Structural And Functional Integration: Why all imaging requires you to be a structural imager

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Structural Information is Critical for Functional (or other) research

- Functional analysis is potentially critically flawed without structural information. Structural data is necessary for:
 - visualization of functional data (view data on an easy to interpret representation of the brain)
 - spatial normalization of functional data (match anatomical locations across subjects)
 - region of interest analysis of functional data (sample data from a specific area of the brain)
 - Correction of measurement error (e.g. partial volume)
 - integrated functional analysis (e.g. volume analysis/anatomical descriptions)







What is the locus coeruleus?



Why Is Structure/Anatomy Critical?

Structure to a large degree dictates what can be done functionally:



- How big are your voxels?
- How much smoothing?
- Physiological contamination?
- How big is your effect?

Why Is Structure/Anatomy Critical?

- Accurate localization of functional results
 - Precision of localization is key to any interesting functional study
 - Don't just say 'dorsolateral prefrontal cortex'
 - Know your limitations (small structures, big voxels; ventricular borders, etc.)
- Understanding the contributions of structural changes to fMRI results in health and disease
 - Are functional changes associated with tissue degeneration within a brain structure?
 - Controlling for group biases/confounds due to structural changes
- Clinical procedures
 - Structural measurements are useful clinically, independent of functional integration (volume/lesion studies)
 - Localization of vital regions of the brain to *avoid* in neurosurgical procedures

Why is integration of anatomical and functional data difficult?

- Various levels of neuroanatomy: gyral, functional, cytoarchitectonic, neurochemical, gene expression, etc.
- Accurate models of the brain are difficult to create
- Differences in distortions/geometry across imaging domains
 - Distortion correction (acquisition/processing)
- Biological variability in anatomy across individuals

Levels of Anatomy



Modified from Devlin and Poldrack, *Neuroimage*, 2007

Commonly Used MR Sequences

- Anatomy/structure AND pathology
- T1-weighted imaging: Good contrast for gray matter/white matter; useful in segmentation of cortex and deep/subcortical gray matter
- T2/FLAIR imaging: Good contrast for segmentation of altered brain tissue such as white matter signal abnormalities (WMSA; hyperintensities; hypointense on T1, but less sensitive)
- Diffusion imaging: Good contrast for anatomy of white matter fascicles (bundles) projecting across neural regions/microstructural properties





Types of 'anatomical' information extracted from MRI data

- Segmentations (extraction of specific structures) brain, cortex, white matter, deep brain structures
- Cortical surface models
- Cortical parcellation (division of the cortex)
- White matter fascicles
- Lesions
- fMRI regions of activation









How is Structure Related to Function?



Brodmann: Primary Visual Cortex



Salat:StructFunct:HST.583:2008

How is Structure Related to Function? (Can folds predict cytoarchitecture?)

- *Ex vivo* imaging for creation of surface models
- Cytoarchitectonic borders defined with histology and mapped to surface models
- Cytoarchitecture showed good correspondence with folds, particularly in primary/secondary areas
- Some limitations of MR for defining microanatomy can be overcome by good macroanatomy



Fischl et al., *Cerebral Cortex*, 2008

How is Structure Related to Function? (Can connectivity predict function?)

- Cortical connectivity can be used to segment the nuclei of the thalamus
- Connectivity based segmentation of thalamic nuclei validated through correspondence to functionally distinct regions
- Several ways to define anatomy with imaging



Figure 1. Individual variation of connectivity-based thalamic segmentation. (*A*) Segmentation of the human thalamus. (*i*) Axial thalamic section from a cytoarchitectonic atlas (Morel *et al.*, 1997). (*ii*) In the same atlas section, major nuclei have been coloured according to their major cortical connection site (as in iii). (*iii*) Cortical subdivisions. Red = PFC; blue = PMC; orange = M1; magenta = S1/S2; green = PPC; cyan = occipital; yellow = temporal. (*iv*) Connectivity-based parcellation of the thalamus in a single subject for an axial slice at the same level as the section in *A* and *B*. Voxels are coloured according to the cortical region with which they show the highest connection probability (as in *iii*). (*B*) Connectivity-based segmentation of the thalamus in eleven subjects. Top left panel indicates location of axial thalamic slices. Each subsequent panel represents data from an individual subject. Thalamic voxels colour coded as in *Alii*. (*C*) Group probability maps. Axial images showing overlap of thalamic sub-regions across subjects in voxels showing >25% probability of connection to selected cortical mask (indicated in top left or each image) using a colour scale running from red (4/11 subjects) to yellow (11/11 subjects). Slices are taken at the average *Z*-coordinate (given in bottom right of each image) across the left and right hemispheres for the centre of gravity of that cluster.

Johansen-Berg et al., *Cerebral Cortex*, 2005

Retinotopy

- Presentation of spoke
 image to macaque
- Measure metabolism using 2DG technique on flattened visual cortex





Tootell, 1982

Slice Based Visual Organization

Spatial organization of the fMRI response to visual stimuli in occipital cortex.



Fischl et al.

Cortical Surface Model, Inflated, Flattened

 Model created of the cortical mantle, computationally manipulated for inflation, cutting and flattening

 Surface Boundaries
 Folded
 Semi-inflated
 Inflated

 Image: Surface Boundaries
 Folded
 Semi-inflated
 Inflated

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Occipital Flat Patch Retinotopy

 Different patterns presented to the visual field create different patterns of activity in occipital cortex

Flat Patch of Occipital Cortex



Tootell et al., PNAS, 1998

Variability in Cortical Anatomy





Fischl, Sereno, Dale, Neuroimage, 1999; Fischl et al., Human Brain Mapping, 1999

Various (imperfect) ways deal with anatomical variability

- Automated spatial normalization (typically whole-brain matching)
 - 'Best guess': morph structure from participant
 1 to participant 2
 - What are the anatomical features?
- Individual labeling of regions/structures
 - Hippocampus may differ in shape and size, but has clear boundaries within individuals
- Functional localizer

Spatial Normalization: Affine/linear averaging

Single subject



Fischl et al.

Average of 40

A Surface-Based Coordinate System

Fischl et al., Neuroimage, 1999; Fischl et al., Human Brain Mapping, 1999

Spherical Versus Volumetric Normalization

- Activations stronger in maps created from surface based averaging
- This demonstrates validity to the idea that function is somewhat predicted by structure (greater statistical power)
- Suggests that some limitations due to variability in anatomy can be overcome with good anatomical models/procedures

Anticevic et al., Neuroimage, 2008; See also Dasai et al., Neuroimage, 2005

Surface Smoothing

Limit smoothing to regions in close proximity on cortical surface

3D Spatial Smoothing: Combines information across gyral/sulcal boundaries Surface Smoothing: Constrains the type of information included

Spatial Smoothing

- 5 mm apart in 3D
- 25 mm apart on surface!
- Kernel much larger
- Averaging with other tissue types (WM, CSF)
 Averaging with other
- functional areas

Greve et al.

Good anatomy makes better function!

Affine registration to MNI305

5mm volume smoothing vs. 10mm surface smoothing

Greve et al.

Once you have a result, high quality atlases are important in localization

- General neuroanatomy
- Structures of Interest (hippocampus, cerebellum)
- 'Talairach' atlas commonly used: not really an atlas of neuroanatomy
- Anatomy should be confirmed for each given individual in a study (automated procedures for labeling individual anatomy exist)
- Template anatomy (using regional labels based on an atlas) can be confounded
 - Registration to the template
 - Disease associated changes

Region of Interest (ROI)

- ROI analysis is typically a secondary step
- Why ROI over maps?
 - Focused data exploration: plot data by condition for each individual/group
 - Control for statistical error by limiting measurements to *a priori* hypothesized regions
 - Limit testing to a defined region
 - Avoid circularity
 - Examine the association between the anatomical structure and function