# Analysis of longitudinal neuroimaging data with OLS & Sandwich Estimator of variance

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#### An example of longitudinal studies in neuroimaging The ADNI study

- Tensor-Based Morphometry (TBM) images from the Alzheimer's Disease Neuroimaging Initiative (ADNI) (Hua et al., 2013; Guillaume et al., 2014)
- Available scans:

	AD	MCI	Ν	Total
0 month	188	400	229	817
6 months	159	346	208	713
12 months	138	326	196	660
18 months	n/a	286	n/a	286
24 months	105	244	172	521
36 months	n/a	170	147	317

#### TBM images?



Determinant of the deformation matrix:

det(J)



det(J) > 1: expansion det(J) < 1: contraction

Results

Summary

References

# The Naive Ordinary Least Squares (N-OLS) model





- Assumes Compound Symmetry (CS):
  - Equal intra-visit variances
  - Equal intra-visit correlations
- No inference possible on between subject effects (e.g., group intercept, gender, age at first visit)

# Compound Symmetry (CS) in the ADNI dataset?

 Box's test of Compound Symmetry (Box, 1950) image thresholded at 5% after Bonferroni correction:



• 56% of the in-mask voxels survived the thresholding!!!

# The Summary Statistics OLS (SS-OLS) model

#### Procedure

- Extraction of summary statistics for each subject
  - E.g., intercept, slope

Use of an OLS model for each summary statistic

- Transformation of correlated data into uncorrelated data
- Important loss of information
  - Will affect negatively the power
  - In general, misbehaviour in unbalanced design
    - E.g., subject with 2 visits vs. subject with 6 visits

# Linear Mixed Effect (LME) models

For each subject *i*:







Fixed effects

Random effects

Random error

Pros

- The gold standard in the biostatistic literature
- Accurate if correctly specified
- Subject-specific inferences on the random effects possible

Cons

- Difficult to specify and validate
  - Only random intercepts? Also, random slopes?
  - Best model may vary across the brain
- Generally not robust against misspecification
  - E.g., random-intercept LME assumes CS like the N-OLS method
- Iterative method
  - Generally slow
  - May fail to converge

References: Bernal-Rusiel et al. (2013a,b); Chen et al. (2013); Guillaume et al. (2014)

#### Other methods could also be considered

- The "SPM procedure"
  - Assumption of a common covariance structure for the whole brain
- Generalised Methods of Moments (Skup et al., 2012)
- Generalised Estimating Equations (Li et al., 2013)
- . . .

#### The Sandwich Estimator (SwE) method

- Use of a simple OLS model (without subject indicator variables)
- The fixed effect parameters  $\beta$  are estimated by

$$\hat{\beta}_{OLS} = \left(\sum_{i=1}^{M} X_i' X_i\right)^{-1} \sum_{i=1}^{M} X_i' y_i$$

• The fixed effect parameters covariance  $var(\hat{\beta}_{OLS})$  are estimated by

$$S = \underbrace{\left(\sum_{i=1}^{M} X'_{i} X_{i}\right)^{-1}}_{\text{Bread}} \underbrace{\left(\sum_{i=1}^{M} X'_{i} \hat{V}_{i} X_{i}\right)}_{\text{Meat}} \underbrace{\left(\sum_{i=1}^{M} X'_{i} X_{i}\right)^{-1}}_{\text{Bread}}$$

#### Property of the Sandwich Estimator (SwE)

$$S = \left(\sum_{i=1}^{M} X_i' X_i\right)^{-1} \left(\sum_{i=1}^{M} X_i' \hat{V}_i X_i\right) \left(\sum_{i=1}^{M} X_i' X_i\right)^{-1}$$

If  $m^{-1} \sum_{i=1}^{m} X'_i \hat{V}_i X_i$  consistently estimates  $m^{-1} \sum_{i=1}^{m} X'_i V_i X_i$ , the SwE tends **asymptotically** (Large samples assumption) towards the true variance var( $\hat{\beta}_{OLS}$ ). (Eicker, 1963; Eicker, 1967; Huber, 1967; White, 1980)

#### The classical (uncorrected) SwE method

•  $V_i$  estimated from the residuals  $e_i = y_i - X_i \hat{\beta}$  by

$$\hat{V}_i = e_i e_i'$$

and the SwE becomes

$$S_{\text{classic}} = \left(\sum_{i=1}^{M} X_i' X_i\right)^{-1} \left(\sum_{i=1}^{M} X_i' r_i r_i' X_i\right) \left(\sum_{i=1}^{M} X_i' X_i\right)^{-1}$$

- Asymptotic test:
  - $H_0: C\hat{\beta} = 0, H_1: C\hat{\beta} \neq 0$ C: contrast matrix of rank q

$$\frac{(C\hat{\beta})'(CSC')^{-1}(C\hat{\beta})}{q} \sim \chi^2(q)$$

- Works well in large samples
- But not in small samples

### Small sample adjustment of the SwE method

- Several adjustments exists
- One of the best combination of adjustment (Guillaume et al., 2014):
  - Use of corrected residuals  $e_{ik}/(1 h_{ik})$  in the estimation of  $V_i$
  - Assumption of homogeneity across subjects within groups (e.g., same covariance structure for all the AD subjects)
  - Use of a statistical test assuming small samples  $H_0: C\hat{\beta} = 0, H_1: C\hat{\beta} \neq 0$ *C*: contrast matrix of rank *q*

C: contrast matrix of rank q

$$\frac{\nu-q+1}{\nu q}(C\hat{\beta})'(CSC')^{-1}(C\hat{\beta}) \sim F(q,\nu-q+1)$$

- Designs considered:
  - ADNI design and 4 of its subsets (817, 408, 204, 103 and 51 subjects)
- Monte Carlo Gaussian null simulation (10,000 realizations)
- For each realization,
  - Generation of longitudinal Gaussian null data (no effect) with intra-visit covariance structures:

Compound Symmetry

Toeplitz

1	1	0.95	0.95	0.95	0.95 \	1	1	0.9	0.8	0.7	0.6	)
	0.95	1	0.95	0.95	0.95	1	0.9	1	0.9	0.8	0.7	
	0.95	0.8	1	0.95	0.95		0.8	0.9	1	0.9	0.8	
	0.95	0.95	0.95	1	0.95		0.7	0.8	0.9	1	0.9	
	0.95	0.95	0.95	0.95	1 /		0.6	0.7	0.8	0.9	1	,
		-										

Statistical test (F-test at 5%) on the parameters of interest and estimation of the FPR

References

# False Positive Rate (FPR) control







Summary

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#### Real ADNI data: use of the SwE toolbox



• Freely available at http://warwick.ac.uk/tenichols/SwE

# Real ADNI data: reminder of the Box's test of CS

 Box's test of Compound Symmetry (Box, 1950) image thresholded at 5% after Bonferroni correction:



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Introduction

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#### Real ADNI data: Visit effect on the brain atrophy



Introduction	The Sandwich Estimator method	Results	Summary	Reference
Summary				

- Longitudinal standard methods not really appropriate to neuroimaging data:
  - N-OLS & LME with random intercepts: issues when CS does not hold
  - Difficulties to specify and validate LME models
  - Convergence issues with LME models
  - Under unbalanced design, SS-OLS may be inaccurate and its power quite poor
- The SwE method
  - Accurate in a large range of settings
  - Easy to specify
  - No iteration needed
    - Quite fast
    - No convergence issues
  - Can accommodate pure between covariates
  - SPM toolbox available
  - But, careful in small samples:
    - Adjustments essential
    - Typically, less powerful than N-OLS or LME models

Results

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Thanks for your attention!