

IPWE comparison

2020-07-01

Setting: nonstandardized, the predictors have non zero mean values

Simulation

The data sets are generated following the below parameter setting:

$$\mathbf{y}_{\mathbf{k}\mathbf{i}} = \mathbf{X}_{\mathbf{i}}(\beta_{\mathbf{k}} + \mathbf{b}_{\mathbf{k}\mathbf{i}} + \mathbf{\Gamma}_{\mathbf{k}}(\alpha' \mathbf{x})) + \epsilon_{\mathbf{k}\mathbf{i}} \quad (1)$$

where

- $\alpha = \alpha_1 + \delta \alpha_2$
 - $\alpha_1 = (1, 2, \dots, p)'$;
 - $\alpha_2 = (\alpha_{21}, \dots, \alpha_{2p})' \sim \text{Exp}(10)$
 - $\|\alpha\|_2^2 = 1$
 - when $\delta = 0$, equation (1) is a GEM model while $\delta \neq 0$, equation (1) is a non GEM model
- Suppose we have $k = \{1, 2\}$ treatments. $k = 1$ represents the placebo group while $k = 2$ represents the drug group.
- $\mathbf{X}_{\mathbf{i}} = [1, t, t^2]$, and $t = [0, 1, 2, 3, 4, 6, 8]$, which is the design matrix for fixed effect and random effect
- $\beta_1 = \beta_2 = [0, -0.1, -0.05]$
- $\mathbf{\Gamma}_1 = [0, \cos(\pi/3)/10, -\sin(\pi/3)/10]$, $\mathbf{\Gamma}_2 = [0, \sin(\pi/3)/10, \cos(\pi/3)/10]$,
- $\mathbf{b}_{1\mathbf{i}} \sim N(0, \mathbf{D}_1)$, $\mathbf{b}_{2\mathbf{i}} \sim N(0, \mathbf{D}_2)$, $\mathbf{D}_1 = \mathbf{D}_2 = \begin{pmatrix} 1 & 0 & 0 \\ 0 & 0.1 & 0 \\ 0 & 0 & 0.05 \end{pmatrix}$
- $\mathbf{x}_{\mathbf{i}} \sim MVN(\mu_x, \Sigma_x)$, $\mu_x = (1, 2, \dots, p)$, Σ_x has diagonal equals to 1 and 0.5 everywhere else.
- $\epsilon_1, \epsilon_2 \sim N(0, 1^2)$
- $p = \{3, 10\}$, which is the dimension of the predictors x_i .

The IPWE are calculated with the estimated treatment assignment by using the purity criterion and by using the log-likelihood criterion. The proportions of correct assignment are also calculated. The whole procedures are repeated for 100 times.



