

Simulation: coefficient estimation

Simulation

The data sets are generated following the below parameter setting:

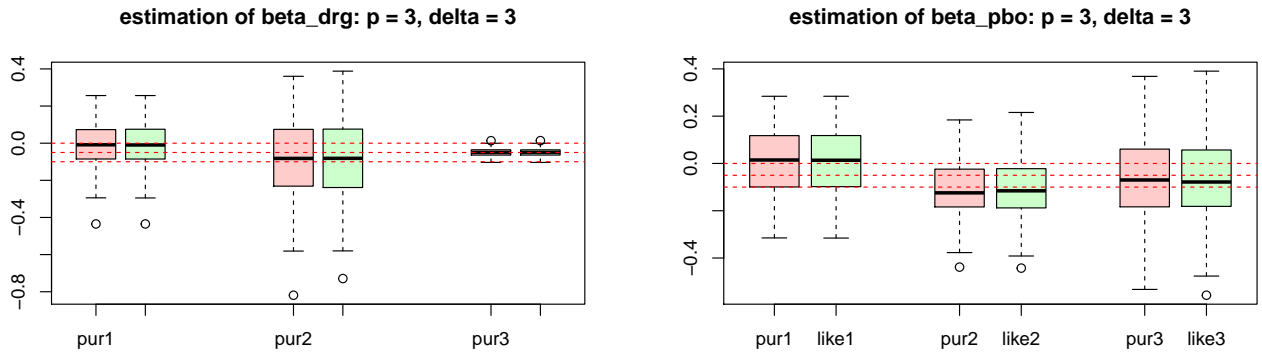
$$\mathbf{y}_{ki} = \mathbf{X}_i(\beta_k + \mathbf{b}_{ki} + \Gamma_k(\alpha' \mathbf{x})) + \epsilon_{ki} \quad (1)$$

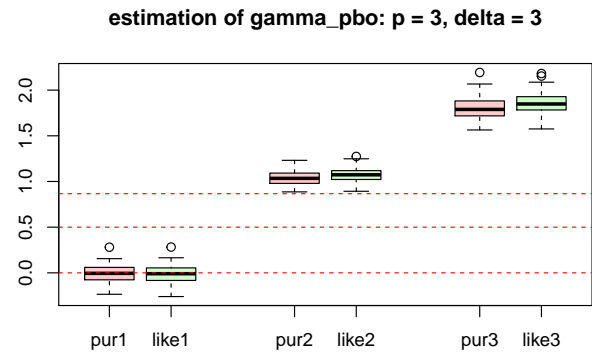
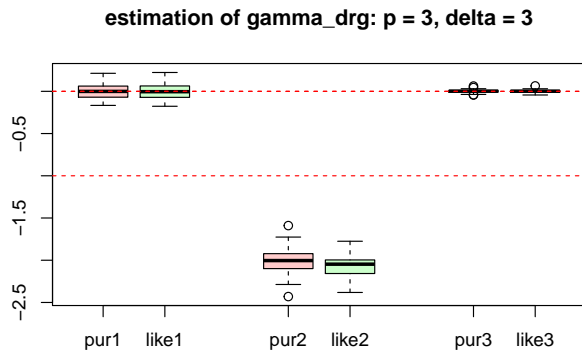
where

- $\alpha = \alpha_1 + \delta \alpha_2$
 - $\alpha_1 = (1, 2, \dots, p)'$;
 - $\alpha_2 = (\alpha_{21}, \dots, \alpha_{2p})' \sim \text{Exp}(4)$
 - $\|\alpha\|_2^2 = 1$
 - 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.
- $\mathbf{X}_i = [1, t, t^2]$, and $t = [0, 1, 2, 3, 4, 6, 8]$, which is the design matrix for fixed effect and random effect
- $\beta_1 = \beta_2 = [0, -0.1, -0.05]$
- $\Gamma_1 = [0, 1, 0]$, $\Gamma_2 = [0, \cos(\pi/3), \sin(\pi/3)]$,
- $\mathbf{b}_{1i} \sim N(0, \mathbf{D}_1)$, $\mathbf{b}_{2i} \sim N(0, \mathbf{D}_2)$, $\mathbf{D}_1 = \mathbf{D}_2 = \begin{pmatrix} 1 & 0 & 0 \\ 0 & 0.1 & 0.05 \\ 0 & 0.05 & 0.05 \end{pmatrix}$
- $\mathbf{x}_i \sim MVN(\mu_x, \Sigma_x)$, $\mu_x = \mathbf{0}_p$, 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 .

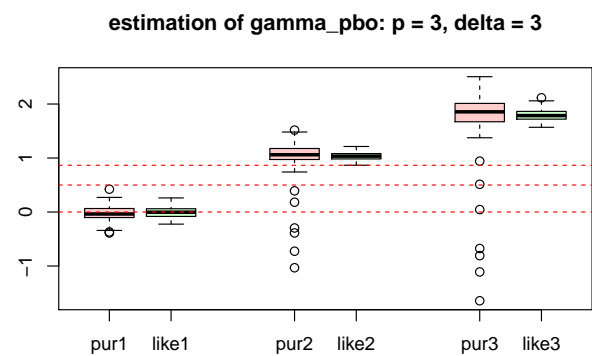
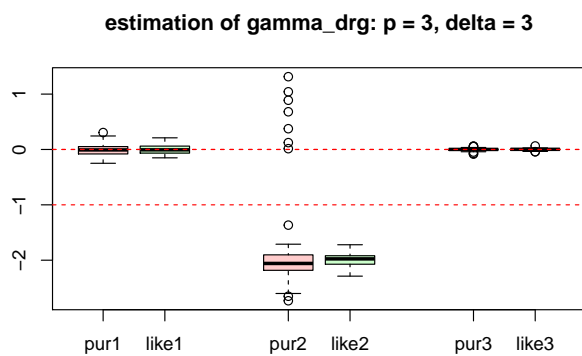
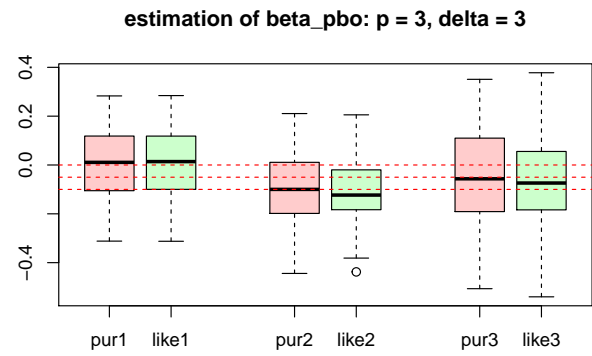
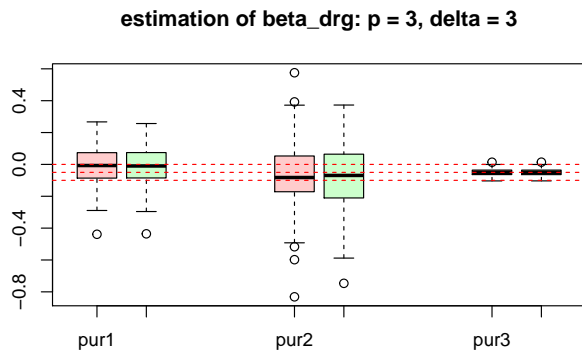
The IPWE are calculated with the estimated treatment assignment by using the purity criterion and by using the log-likelihood criterion. The proportions of correct assignment are also calculated. The whole procedures are repeated for 100 times.

Comparison of coefficient estimation: $p = 3$, $\delta = 0$

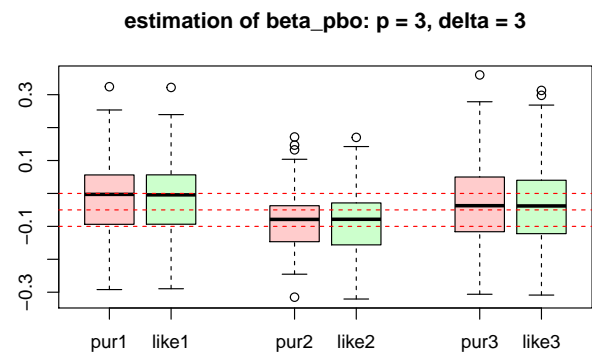
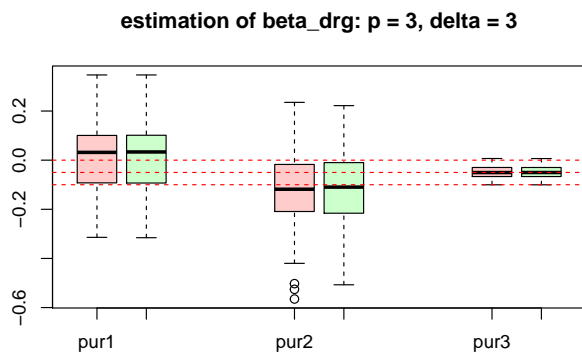


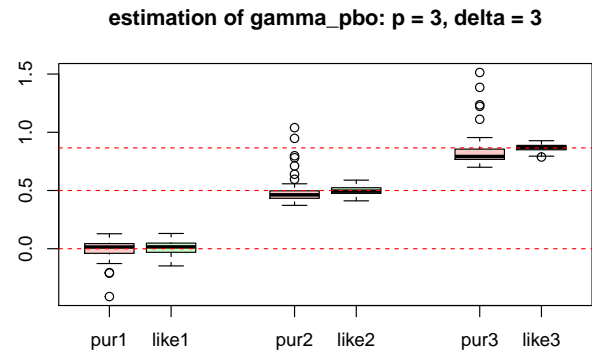
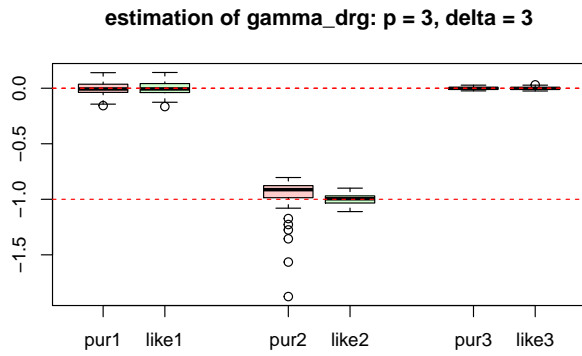


Comparison of coefficient estimation : $p = 3, \delta = 3$



Comparison of coefficient estimation : $p = 10, \delta = 0$





Comparison of coefficient estimation : $p = 10, \delta = 3$

