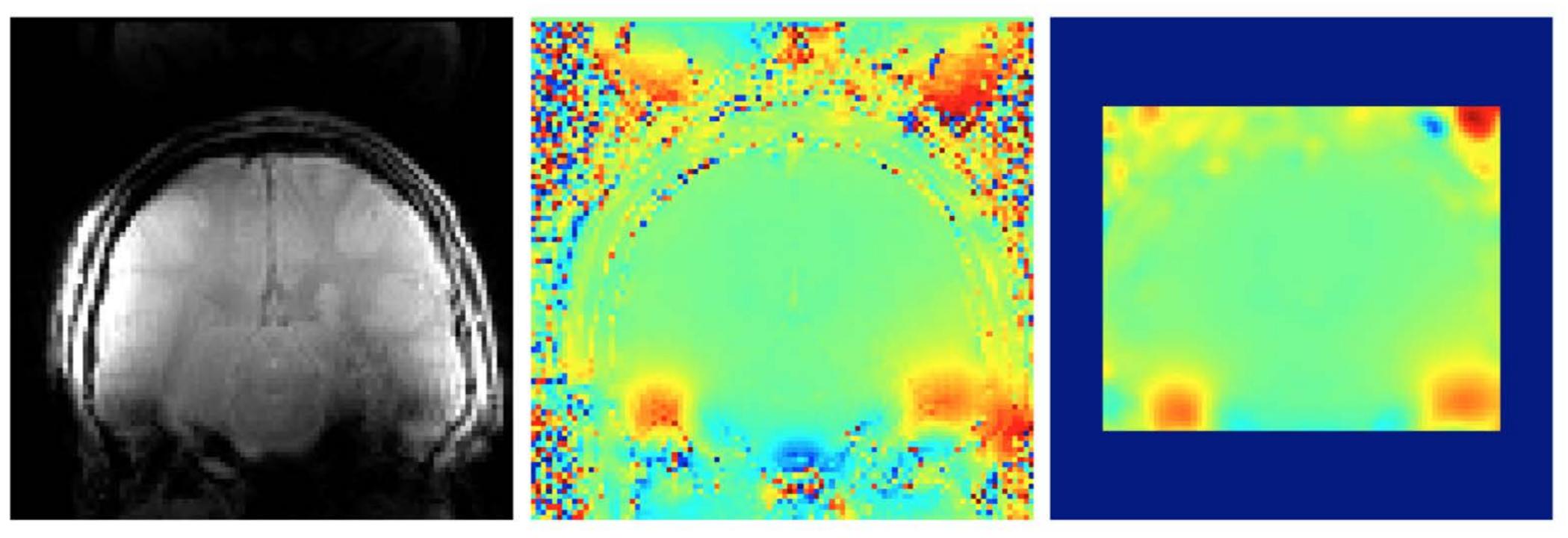
#### http://github.com/kendrickkay/cvncode/









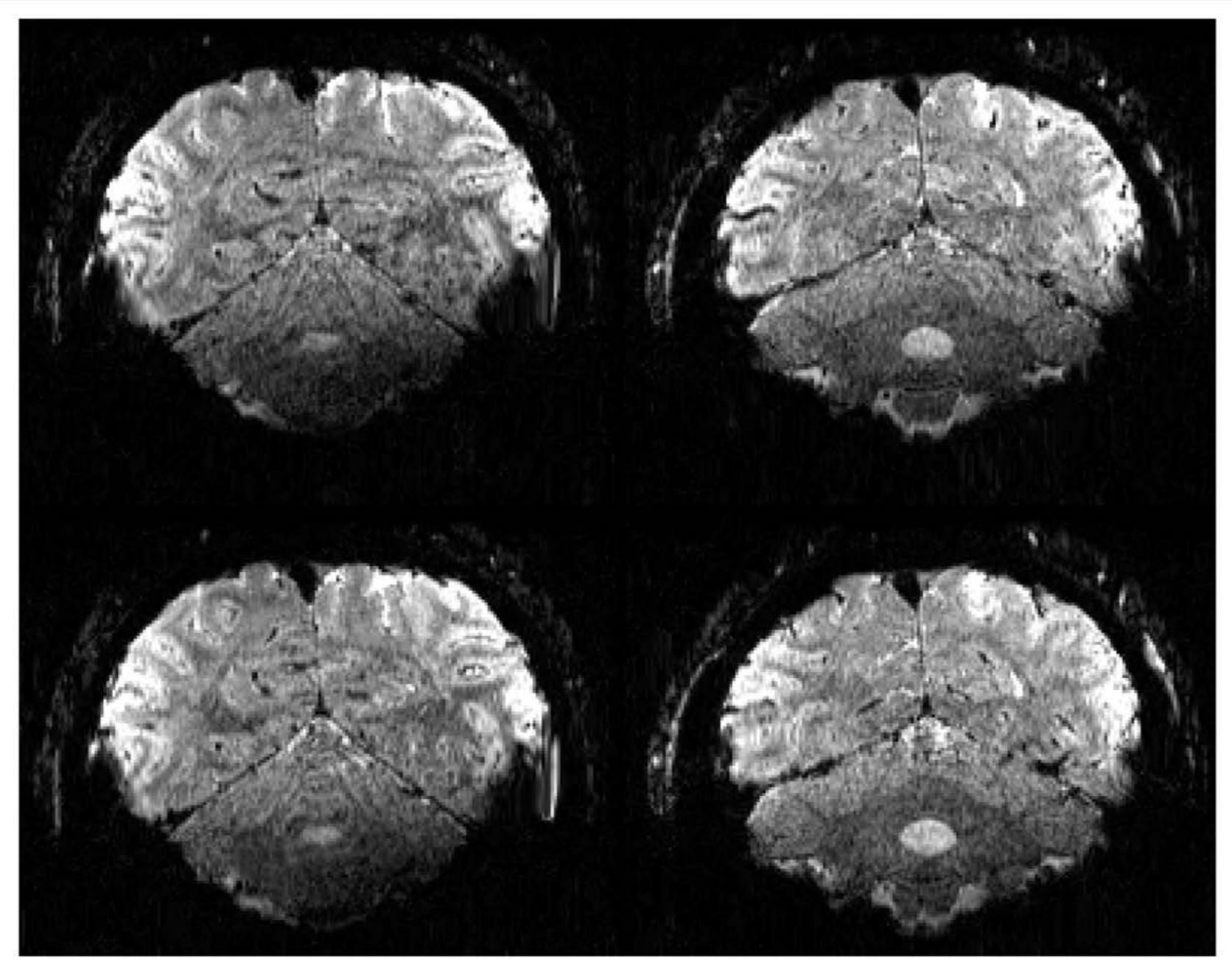


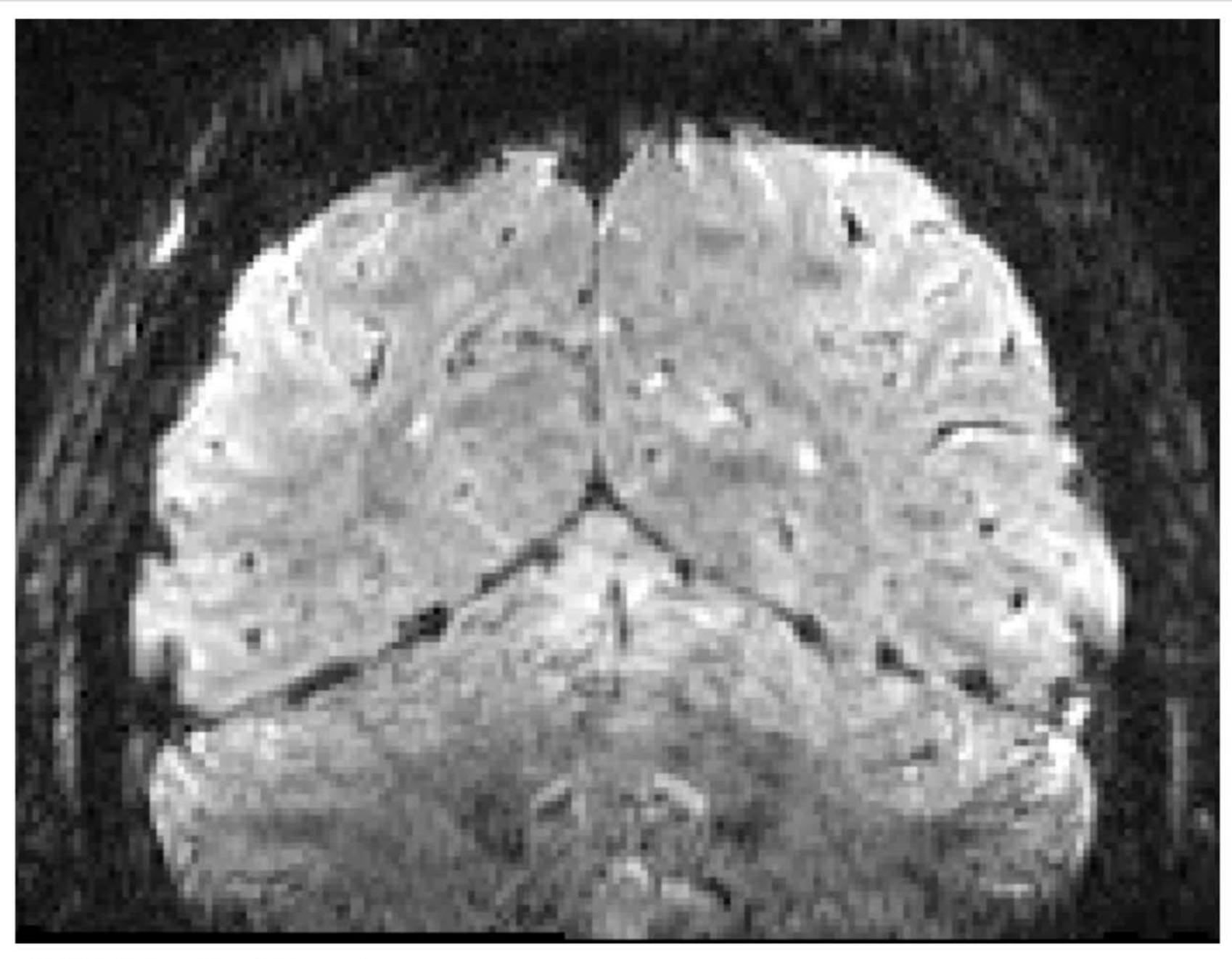
#### Fieldmap magnitude

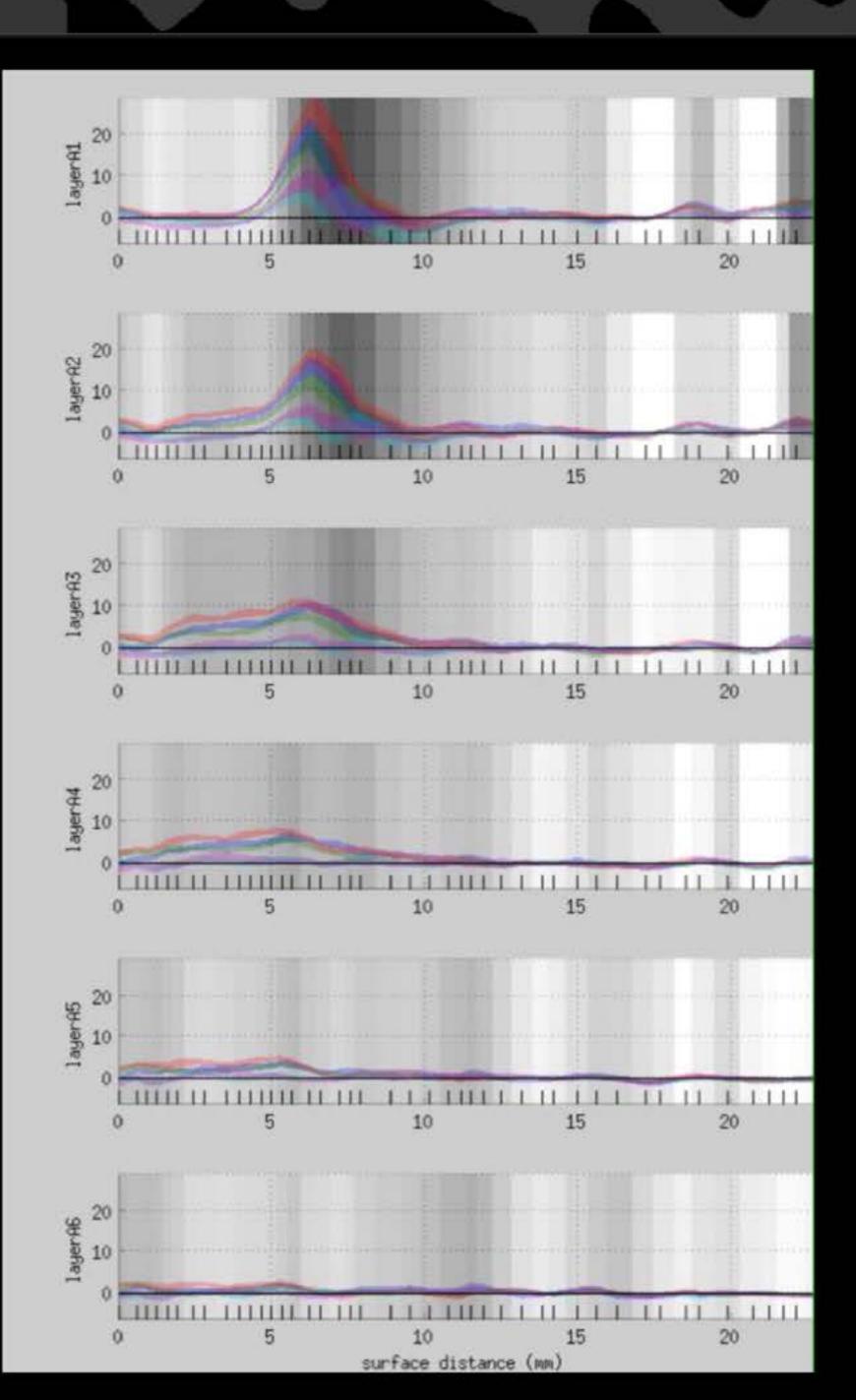
Fieldmap phase

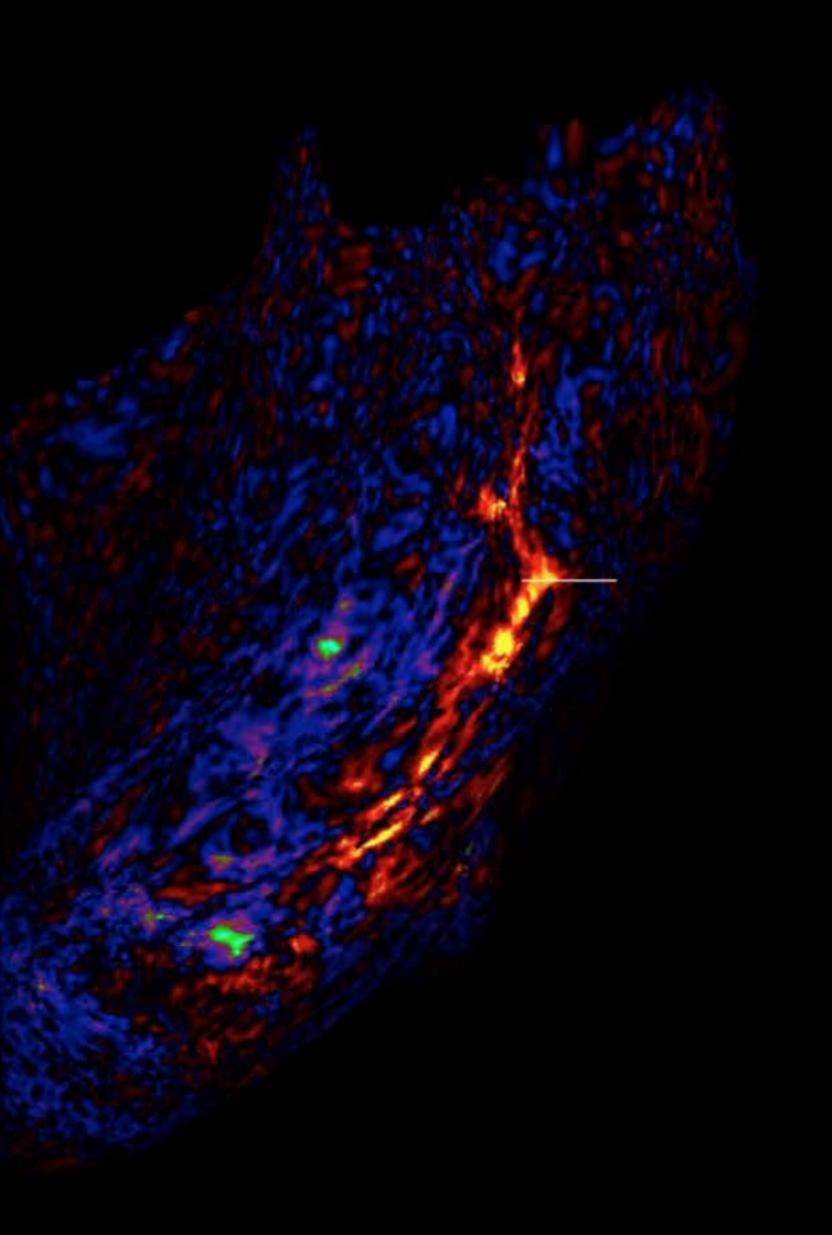
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# Fieldmap phase (regularized)









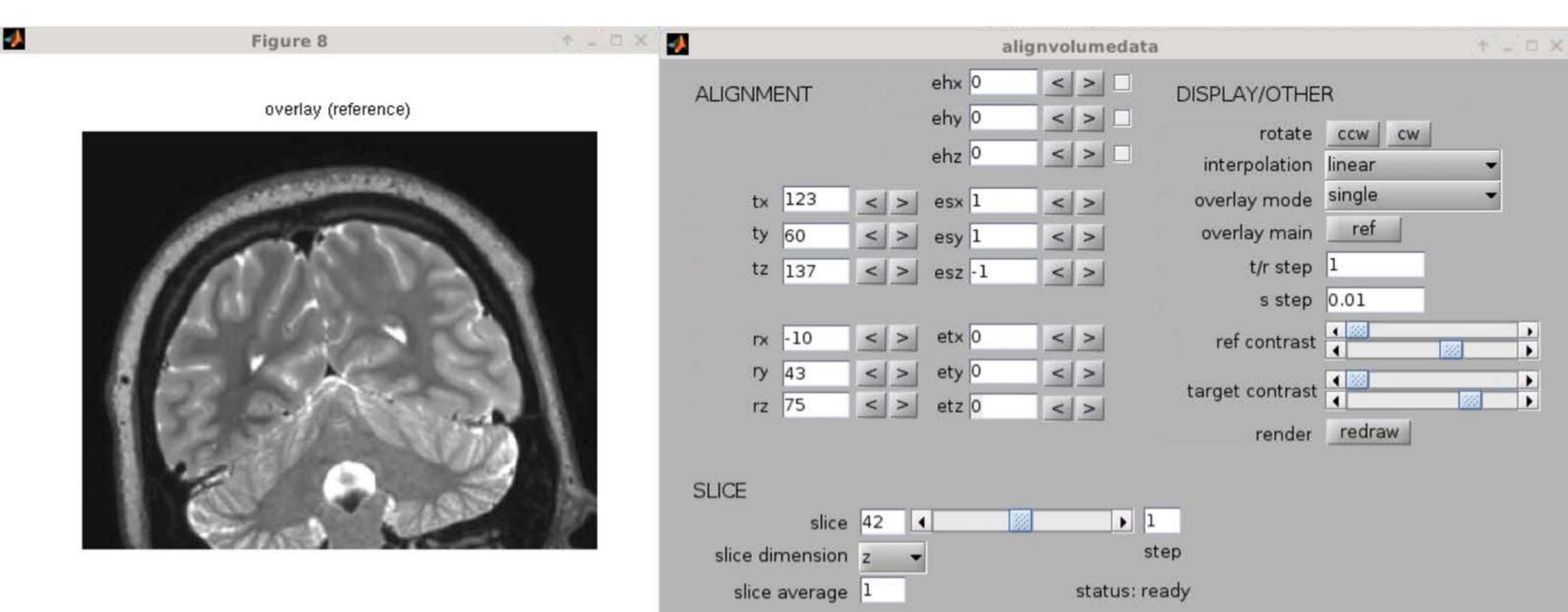
### Volume co-registration (4/7)

http://github.com/kendrickkay/alignvolumedata/

- Flexible inputs (any two volumes)
- Manual adjustment or automatic optimization
- Can use spatial mask
- Rigid-body or affine transformation

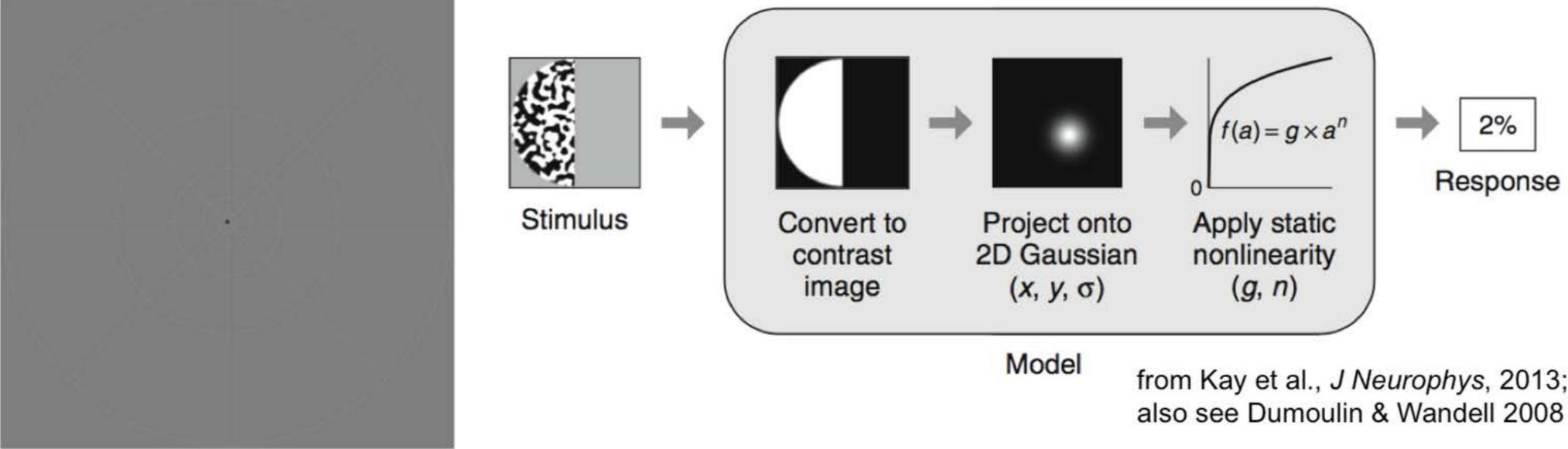
### Volume co-registration (4/7)

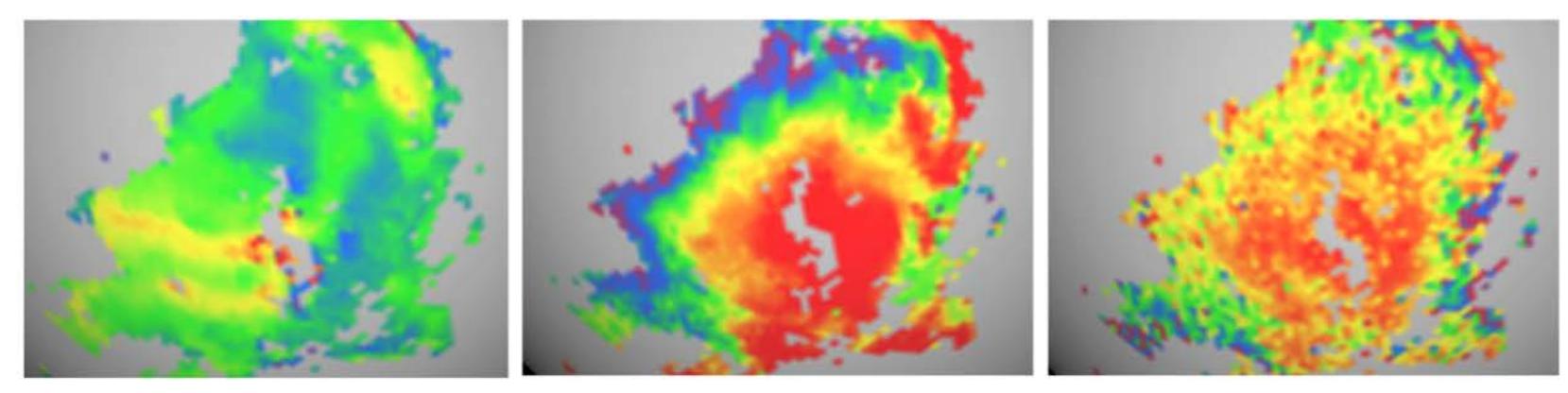
#### http://github.com/kendrickkay/alignvolumedata/



# analyzePRF (5/7) http://kendrickkay.net/analyzePRF/

Fit a parametric model that characterizes stimulus-response mapping





Angle

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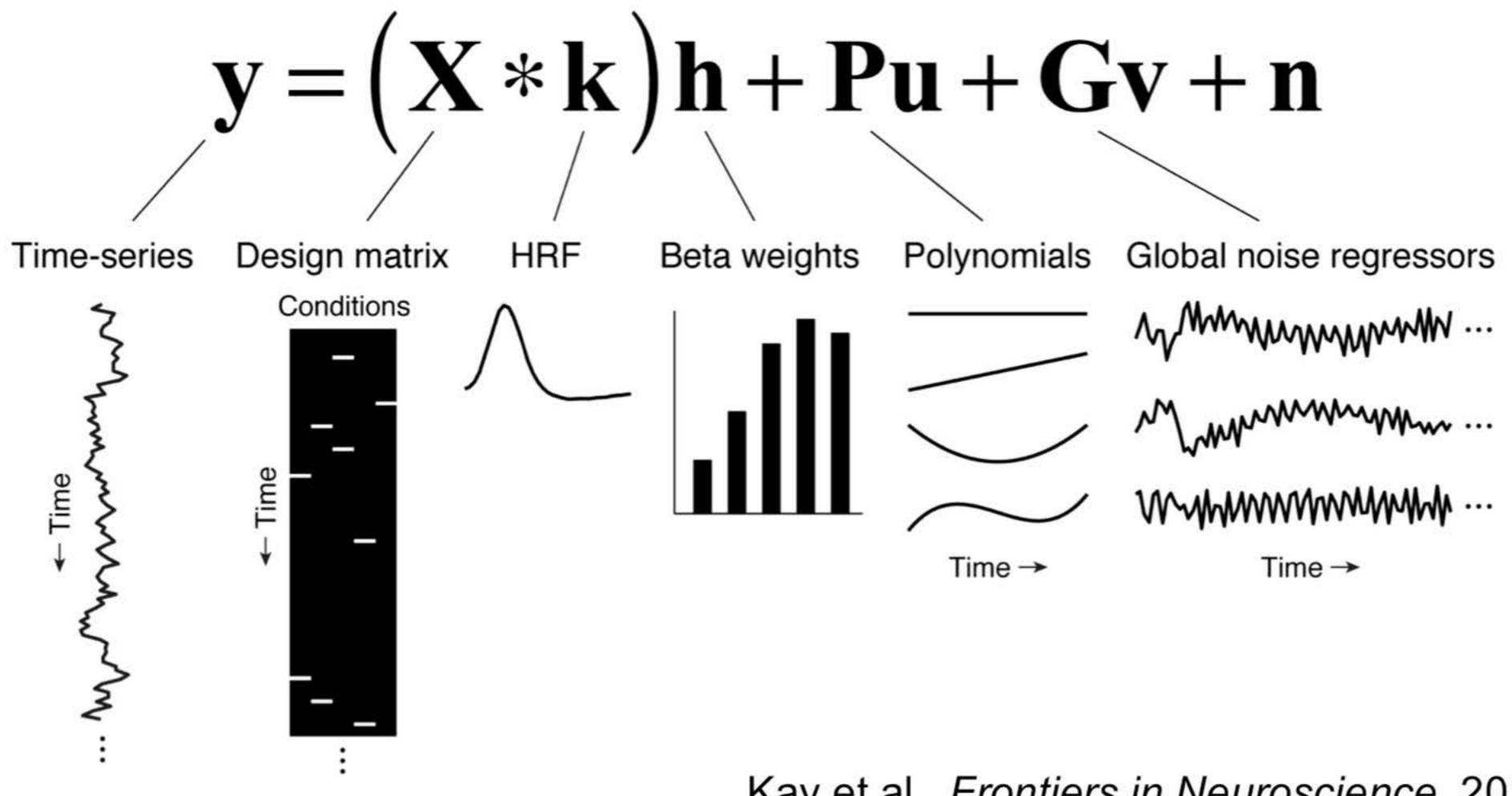
Eccentricity

from Kay et al., J Neurophys, 2013; also see Dumoulin & Wandell 2008

**RF** size

## GLMdenoise (6/7)

Fit a GLM that derives noise regressors and produces denoised beta weights

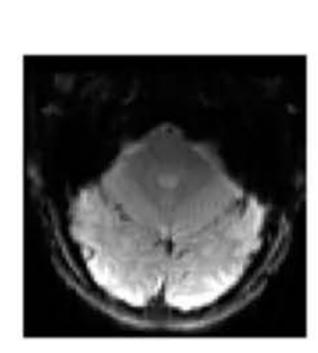


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http://kendrickkay.net/GLMdenoise/

Kay et al., Frontiers in Neuroscience, 2013

## GLMdenoise (6/7)



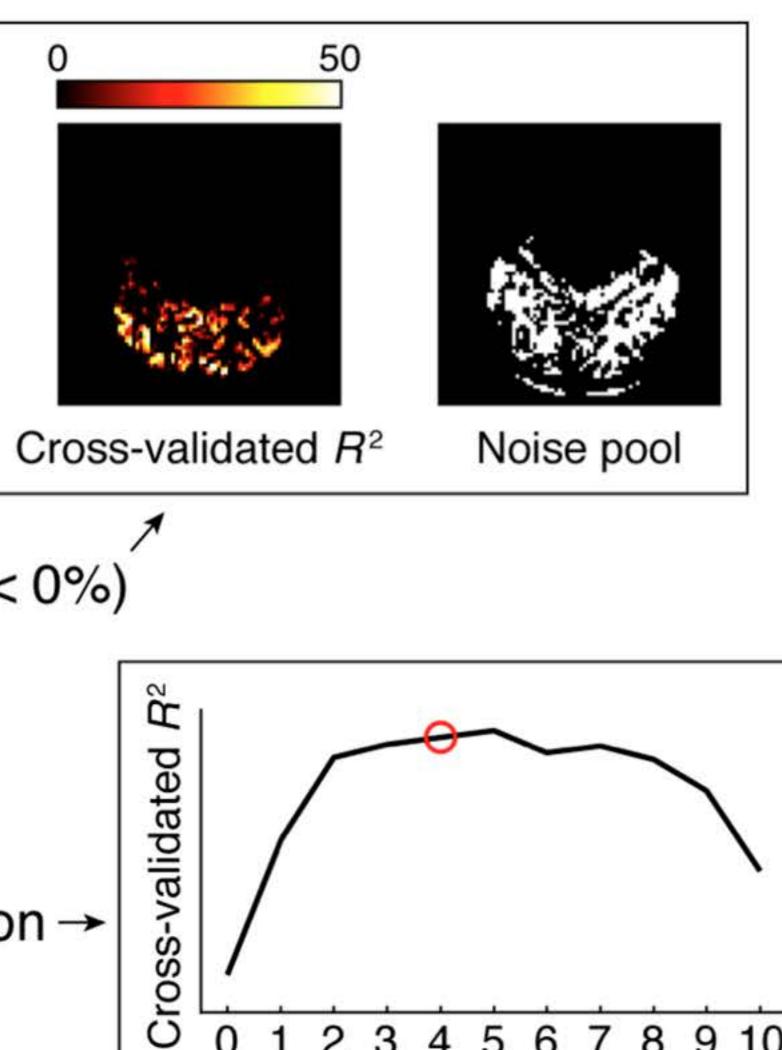
Perform initial model fit

Raw MR signal

- 2. Determine noise pool (cross-validated  $R^2 < 0\%$ )
- 3. Perform PCA on noise pool
- 4. Add PCs into the model, one at a time
- Select number of PCs using cross-validation →
- 6. Fit final model to the full dataset

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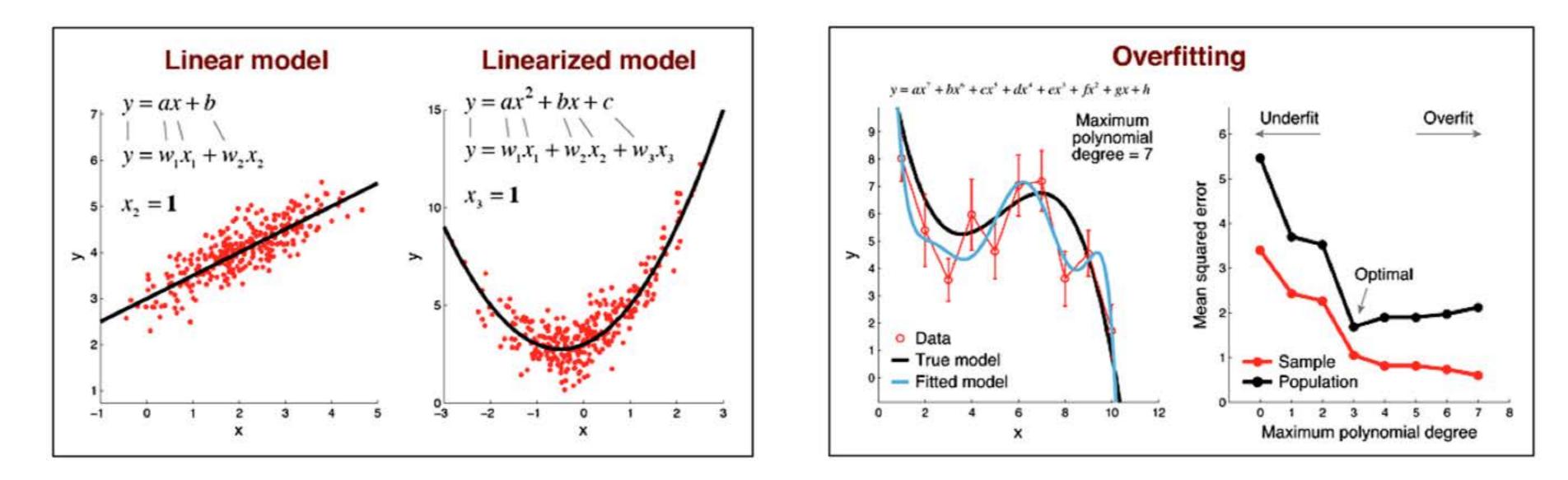
http://kendrickkay.net/GLMdenoise/

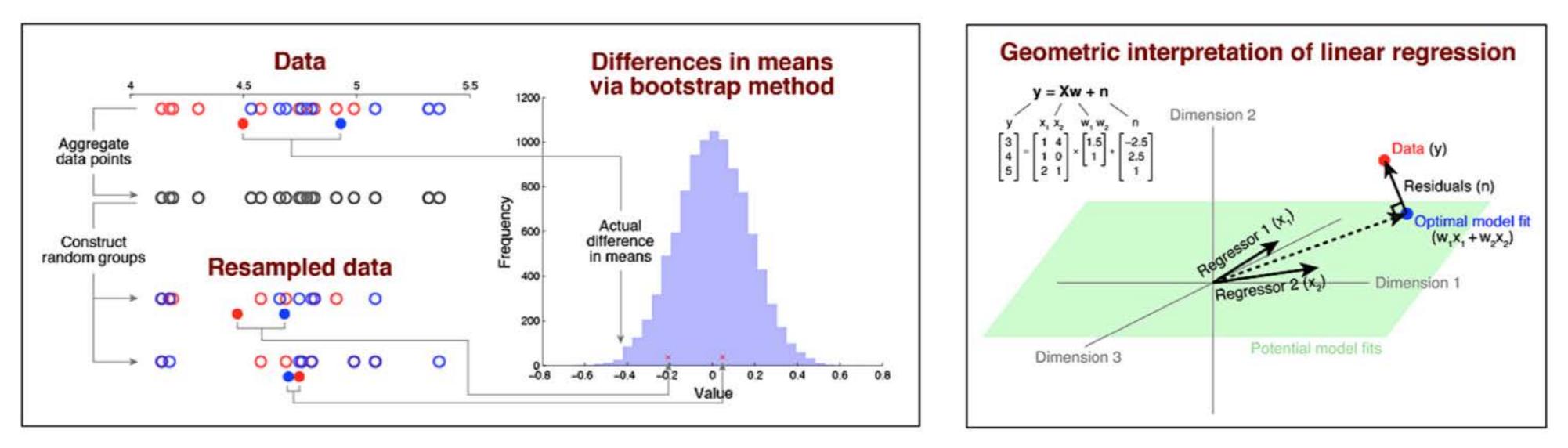


Kay et al., Frontiers in Neuroscience, 2013

Number of PCs

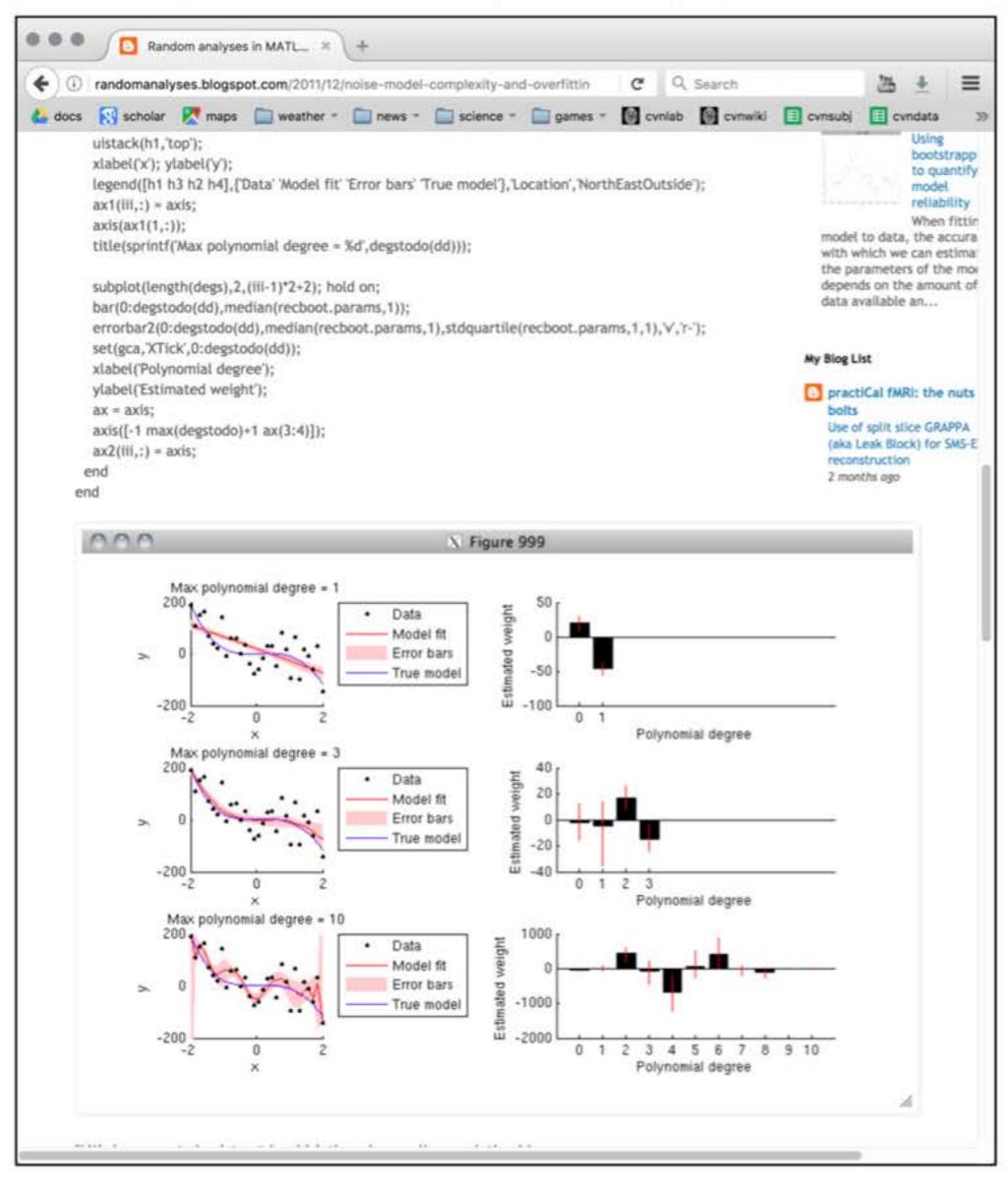
- Why statistical simulations?
  - They help teach concepts
  - They help check code correctness



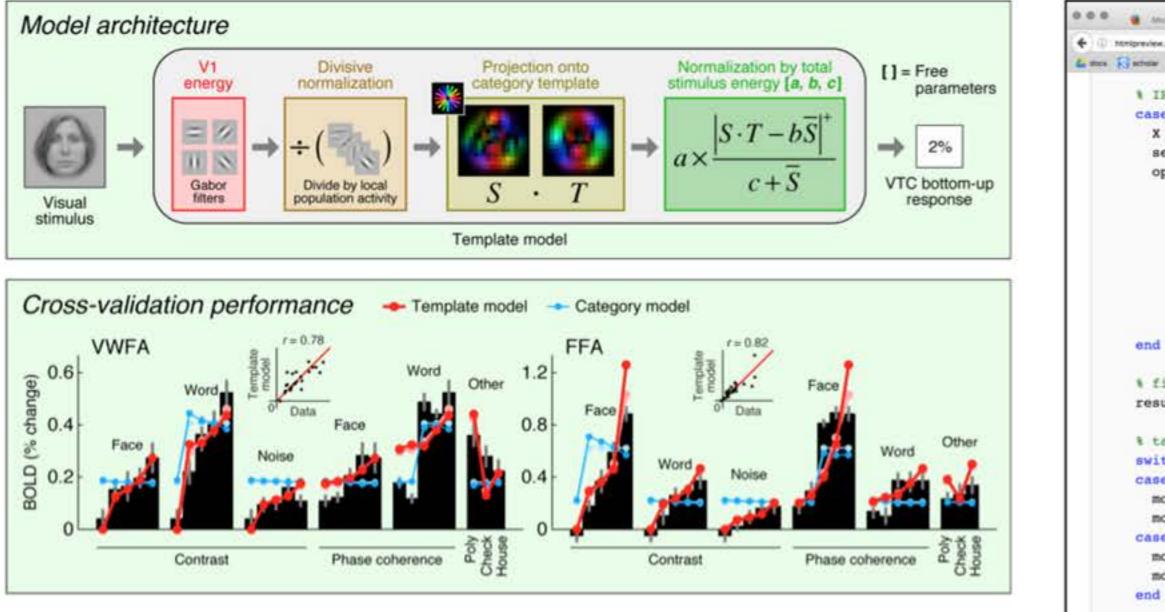


Materials at http://kendrickkay.net/psych5007/

#### http://randomanalyses.blogspot.com



#### Kay & Yeatman, eLife, 2017



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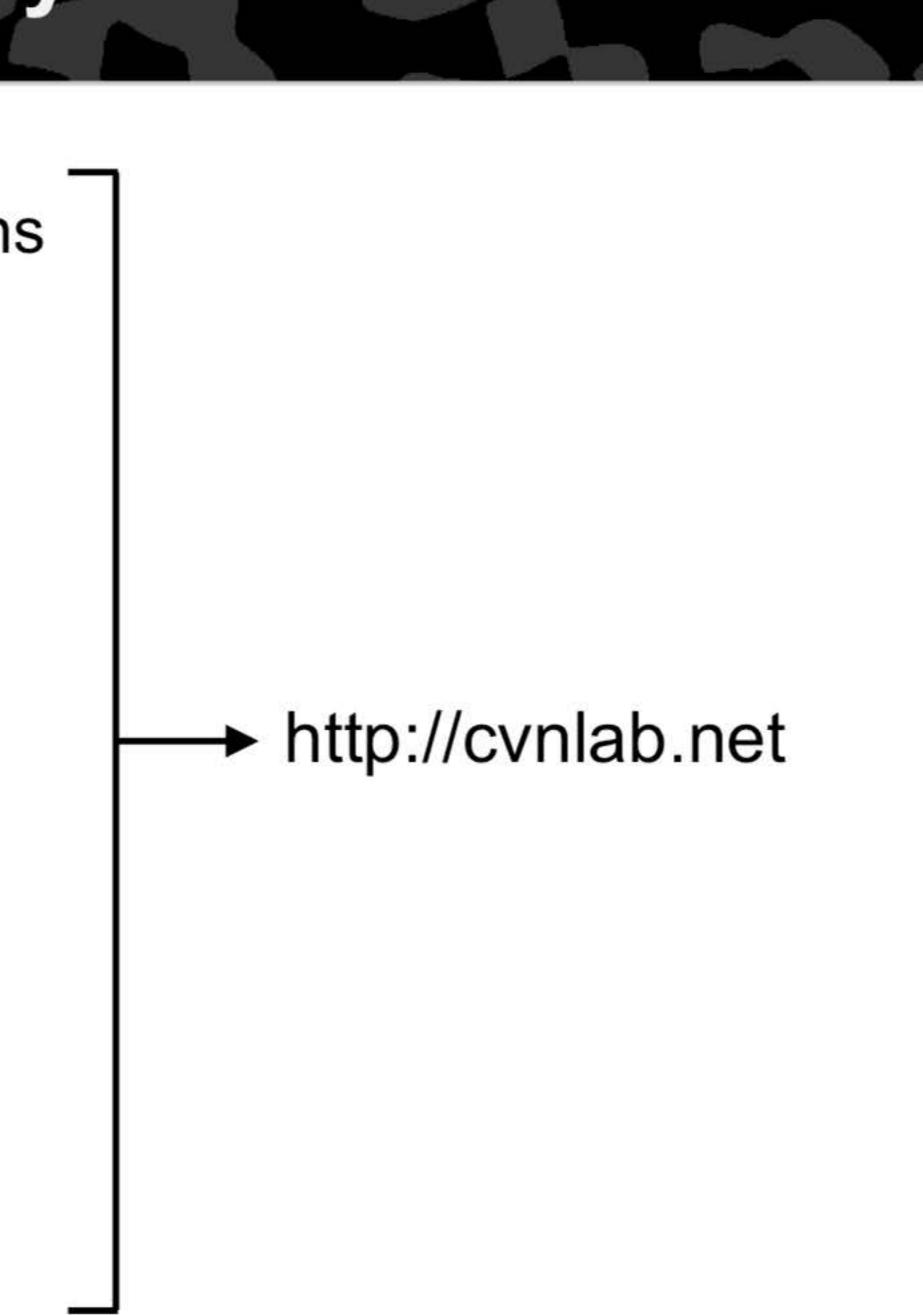
#### http://cvnlab.net/vtcipsmodel/

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        IPS-scaling model
        case 10
          X = [repmat(eye(n),[3 1]) (1:3*n)'];
          seed0 = @(ix) [data(1:n,ix)' datatopdown(:,ix)' 0 1];
          optl = struct('stimulus',X,'data', @(ix) data(:,ix),'vxs', l:size(data,2), ...
                          'model',{ ...
                                   {{[] [NaN(1,n) NaN(1,3*n) -Inf -Inf; Inf(1,n+3*n+2)] ...
                            @(p,x) x(:,1:n)*p(1:n)' .* nanreplace(p(n+3*n+1)*p(n+x(:,n+1))'+p(n+3*n+2),1)) ...
                                    (@(ss) ss [-Inf(1,n) NaN(1,3*n) -Inf -Inf; Inf(1,n+3*n+2)] ...
                     @(ss) @(p,x) x(:,1:n)*p(1:n)' .* nanreplace(p(n+3*n+1)*p(n+x(:,n+1))'+p(n+3*n+2),1)}}, ...
                          'seed', seed0, 'resampling', xvalscheme, 'metric', metricfun, ...
                          'optimoptions', {{'Display', 'off'}}, extraopt{:});
       % finally, fit the model
        results = fitnonlinearmodel(opt1);
        % take the results and store them
        switch xx
        case 1
                                    = squish(results.modelfit(1,:,:),2);
          modelfit(:,:,mm)
          modelparams (mm)
                                    = squish(results.params(1,:,:),2);
        case 2
                                    = results.modelpred;
          modelpred(:,:,mm)
          modelperformance(:,mm) = results.aggregatedtestperformance(1,:);
```

### Summary

- 1. Automated surface visualizations
- 2. Brain art
- 3. High-res fMRI pre-processing
- 4. Volume co-registration
- 5. analyzePRF
- 6. GLMdenoise

### 7. Statistics, model fitting



## Acknowledgments

Keith Jamison (UMN)

### Eshed Margalit (Stanford)



