TECHNICAL REPORT

Method-of-moments computations for meta-analysis using summary (trial-level) data when applied to binary surrogate and true endpoints

Supplement to: A simple meta-analytic approach for using a binary surrogate endpoint to predict the effect of intervention on true endpoint *Biostatistics* 2005

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We adapted the trial-level meta-analysis of surrogate endpoints (Gail et al, 2000, Buyse et al, 2000) to binary surrogate and true endpoints using a relatively simple method-of-moments approach for estimation. Let Z denote randomization group, S denote a surrogate endpoint with realizations of 0 or 1, and T denote a true endpoint with realizations 0 or 1. Also let I=0 denote the application or validation trial and I=i = 1, 2, k index previous trials. We define

 $\phi_{0zT}=pr(T=1|Z=z,I=0)=$ the probability the true endpoint equals 1 in arm z of the application or validation trial,

 $\phi_{0zS} = pr(S=1|Z=z, I=i) = \text{ the probability the surrogate endpoints equals 1}$ in arm z of the application or validation trial,

 $\phi_{izS} = pr(S=1|Z=z, I=i)$, = the probability the true endpoint equals 1 in arm z of previous trial i,

 $\phi_{izT}=pr(T=1|Z=z,\,I=i)=$ the probability the true endpoint equals 1 in arm z of previous trial i,

The parameter of interest is the treatment effect in the new trial $\Delta_0 = \phi_{01T} - \phi_{00T}$. We construct a joint random effects model for the estimate of ϕ_{izS} and ϕ_{izT} based on data from previous trials $i=1,\,2,\,k$ and use the model along with the estimates in new trial of the probability of surrogate endpoint to estimate the treatment effect Δ_0 in the new trial. Let n_{izst} denote the number of subjects in group z of trial i who have surrogate outcome s and true outcome t. The estimates under a binomial model are $\widehat{\phi}_{izS} = n_{iz1+}/n_{iz++}$ and $\widehat{\phi}_{izT} = n_{iz+1}/n_{iz++}$. Let $\phi_i = (\Delta_i, \phi_{i0S}, \phi_{i1S})^T$, where $\Delta_i = \phi_{i1T} - \phi_{i0T}$. We assume that $\widehat{\phi}_i \sim N(\phi_i, V_{\text{sampling}(i)})$. Let $\theta_{izst} = pr(S=s, T=t|i,z)$. Applying the delta method

$$V_{\text{sampling}(i)} = \begin{pmatrix} \sigma_{i\Delta\Delta}^2 & \sigma_{i\Delta S0}^2 & \sigma_{i\Delta S1}^2 \\ \sigma_{i\Delta S0}^2 & \sigma_{iS0S0}^2 & 0 \\ \sigma_{i\Delta S1}^2 & 0 & \sigma_{iS1S1}^2 \end{pmatrix} = A \cdot W_i \cdot A^T$$
, where

$$W_{i} = \begin{pmatrix} w_{i0SS} & w_{i0ST} & 0 & 0 \\ w_{i0ST} & w_{i0TT} & 0 & 0 \\ 0 & 0 & w_{i1SS} & w_{i1ST} \\ 0 & 0 & w_{i1ST} & w_{i1TT} \end{pmatrix}, A = \begin{pmatrix} 0 & -1 & 0 & 1 \\ 1 & 0 & 0 & 0 \\ 0 & 0 & 1 & 0 \end{pmatrix},$$
(1)

$$\begin{split} w_{izSS} &= \phi_{izS} (1 - \phi_{izS}) \, / \, n_{iz++}, \\ w_{izTT} &= \phi_{izT} (1 - \phi_{izT}) \, / \, n_{iz++}, \\ w_{izST} &= cov(\phi_{izS}, \phi_{izT}) = cov(\theta_{iz10} + \theta_{iz11}, \theta_{iz01} + \theta_{iz11}) \\ &= (1 \quad 0 \quad 1) \begin{pmatrix} \theta_{iz01} (1 - \theta_{iz01}) & - \theta_{iz01} \theta_{iz10} & - \theta_{iz01} \theta_{iz11} \\ - \theta_{iz01} \theta_{iz10} & \theta_{iz10} (1 - \theta_{iz10}) & - \theta_{iz10} \theta_{iz11} \\ - \theta_{iz01} \theta_{iz11} & - \theta_{iz10} \theta_{iz11} & \theta_{iz11} (1 - \theta_{iz11}) \end{pmatrix} / n_{iz++} \begin{pmatrix} 0 \\ 1 \\ 1 \end{pmatrix} \\ &= \left\{ \theta_{iz11} \, \left(1 - \phi_{izS} \right) - \theta_{iz01} \, \phi_{izS} \right\} / n_{iz++}, \end{split}$$

For estimation, $\widehat{\theta}_{iz11} = n_{iz11}/n_{iz++}$ and $\widehat{\theta}_{iz01} = n_{iz01}/n_{iz++}$.

Following the general approach of Gail et al (2000) and Buyse et al (2000), to allow for extra variability, we superimpose a random effects models, $\phi_i \sim N(\phi, V_{\rm random})$, where

$$V_{\text{random}} = \begin{pmatrix} d_{\Delta\Delta} & d_{\Delta S0} & d_{\Delta S1} \\ d_{\Delta S0} & d_{S0S0} & d_{S0S1} \\ d_{\Delta S1} & d_{S0S1} & d_{S1S1} \end{pmatrix}.$$
(2)

Let $\omega_0 = (\Delta_0, \widehat{\phi}_{00S}, \widehat{\phi}_{01S})$, where $\Delta_0 = \phi_{01T} - \phi_{00T}$ is the parameter of interest. Based on (B1) and (B2), we assume a joint normal distribution for ω_0 with mean $(E(\Delta_0), E(\phi_{00S}), E(\phi_{01S}))$ and variance

$$\begin{pmatrix} d_{\Delta\Delta} & d_{\Delta S0} & d_{\Delta S1} \\ d_{\Delta S0} & d_{S0S0} + \sigma_{0S0S0}^2 & d_{S0S1} \\ d_{\Delta S1} & d_{S0S1} & d_{S1S1} + \sigma_{0S1S1}^2 \end{pmatrix}$$

$$(3)$$

Based on the formula for conditional joint normal distribution (e.g. Morrison, 1980, p. 97),

$$E(\Delta_{0} \mid \widehat{\phi}_{00S}, \widehat{\phi}_{01S}) = E(\Delta_{0}) + (d_{\Delta S0}, d_{\Delta S1}) \begin{pmatrix} d_{S0S0} + \sigma_{0S0S0}^{2} & d_{S0S1} \\ d_{S0S1} & d_{S1S1} + \sigma_{0S1S1}^{2} \end{pmatrix}^{-1} \begin{pmatrix} \widehat{\phi}_{00S} \\ \widehat{\phi}_{01S} \end{pmatrix} - \begin{pmatrix} E(\phi_{00S}) \\ E(\phi_{01S}) \end{pmatrix}$$

$$(4)$$

The formula in Buyse et al (2000) is (4) with $\sigma_{0S0S0}^2 = \sigma_{0S1S1}^2 = 0$, which ignores the variability in the estimates from the application or validation trial, although, in practice, we found this made little difference.

We plug the following estimates into (4). We estimate $E(\Delta_0)$ by $\Sigma_i \ \widehat{\Delta}_i \ / k$, and we estimate $E(\widehat{\phi}_{0zS})$ by $\Sigma_i \ \widehat{\phi}_{izS} \ / k$. We estimate the components of $\widehat{V}_{\rm random}$ using the following method-of-moments approach. Let $\overline{\phi}_i = \Sigma i \ \widehat{\phi}_i \ / k$. Without imposing any

model the estimated variance of $\overline{\phi}$ is $\widehat{V}_{\text{total}} = (\widehat{\phi}_i - \overline{\phi}_i)^T \cdot (\widehat{\phi}_i - \overline{\phi}_i)/(k(k-1))$. The sampling variance for $\overline{\phi}$ is $\widehat{V}_{\text{sampling}} = \Sigma_i \widehat{V}_{\text{sampling}(i)}/k^2$, where $\widehat{V}_{\text{sampling}(i)}$ is simply $V_{\text{sampling}(i)}$ with the estimates substituted for the parameters. The method of moments estimate of $\widehat{V}_{\text{random}}$ is $\widehat{V}_{\text{random}} = \widehat{V}_{\text{total}} - \widehat{V}_{\text{sampling}}$. However for $\widehat{V}_{\text{random}}$ to be a valid estimated covariance matrix, it must be positive definite. If $\widehat{V}_{\text{random}}$ is not positive definite, we compute an adjusted covariance matrix V_{random}^* as follows. We first write $\widehat{V}_{\text{random}} = M \, Diagonal \, (\underbrace{\lambda}_i) \, M$, where $\underbrace{\lambda}_i = (\lambda_1, \lambda_2, \ldots)$ are the eigenvalues, Diagonal indicates a diagonal matrix, and M is the normalized eigenmatrix. We then replace eigenvalues that are less than or equal to zero with a small positive value to obtain a new vector of eigenvalues λ_i^* . We then computed the adjusted positive definite covariance matrix $V_{\text{random}}^* = M \, Diagonal \, (\lambda_i^*) \, M$.

Additional Reference

Morrison DF. (1976). *Multivariate Statistical Methods*. New York: McGraw-Hill Book Company.