

Population models in design and analysis of dose finding experiments for binary responses

V. Fedorov, GSK

June, 2009

Outline

- Two types of mixing models
- Moment based approach
- Iterated estimators
- Design

Two mixing models

Y is the binary response and

$$\pi_{ij} = P(Y = 1 \mid i, j)$$

j stands treatment (dose), i – experimental unit (patient, center).

Two types of “mixing”

Type A

$$\pi_{ij} = \eta(x_j, \gamma_i), \quad \gamma_i \sim \phi(\gamma \mid \theta)$$

Type B

$$\pi_{ij} \sim \psi(\pi, \gamma(x_j, \theta))$$

Type A: G-probit model

$$\pi_{ij} = \pi(x_j, \gamma_i) = \int_{-\infty}^{\eta(x_j, \gamma_i)} \psi(u|\alpha) du, \quad \gamma_i \sim \varphi(\gamma|\theta)$$

- ⊕ Probabilities π_{ij} are dependent for different j-s

Likelihood function:

$$L(\theta|\mathbf{Y}) = \prod_{i=1}^N \mathbb{E} \left[\prod_{j=1}^K \pi_{ij}^{Y_{ij}} (1 - \pi_{ij})^{n_{ij} - Y_{ij}} \mid \theta \right]_{\pi}$$

- ⊖ Easy to write difficult to compute!
- ⊖ Problem with building information matrix.

Type B. Beta-binomial model

$$\pi_{ij} \sim \text{Beta}(a, b)$$

$$a_{ij} = \exp[\boldsymbol{\theta}_a^T f_a(x_{ij})], \quad b_{ij} = \exp[\boldsymbol{\theta}_b^T f_b(x_{ij})]$$

Marginal distribution of Y_{ij} .

$$\psi(y; n_{ij}, a, b) = \binom{n_{ij}}{y} \frac{B(y + a_{ij}, n_{ij} - y + b_{ij})}{B(a, b)}$$

$$L(\boldsymbol{\theta} | \mathbf{Y}) = \prod_{i=1}^N \prod_{j=1}^K \binom{n_{ij}}{Y_{ij}} \frac{B(Y_{ij} + a_{ij}, n_{ij} - Y_{ij} + b_{ij})}{B(a_{ij}, b_{ij})}$$

⊕ Easy to compute MLE and information* matrices

⊖ No dependence between results on different doses

$$E[Y_{ij}] = n_{ij} \frac{a}{a+b} = n_{ij} p$$

$$\text{Var}[Y_{ij}] = n_{ij} \frac{ab(a+b+n_{ij})}{(a+b)^2(a+b+1)}$$

$$= n_{ij} p(1-p) \left\{ 1 + \frac{\tau}{1+\tau} (n_{ij} - 1) \right\}$$

*Sudhir R., etc. Biometrical Journal 47 (2005) 2, 230–236

Quasi-linear or moment based methods I

Very modest assumptions:

$$\mathbf{E}[\boldsymbol{\pi}_i] \equiv \mathbf{p} \quad \text{and} \quad \text{Var}[\boldsymbol{\pi}_i] = \mathbf{V}$$

Estimators for elemental parameters:

$$\hat{\pi}_{ij} = \bar{Y}_{ij} = \frac{Y_{ij\cdot}}{n_{ij}}$$

Their variance*: $\text{Var}[\hat{\boldsymbol{\pi}}_i] = \mathbf{V}_i = \mathbf{M}_i^{-1} + \mathbf{V}$

$$\hat{\boldsymbol{\pi}}_i = \{\hat{\pi}_{ij}\}_1^K \quad \text{and} \quad M_{i,jj'} = \delta_{jj'} \frac{n_{ij}}{p_j(1-p_j) - V_{jj}}$$

We are interested in estimation of \mathbf{p} and \mathbf{V} .



Simple computing, easy to introduce correlation between results at different doses



Potential loss of information

Quasi-linear or moment based methods II

Previous formulae are based on the formula for marginal variance:

$$\text{Var}[\hat{\pi}_i] = E [\text{Var} [\hat{\pi}_i | \pi_i]] + \text{Var} [E [\hat{\pi}_i | \pi_i]]$$

Note.

For any continuous p.d.f. $f(p)$ with the support set $[0, 1]$, mean μ_p and variance σ_p^2 , we have $\sigma_p^2 < \mu_p(1 - \mu_p)$, with equality only for the atomized distribution at points 0 and 1 with weights $1 - \mu_p$ and μ_p , respectively. Hence, $p_j(1 - p_j) - V_{jj} > 0$.

Iterated quasi-linear estimator I

Mimicking the best linear estimator leads to:

$$\hat{\mathbf{p}} = \left(\sum_{i=1}^N \mathbf{W}_i \right)^{-1} \sum_{i=1}^N \mathbf{W}_i \hat{\pi}_i$$

where $\mathbf{W}_i = \mathbf{V}_i^{-1}$, when \mathbf{M}_i is regular and $\mathbf{W}_i = \mathbf{M}_i - \mathbf{M}_i(\mathbf{M}_i + \mathbf{V}^{-1})^{-1}\mathbf{M}_i$ otherwise.

$$\text{Var}[\hat{\mathbf{p}}] = \left(\sum_{i=1}^N \mathbf{W}_i \right)^{-1} = \left(\sum_{i=1}^N \mathbf{V}_i^{-1} \right)^{-1}$$

⊖ Problem: \mathbf{p} and \mathbf{V} are unknown ☹

Iterated quasi-linear estimator II

$$\hat{\mathbf{p}}^{(t+1)} = \left(\sum_{i=1}^N \mathbf{W}_i^{(t)} \right)^{-1} \sum_{i=1}^N \mathbf{W}_i^{(t)} \hat{\boldsymbol{\pi}}_i$$

Where:

$$\mathbf{W}_i^{(t)} = \left[\left(\mathbf{M}_i^{(t)} \right)^{-1} + \mathbf{V}^{(t)} \right]^{-1}$$

$$\left(\mathbf{M}_i^{(t)} \right)^{-1} = \begin{pmatrix} \frac{p_1^{(t)}(1-p_1^{(t)})-V_{11}^{(t)}}{n_{i1}} & 0 & \dots \\ 0 & \frac{p_2^{(t)}(1-p_2^{(t)})-V_{22}^{(t)}}{n_{i2}} & \dots \\ \dots & \dots & \dots \end{pmatrix}$$

$$\mathbf{V}^{(t)} = \frac{1}{N} \sum_{i=1}^N \begin{pmatrix} \frac{Y_{i1} \cdot Y_{i1} - 1}{n_{i1} \cdot n_{i1} - 1} & \frac{Y_{i1} \cdot Y_{i2}}{n_{i1} \cdot n_{i2}} & \dots \\ \frac{Y_{i1} \cdot Y_{i2}}{n_{i1} \cdot n_{i2}} & \frac{Y_{i2} \cdot Y_{i2} - 1}{n_{i2} \cdot n_{i2} - 1} & \dots \\ \dots & \dots & \dots \end{pmatrix} - \hat{\mathbf{p}}^{(t)} \hat{\mathbf{p}}^{(t)T}$$

Motivation for the estimator selection

The iterative estimator for \mathbf{V} is motivated by the fact that

$$E \left(\frac{Y_{ij} \cdot Y_{ij} \cdot - 1}{n_{ij} n_{ij} - 1} \mid \pi_{ij} \right) = \pi_{ij}^2,$$
$$E \left(\frac{Y_{ij} \cdot Y_{ij'} \cdot}{n_{i1} n_{i2}} \mid \pi_{ij}, \pi_{ij'} \right) = \pi_{ij} \pi_{ij'},$$

while

$$E(\pi_{ij} \pi_{ij'}) = V_{jj'} + p_j p_{j'},$$
$$E(\pi_{ij}^2) = V_{jj} - p_j^2.$$

Locally optimal designs

Optimality criterion:

$$\text{Var}[\ell^T \hat{\mathbf{p}}] = \ell^T \left(\sum_{i=1}^N \mathbf{V}_i^{-1} \right)^{-1} \ell$$

Two controls:

1. number of patients in each center
2. distribution of patients between treatments

$$\text{Var}[\ell^T \hat{\mathbf{p}}] = \ell^T \left(\sum_{i=1}^N \mathbf{V}_i^{-1} \right)^{-1} \ell \geq \ell^T \left(\sum_{i=1}^N (\mathbf{M}^{-1} + \mathbf{V})^{-1} \right)^{-1} \ell = \frac{1}{N} \ell^T (\mathbf{M}^{-1} + \mathbf{V}) \ell$$

Thus optimal design should be balanced across all centers, i.e. $n_{ij} = n_j$ and straightforward optimization leads to

$$n_{ij} \equiv n_j \sim \sqrt{\ell_j^2 [p_j(1-p_j) - V_{jj}]}$$

How to select population means and variances I

Response models with random parameters

Link function

$$\lambda(\pi_{ij}) = \gamma_i^T \mathbf{f}(x_j) + \varepsilon_{ij}$$

$$\gamma_i \sim \mathcal{N}(\gamma, \Sigma), \quad \varepsilon_{ij} \sim \mathcal{N}(0, \sigma^2)$$

Population means:
$$p_j \simeq \lambda^\ominus \left[\gamma^T \mathbf{f}(x_j) - \frac{1}{2} \ddot{\lambda}(\gamma^T \mathbf{f}(x_j)) V_{jj} \right]$$

Var-Cov matrix of \mathbf{p} :
$$\mathbf{V} = \text{Var}(\boldsymbol{\pi}) \simeq \dot{\Lambda}^{-1} (\mathbf{F}^T \boldsymbol{\Sigma} \mathbf{F} + \sigma^2 \mathbf{I}) \dot{\Lambda}^{-1}$$

Where:
$$\mathbf{F} = \{\mathbf{f}(x_j)\}_1^K, \quad \dot{\Lambda} = \{\text{diag} \dot{\lambda}(\gamma_i^T \mathbf{f}(x_j))\}$$

How to select population means and variances II

Derivations are based on Taylor's expansion:

$$\lambda(\pi_{ij}) = \gamma_i^T \mathbf{f}(x_j) + \varepsilon_{ij} \simeq \lambda(p_j) + \dot{\lambda}(p_j)(\pi_{ij} - p_j) + \frac{1}{2}\ddot{\lambda}(p_j)(\pi_{ij} - p_j)^2$$

For logit link: $\lambda(p) = \ln \frac{p}{1-p} = u, \quad p(u) = \lambda^\ominus(u) = \frac{e^u}{1+e^u}$

Response described by autoregression model:

$$\lambda(\pi_{ij}) = [\gamma_i(x_j - x_{j-1})]\lambda(\pi_{i,j-1}) + \varepsilon_{ij}$$

References

- Agresti, A. and Hartzel, J. (2000). Strategies for comparing treatments on a binary response with multi-centre data. *Statistics in Medicine*, **19**: 1115–1139.
- Dragalin V., Fedorov V. (2006) Design of multi-centre trials with binary response. *Statistics in Medicine*, **25**:2701–2719.
- Sudhir P., Uditha B., and Tathagata B. (2005). Fisher Information Matrix of the Dirichlet-multinomial Distribution *Biometrical Journal*, **47**, 230–236.