Biostat 743 Lecture 10/12/17

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October 15, 2017

Outline

- Clustered Data
 - Notation
 - Example
- Marginal Models
- Generalized Estimating Equations (GEE)
 - Sandwich Estimator
- Generalized Linear Mixed Models

Clustered Data

- Set of observations in correlated sets (or clusters), e.g., individuals in a family, geographical unit, hierarchical data, teeth in mouth, repeated measures on units (longitudinal data)
- Why do we need to think of this data differently than that of independent data?
 - Observations in clusters breaks our assumption of independence
 - Clustered data gives less information than independent data

Notation

 $Y_{ij} = \text{observations/outcome for unit (individual)} i \text{ at observation } j$

$$j = 1, 2, ..., n_i$$

 $i = 1, 2, ..., n$
 $N = \sum n_i$

Example

 Y_{ij} is the number of cases of some disease X in state i and district j.

Treatment
$$Trt_{ij} = \begin{cases} 1, & \text{if } Trt \text{ is in i, j} \\ 0, & \text{otherwise} \end{cases}$$

$$Y_{ij} \sim Poisson(\lambda_{ij}, P_i)$$

$$log(\lambda_{ij}) = \beta_0 + \beta_1 Trt_{ij} + \beta_2 X$$

- X is a possible confounder; e.g., past incidence or socioeconomic status
 - We want to make inference about β_1 . What is the treatment effect?

- Need to adjust for clustering because Y_{ij} is not independent from $Y_{ij'}$
- One approach to adjusting for clustering: marginal models

Marginal Models

- Directly model $E(Y_{ij}|X_{ij})$
- Effects can be interpreted as population level average effects

Ingredients for Marginal Models

1. Model expected response linked to covarates using a GLM

$$E(Y_{ij}|X_{ij}) = \mu_{ij}$$
$$g(\mu_{ij}) = \eta(X_{ij}) = X_{ij}\beta$$

2. Marginal variance is a function of the marginal mean

$$Var(Y_{ij}|X_{ij}) = \phi v(\mu_{ij})$$

- where ϕ is a scale parameter and $v(\mu_{ij})$ a known function.
- e.g.
- (a) Gaussian/continuous outcome: $v(\mu_{ij}) = 1$, $\phi = \sigma^2$
- (b) Poisson/binomial GLM: $v(\mu_{ij}) = \mu_{ij}$ (poisson), $\phi = 1$ ($\phi > 1$ in overdispersed model)
- 3. Specify within unit correlation as a function of μ_{ij} and maybe other parameters
 - e.g., $Cov(Y_{ij}, Y_{ij'}) = sd(Y_{i\ j}) * Corr(Y_{i\ j}, Y_{ij'}) * sd(Y_{ij'})$ where $Corr(Y_{ij}, Y_{ij'})$ is parameterized.

Examples of complete specification of marginal model

• Normal outcome:

1.
$$\mu_{ij} = X_{ij}\beta$$

2. $v(\mu_{ij}) = \phi$
3. $Corr(Y_{ij}, Y_{ij'}) = \alpha_{jj'}$

- Unstructured correlation and $\alpha_{jj'}$ is different for every pair of (j, j').
- Covariance structure of data
- For n_i , i = 1, 2, 3, ...,

$$V = \phi \begin{pmatrix} R_{n_1} & 0 & 0 & \dots \\ 0 & R_{n_2} & 0 & \dots \\ 0 & 0 & R_{n_3} & \dots \\ \vdots & \vdots & \vdots & \ddots \end{pmatrix} \text{ where, say... } R = \begin{pmatrix} 1 & \alpha_{12} & \alpha_{13} \\ \alpha_{21} & 1 & \alpha_{23} \\ \alpha_{31} & \alpha_{32} & 1 \end{pmatrix}$$
 or, say...
$$R = \begin{pmatrix} 1 & \alpha & \alpha \\ \alpha & 1 & \alpha \\ \alpha & \alpha & 1 \end{pmatrix} \text{ (compound symmetry)}$$

or, say... R = exponential correlation structure, etc...

or, say... R = autoregressive correlation structure, etc...

or, say... R = unstructured correlation structure, etc...

or, say... R = Toeplitz correlation structure, etc...

etc...

etc...

Poisson/count outcome:

1.
$$ln(\mu_{ij}) = X_{ij}\beta$$

2.
$$v(\mu_{ij}) = \phi \ \mu_{ij}$$

overdispersed model if $\phi > 1$

3.
$$Corr(Y_{ij}, Y_{ij'}) = \alpha$$

Generalized Estimating Equations (GEE)

- Quasilikelihood approach
- Marginal models can be fit using GEE

$$\mu(\beta) = \sum_{i=1}^{N} \left(\frac{\delta \mu_i}{\delta \beta} \right)^T V_i^{-1}(y_i - \mu_i) = 0, \text{ where } \mu_i = g^{-1}(X_i \beta)$$

• In practice, V_i^{-1} is a scalar value; working covariance matrix - means approximate, i.e., know not exact truth

Key Points about GEE

- 1. β estimates are equivalent to ML estimates from GLM's when data are from an exponential family
- **2.** Even though this is quasilikelihood, there still are nice asymptotic results: $\hat{\beta} \to \beta$ even if variance is unspecified, estimate $\hat{\beta}$ ignoring correlation structure and instead use...
- 3. Sandwich Estimator for variance to get our SE's to account for possible misspecification of variance
 - Model based covariance of estimator:

$$V_{model} = \left[\sum_{i=1}^{n} \left(\frac{\delta \mu_i}{\delta \beta} \right)^T V(\mu_i)^{-1} \left(\frac{\delta \mu_i}{\delta \beta} \right) \right]^{-1}$$

• in practice, we use the sandwich estimator:

$$V_{sandwich} = V_{model} * \left[\sum_{i=1}^{n} \left(\frac{\delta \mu_i}{\delta \beta} \right)^T * [v(\mu_i)]^{-1} * var(Y_i) * [v(\mu_i)]^{-1} \left(\frac{\delta \mu_i}{\delta \beta} \right) \right] * V_{model}$$

- $var(Y_i)$ is the data driven estimate of variance
- **4.** Interpretations of $\beta's$ are for the population as a whole.
 - GEE setting: $g(E[Y_{ij}|X_{ij}]) = X_{ij}\beta$ where g(.) is the link
 - Correlations are considered "nuisance" parameters
 - Allows us to "fix" SE's post-hoc

Generalized Linear Mixed Models (GLMM's)

- Just a different way of accounting for correlation structure
- Model the correlations directly as part of our mean model

$$g\left(E[Y_{ij}|X_{ij},b_i]\right)=X_{ij}\beta+Z_{ij}b_i$$
 ,
where $X_{ij}\beta$ are fixed effects and $Z_{ij}b_i$ are random effects

- Simple case: $Z_{ij}=1 \implies Z_{ij}b_i=\alpha_i$, where the α_i are a set of random intercepts, $\alpha_i \sim N(0,\tau^2)$
- In a marginal model, when we don't have random effects, the $\beta's$ refer to overall population level effect, but when we use random intercepts, we have unit/subject specific interpretation or treatment effect e^{β} .