

No Limits: AN(C)OVA for Group Analysis with a Large Number of Factors and Covariates

Gang Chen¹, Ziad Saad¹, Johanna Jarcho², Daniel Pine², Robert Cox¹



¹Scientific and Statistical Computing Core

²Section on Development and Affective Neuroscience

NIMH & NINDS / NIH / DHHS / Bethesda MD USA

Contact: gangchen@mail.nih.gov <http://afni.nimh.nih.gov>



Traditional FMRI Group Analysis Approaches

General Linear Model (GLM)

- Take effect estimates (β 's) only from individual subject analyses
- Assume that effect estimates have same precision across subjects
- Models in widespread use
 - Student t-tests: one-, two-sample, and paired
 - Simple AN(C)OVA
- Programs in AFNI: 3dttest++, 3dANOVA2/3, GroupAna
- Limitations
 - Limited number of explanatory variables (factors and covariates)
 - Rigid data structure: no missing data allowed
 - Rigid model: all possible effects included even if not significant
 - Rigid assumptions: sphericity; generic covariance structure for random effects and residuals unavailable

Linear Mixed-Effects (LME) Model

- Take effect estimates (β 's) only from individual subject analyses
- Assume effect estimates have same precision across subjects
- Flexibility of specifying covariance structures and heterogeneity
- No limit on number of explanatory variables
- Missing data allowed
- Program in AFNI: 3dLME
- Limitations: difficult in assigning degrees of freedom

Mixed-Effects Multi-level Analysis (MEMA)

- Consider precision information from individual level
- More accurate group effect estimate and significance test
- Program in AFNI: 3dMEMA
- Limitations: only able to analyze paired, one-, and two-sample types

Multivariate Modeling (MVM) Paradigm

Repeated Measures (RM) Designs

- Within-subject or RM factor
 - Categorical variable: e.g., 3 levels of emotion (positive, negative, neutral)
 - Effect estimates available at all levels from every subject
 - Traditionally modeled as an explanatory variable (factor)
 - One-way within-subject or RM ANOVA
- RM designs are popular in psychology and FMRI experiments
- Typically formulated as ANOVA: e.g., one-way within-subject or RM ANOVA

$$y_{ij} = \mu_j + b_i + \epsilon_{ij}$$

where y_{ij} is the response at the j th factor level for the i th subject, μ_j is the effect at the j th factor level, b_i is the deviation of the i th subject at the j th factor level, and ϵ_{ij} is the residual of the i th subject at the j th factor level, $i = 1, \dots, n$, $j = 1, \dots, m$. The distributional assumptions are $b_i \sim N(0, \tau^2)$ and $\epsilon_{ij} \sim N(0, \sigma^2)$, where τ^2 and σ^2 are between- and within-subject variances respectively.

- Efficient when sphericity holds, but sensitive to sphericity violation: inflated significance
- May involve between-subjects (or subject-grouping) factors

Multivariate modeling (MVM) approach

- All levels (or their contrasts) of RM factor: multiple response variables
- One-way RM ANOVA transforms to a special MANOVA

$$\mathbf{y}_i = \mathbf{u} + \mathbf{e}_i$$

where $\mathbf{y}_i = (y_{i1}, \dots, y_{im})^T$, $\mathbf{u} = (\mu_{i1}, \dots, \mu_{im})^T$, and $\mathbf{e}_i = (\epsilon_{i1}, \dots, \epsilon_{im})^T$, $i = 1, \dots, n$. The distributional assumption is $\mathbf{e}_i \sim N_m(0, \Sigma)$, where Σ is a positive definite and symmetric matrix that is constant across subjects, and \mathbf{e}_i 's are independent from each other. Unlike ANOVA, the covariance matrix Σ is estimated from the data instead of being assumed spherical.

- Traditional AN(C)OVA becomes MAN(C)OVA: only between-subjects factors and quantitative variables are treated as explanatory variables
- Immune to sphericity violation, but may lose power when sphericity holds

Implementation in AFNI: 3dMVM

- Program written in R [1] with package afex [2] for MVM
- Post-hoc tests performed through symbolic coding with labels in R package phia [3]
- Currently implemented as shell scripting with parallel computing capability through package snow [4]

Assessment of MVM Approach

Advantages of MVM: Flexibility in Modeling

- Multivariate aspect: when BOLD response shape is modeled with multiple basis functions, the shape integrity can be maintained at group level using MVM without presuming the covariance structure across the multiple effects
 - Alternative approach: LME with AR covariance matrix for residuals – 3dLME
- Univariate aspect: MVM platform allows for GLM capability
 - Huynh-Feldt and Greenhouse-Geisser corrections available when sphericity is violated
- Allows for unbalanced designs (unequal subjects across groups)
- Explanatory variables can be categorical or quantitative
- No limit on the number of explanatory variables (factors and covariates)
- Post-hoc tests through symbolic coding with labels instead of direct dummy coding

Disadvantages of MVM

- Under some circumstances bigger sample size (subjects) required
- Computationally heavy
- Unable to deal with missing data
 - Solution: 3dLME
- Incapable of handling within-subject quantitative variables
 - Solution: 3dLME

Sample Application

- 6 categorical variables (factors) in FMRI experiment: 6-way ANOVA
 - 3 between-subjects factors: age (adult, child), diagnosis (healthy, anxious), and scanner (scanners 1 and 2)
 - 3 within-subject factors: subject's interest in a target (interest, no interest), target's interest in a subject (interest, no interest), accuracy (accurate, inaccurate)
- 89 subjects: healthy children (24) and adults (32), anxious children (15) and adults (15)
 - Unbalanced design: unequal number of subjects across groups
- 259 post-hoc tests in addition to F-stat for main effects and interactions

Acknowledgements

The research was supported by the NIMH Intramural Research Program of the NIH.

References

- [1] R Core Team (2013). R: A language and environment for statistical computing. R Foundation for Statistical Computing, Vienna, Austria. ISBN 3-900051-07-0, URL <http://www.R-project.org/>.
- [2] Henrik Singmann (2013). afex: Analysis of Factorial Experiments. R package version 0.4-57. <http://CRAN.R-project.org/package=afex>
- [3] Helios De Rosario-Martinez (2012). phia: Post-Hoc Interaction Analysis. R package version 0.1-0. <http://CRAN.R-project.org/package=phia>
- [4] Luke Tierney, A. J. Rossini, Na Li and H. Sevcikova (2013). Snow: Simple Network of Workstations. R package version 0.3-12. <http://CRAN.R-project.org/package=snow>