# Qdec:

## A group-analysis tool for Freesurfer

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## What is Qdec?

- An application included in the FreeSurfer software package used to conduct a group analysis of morphometric and other types of data
- Graphical User Interface (GUI)
- Powered by a GLM-fitter
- Optimized for display of results on the cortical surface

## What problem does it solve?

- There is an ever-growing collection of subjects processable by Freesurfer's 'recon-all' stream: eg. 2000+ ADNI subjects, 400+ OASIS subjects, 1000+ FHS subjects
- Researchers need a GUI tool to allow easy exploration of the morphometric data found in this and other subject groups

## Explore data correlations

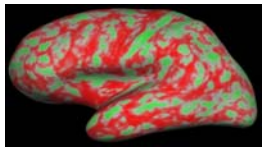
- Does subject age correlate with cortical thickness?
- Does hippocampal volume correlate with cortical thickness in a particular region?
- Is there a gender-diagnosis interaction in the thickness-age correlation?

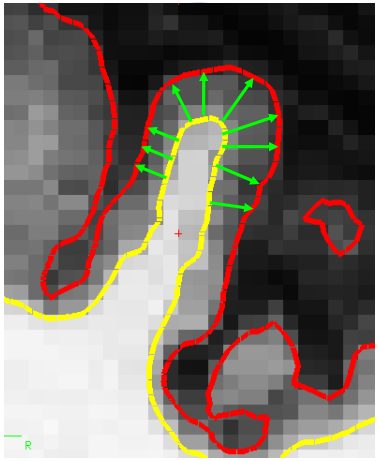
## What is FreeSurfer?

- Cortical reconstruction and subcortical segmentation of T1 MRI images (single subject)
- Morphometric data: cortical thickness, subcortical structure volume
- Surface-based registration: alignment to another subject's cortical folds

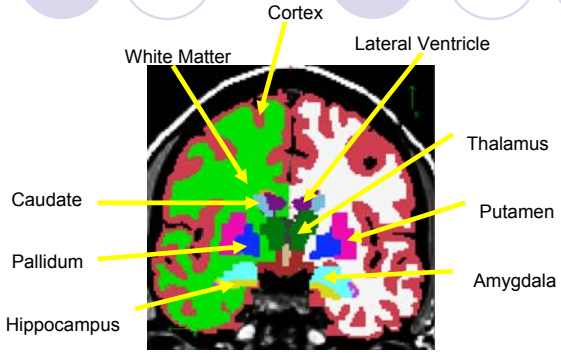
## Cortical thickness

- Distance between white and pial surfaces
- One value per vertex

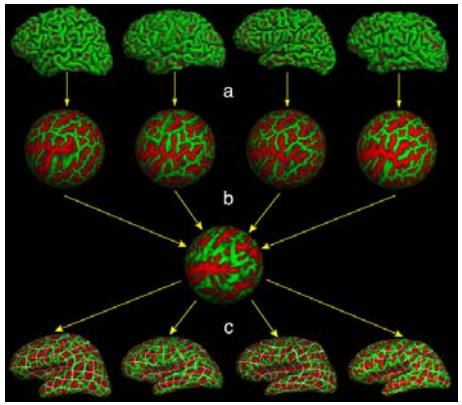




## Subcortical structures




## Surface-based registration

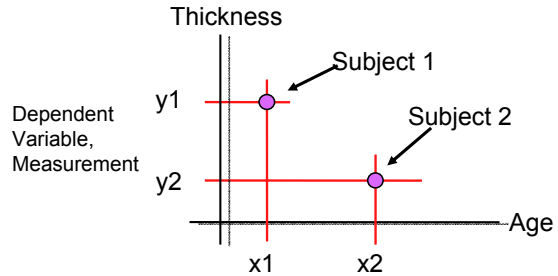


## So what is QDEC???

- Query (a subject database)
- Design (matrix, of data points, for GLM)
- Estimate (regression coefficients, GLM)
- Contrast (matrix, hypothesis)

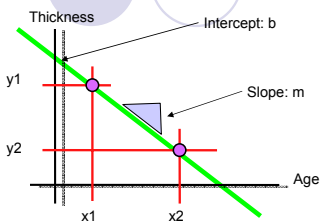
## GLM theory

Is Thickness correlated with Age?




Independent Variable

## Linear model




Intercept = Offset

X = Design Matrix
β = Regression Coefficients
= Parameter estimates
= "betas"
= Intercepts and Slopes
= beta.mgh (mri_glmfit)

System of Linear Equations
y1 = 1*b + x1*m
y2 = 1*b + x2*m

Matrix Formulation

$$\begin{bmatrix} y1 \\ y2 \end{bmatrix} = \begin{bmatrix} 1 & x1 \\ 1 & x2 \end{bmatrix} * \begin{bmatrix} b \\ m \end{bmatrix}$$

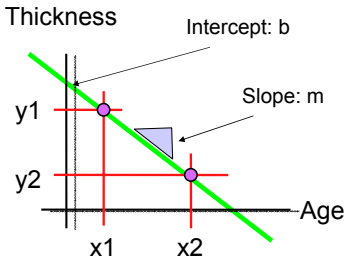
$$\boxed{Y = X*\beta} \qquad \beta = \begin{bmatrix} b \\ m \end{bmatrix}$$

## Hypotheses and contrasts

$$\begin{bmatrix} y1 \\ y2 \end{bmatrix} = \begin{bmatrix} 1 & x1 \\ 1 & x2 \end{bmatrix} * \begin{bmatrix} b \\ m \end{bmatrix}$$

Is Thickness correlated with Age?
Does m = 0?
Null Hypothesis: H0: m=0

$$m = [0\ 1] * \begin{bmatrix} b \\ m \end{bmatrix}$$




$$\beta = \begin{bmatrix} b \\ m \end{bmatrix} \qquad \boxed{\gamma = C*\beta \stackrel{?}{=} 0}$$

C=[0 1]: Contrast Matrix

## Two groups

Do groups differ in Intercept?
Does b1=b2?
Does b1-b2 = 0?
C = [+1 -1 0 0], $\gamma = C*\beta$

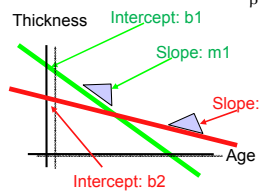
$$\begin{bmatrix} y11 \\ y12 \\ y21 \\ y22 \end{bmatrix} = \begin{bmatrix} 1 & 0 & x11 & 0 \\ 1 & 0 & x12 & 0 \\ 0 & 1 & 0 & x21 \\ 0 & 1 & 0 & x22 \end{bmatrix} * \begin{bmatrix} b1 \\ b2 \\ m1 \\ m2 \end{bmatrix}$$

Do groups differ in Slope?
Does m1=m2?
Does m1-m2=0?
C = [0 0 +1 -1], $\gamma = C*\beta$

Y = X*β

$$\beta = \begin{bmatrix} b1 \\ b2 \\ m1 \\ m2 \end{bmatrix}$$

Is average slope different than 0?
Does (m1+m2)/2 = 0?
C = [0 0 0.5 0.5], $\gamma = C*\beta$




## t-test and p-values

Y = X*β

γ = C*β

$$t = \frac{C*\beta}{\sqrt{\sigma^2 C*(X^T X)^{-1} C^T}}$$

p-value/significance
- value between 0 and 1
- closer to 0 means more significant

FreeSurfer stores p-values as –log10(p):
- $0.1=10^{-1}$→sig=1, $0.01=10^{-2}$→sig=2
- sig.mgh files
- Signed by sign of γ
- p-value is for an unsigned test

## More GLM stuff…

- For a more thorough treatment of the GLM, and FreeSurfer's implementation of it in the 'mri_glmfit' utility, see these resources:
  - Group analysis PowerPoint slides
  - Group analysis mri_glmfit tutorial

## Qdec demo

- Flash movie on next slide…
- Also see:
  - http://surfer.nmr.mgh.harvard.edu/fswiki/FsTutorial/QdecGroupAnalysis


## Future features

- Greater number of factors
- Contrast selection
- Subcortical structure volumes, and volumetric maps
- fMRI, PET, EEG/MEG surface data

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