

# Results

2020-06-05

## Outline

- 1. Trajectories with different  $w = \alpha'x$  value
- 2. Different  $\alpha_1$  and  $\alpha_2$  values
- 3. Add L1 penalty in purity calculation function

## Trajectory

### Data generation

If the true outcome generation models are (with different true  $\alpha$ ):

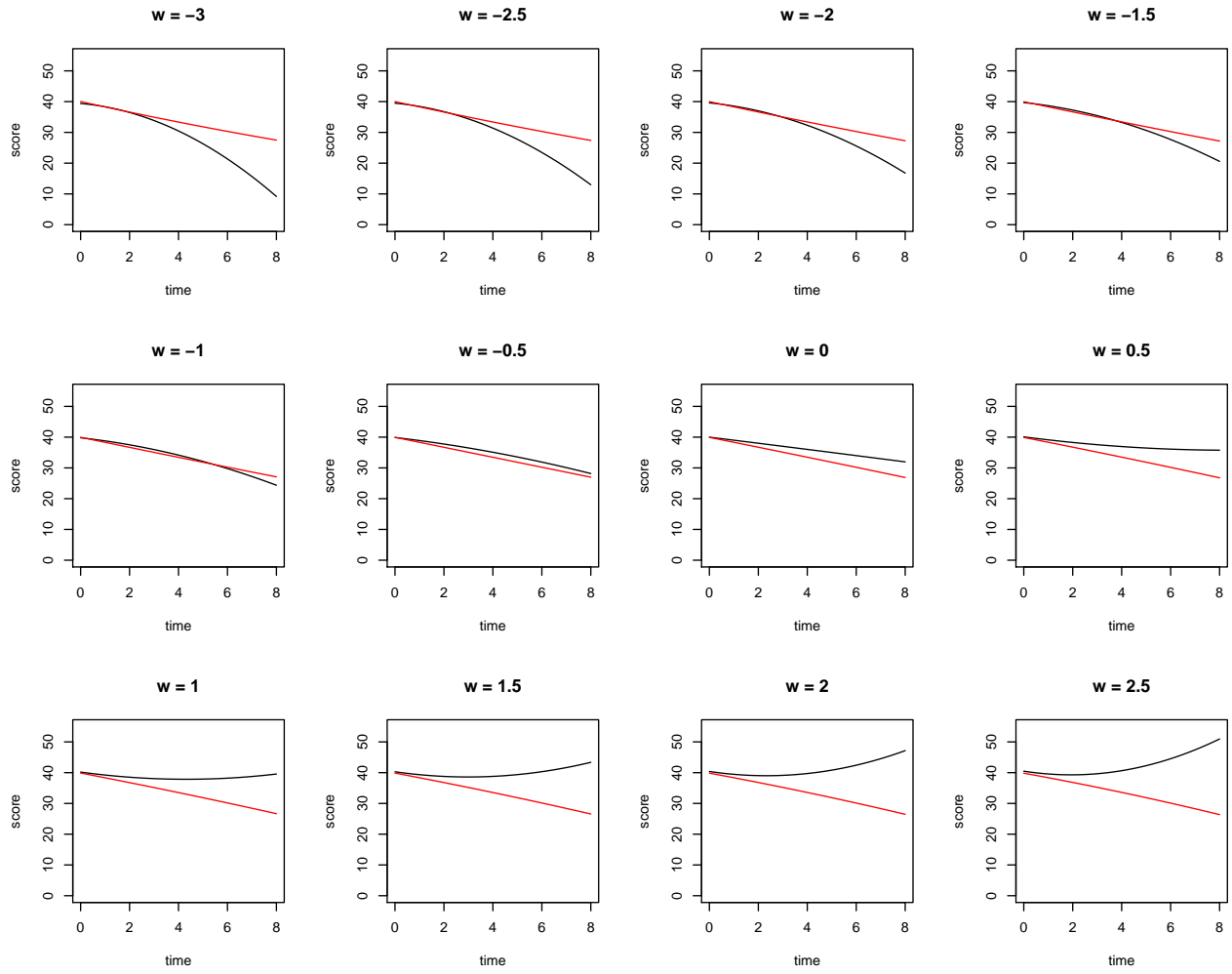
$$Y_{drg} = S(\beta_{drg} + \Gamma_{drg}(\alpha'_{drg}x)) + Sb_{drg} + \epsilon_{drg}$$

$$Y_{pbo} = S(\beta_{pbo} + \Gamma_{pbo}(\alpha'_{pbo}x)) + Sb_{pbo} + \epsilon_{pbo}$$

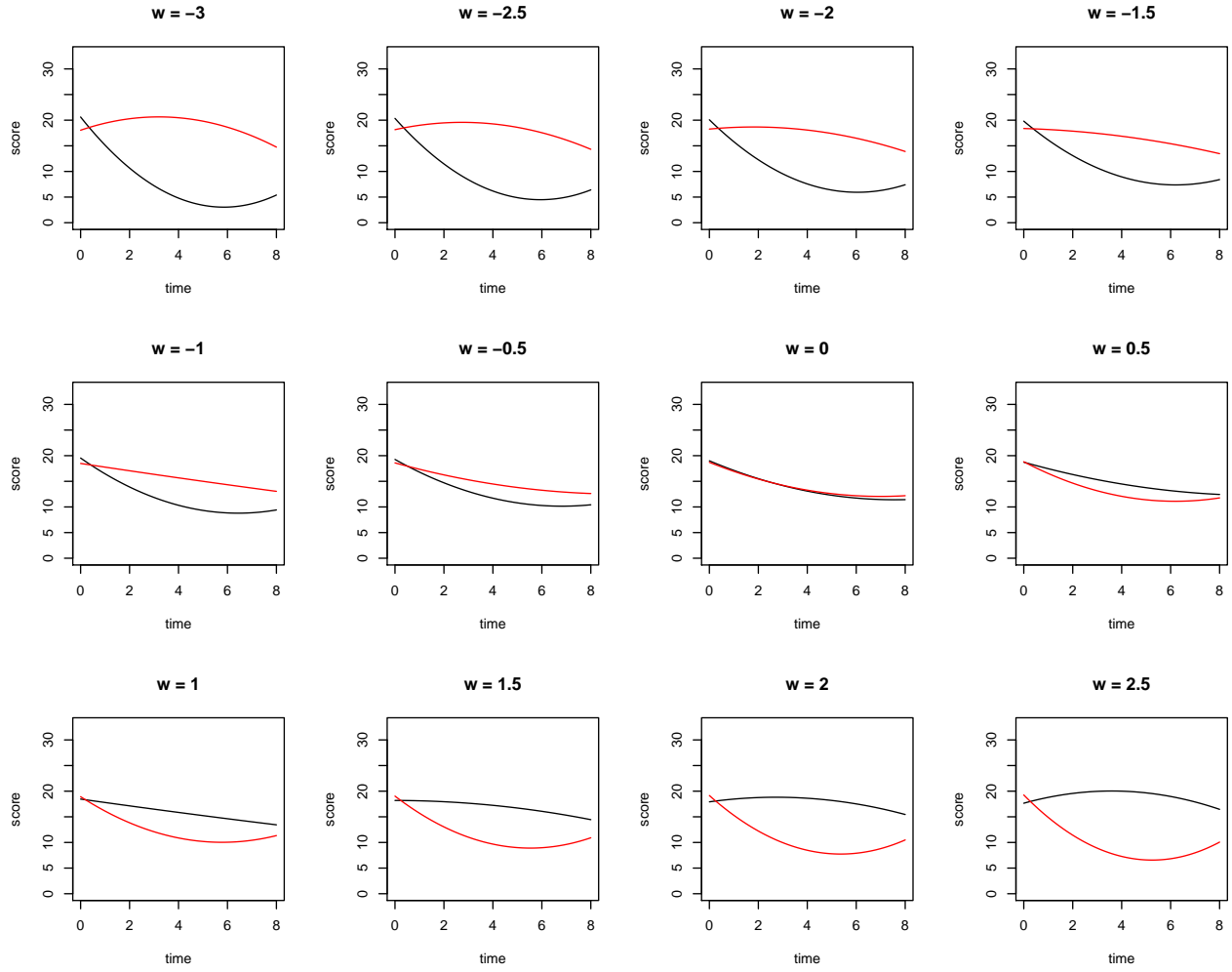
where

- $S = [1, t, t^2]$ ,  $t = [0, 1, 2, 3, 4, 6, 8]$  is the design matrix for fixed effect and random effect
- $x_1 \sim MVN(\mu_x, \Sigma_x)$ ,  $\mu_x = \mathbf{0}_p$ ,  $\Sigma_x$  has diagonal equals to 1 and 0.5 anywhere else. The dimension of  $x$  is set to be  $p = 20$ .
- 80 noises are also added in the dataset, each of them are generated from  $x_{2,j} \sim N(0, 1)$ . Therefore, there are  $np = 100$  covariates in total.
- $\beta_{drg} = [40, -1, -0.02]$ ,  $\beta_{pbo} = [40, -1.1, -0.02]$
- $\Gamma_{drg} = [0, 0.1 \cos(\theta), 0.1 \sin(\theta)]$ ,  $\Gamma_{pbo} = [0, 0.01, 0]$ ,  $\theta = \frac{\pi}{2}$
- $b_{drg} = \begin{bmatrix} 1 & 0 & 0 \\ 0 & 0.01 & 0 \\ 0 & 0 & 0.02 \end{bmatrix}$ ,  $b_{pbo} = \begin{bmatrix} 1 & 0 & 0 \\ 0 & 0.01 & 0.005 \\ 0 & 0.005 & 0.01 \end{bmatrix}$
- $\epsilon_{drg}, \epsilon_{pbo} \sim N(0, 3^2)$
- $\alpha = 1_p$

The trajectory plot with different  $w$  value, black line is the drug group while the red line presents the placebo group.



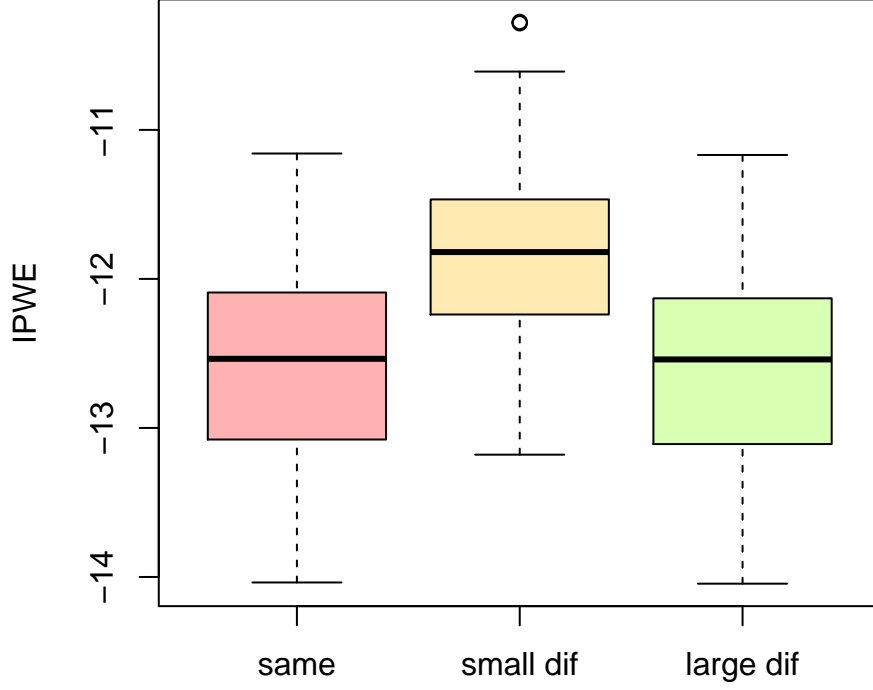
The trajectory plot for EMBARC dataset, used the covariates selected by the forward and backward selection.



## Diff alph

- same alpha:  $\alpha_1 = \alpha_2 = I_p$
- small difference:
  - $\alpha_1 = I_p$
  - $\alpha_2 = [I_{\frac{p}{2}}, 1 : \frac{p}{2}]$
- big difference:
  - $\alpha_1 = I_p$
  - $\alpha_2 = 1 : p$

## LSI: p: 20 ; angle: 90



## LASSO

- Penalty added in the Purity function  $G(\alpha)$

$$G_p(\alpha) = A_0 + A_1 \hat{\mu}'_x \alpha + \frac{A_2}{2} [\alpha' \hat{\Sigma}_x \alpha + \alpha' \hat{\mu}_x \hat{\mu}'_x \alpha] - \lambda \|\alpha\|_1$$

where

$$A_0 = -q + \frac{1}{2} \text{tr}(\hat{D}_2^{-1} \hat{D}_1) + \frac{1}{2} \text{tr}(\hat{D}_1^{-1} \hat{D}_2) + \frac{1}{2} (\hat{\beta}_1 - \hat{\beta}_2)' (\hat{D}_1^{-1} + \hat{D}_2^{-1}) (\hat{\beta}_1 - \hat{\beta}_2)$$

$$A_1 = (\hat{\beta}_1 - \hat{\beta}_2)' (\hat{D}_1^{-1} + \hat{D}_2^{-1}) (\hat{\Gamma}_1 - \hat{\Gamma}_2)$$

$$A_2 = (\hat{\Gamma}_1 - \hat{\Gamma}_2)' (\hat{D}_1^{-1} + \hat{D}_2^{-1}) (\hat{\Gamma}_1 - \hat{\Gamma}_2)$$

$q$  is the dimension of  $D$  matrix.

- Calculation derivation:
  - approximation
  - sign function
- Data set
  - Simulation (100 covariates, 80 noises)
  - EMBARC (215 covariates)

## Attempt 1

1. Initial  $\alpha^{(0)}$  and  $\lambda$
2. Fit linear mixed effect model and estimate  $\hat{\beta}_1^{(0)}, \hat{\beta}_2^{(0)}, \hat{\Gamma}_1^{(0)}, \hat{\Gamma}_2^{(0)}, \hat{D}_1^{(0)}, \hat{D}_2^{(0)}$  and then associated  $\hat{A}_0^{(0)}, \hat{A}_1^{(0)}, \hat{A}_2^{(0)}$

3. Update  $\alpha$ :

- Maximize the penalized purity function
- Since no class from, an approximation of  $\|\alpha\|_1$  is conducted

$$G_p(\alpha) = A_0 + A_1 \hat{\mu}'_x \alpha + \frac{A_2}{2} [\alpha' \hat{\Sigma}_x \alpha + \alpha' \hat{\mu}_x \hat{\mu}'_x \alpha] - \lambda \|\alpha\|_1$$

4. Repeat 2-3 until converge.

5. Tune parameter

- For a given sequence of  $\lambda \in \{0.001, 0.01, 0.1, 1, 10, 100\}$ , find the  $\hat{\alpha}$  that maximizes the penalized purity function.
- Do 10 fold cross validation. Use 9 of the part to train the data, fit LME with  $\hat{\alpha}'x$  and calculate  $\hat{\beta}_1, \hat{\beta}_2, \hat{\Gamma}_1, \hat{\Gamma}_2, \hat{D}_1, \hat{D}_2$ .
- Calculate the IPWE for each of the 10 fold cross validation
- Repeat IPWE calculation for 100 times
- Choose the  $\lambda$  that achieve the max IPWE.

6.

- $\lambda \in \{0.001, 0.01, 0.1, 1, 10, 100\}$ , the one with the largest IPWE  $l_1$
- $\lambda \in seq(l_1/10, l_1 * 10, l_1)$ , the one with the largest IPWE  $l_2$
- $\lambda \in seq(l_2/10, l_2 * 10, l_2)$ , the one with the largest IPWE  $l_3$ , which is the tuned parameter  $\hat{\lambda}$

For the penalty, we use the approximation

$$|\alpha_i^{(j+1)}| = |\alpha_i^{(j)}| + \frac{1}{2|\alpha_i^{(j)}|} \{(\alpha_i^{(j+1)})^2 - (\alpha_i^{(j)})^2\}$$

And

$$\|\alpha^{(j+1)}\|_1 \approx \|\alpha^{(j)}\|_1 + \frac{1}{2} \sum_{i=1}^p \frac{(\alpha_i^{(j+1)})^2}{|\alpha_i^{(j)}|} - \frac{1}{2} \sum_{i=1}^p \frac{(\alpha_i^{(j)})^2}{|\alpha_i^{(j)}|} \propto \frac{1}{2} \sum_{i=1}^p \frac{(\alpha_i^{(j+1)})^2}{|\alpha_i^{(j)}|}$$

Therefore,

$$\begin{aligned} \operatorname{argmax} G_p(\alpha) &\approx \operatorname{argmax} [A_0 + A_1 \hat{\mu}'_x \alpha^{(j+1)} + \frac{A_2}{2} [(\alpha^{(j+1)})' \hat{\Sigma}_x \alpha^{(j+1)} + (\alpha^{(j+1)})' \hat{\mu}_x \hat{\mu}'_x \alpha^{(j+1)}] - \frac{\lambda}{2} \sum_{i=1}^p \frac{(\alpha_i^{(j+1)})^2}{|\alpha_i^{(j)}|}] \\ &= \operatorname{argmax} [A_0 + A_1 \hat{\mu}'_x \alpha^{(j+1)} + \frac{A_2}{2} [(\alpha^{(j+1)})' \hat{\Sigma}_x \alpha^{(j+1)} + (\alpha^{(j+1)})' \hat{\mu}_x \hat{\mu}'_x \alpha^{(j+1)}] - \frac{\lambda}{2} (\alpha^{(j+1)})' \Psi \alpha^{(j+1)}] \end{aligned}$$

Calculation the derivation:

$$\begin{aligned} \frac{\partial A_1 \hat{\mu}'_x \alpha^{(j+1)}}{\partial \alpha^{(j+1)}} &= A_1 \hat{\mu}'_x \\ \frac{\partial (\frac{A_2}{2} [(\alpha^{(j+1)})' (\hat{\Sigma}_x + \hat{\mu}_x \hat{\mu}'_x) \alpha^{(j+1)}])}{\partial \alpha^{(j+1)}} &= A_2 (\alpha^{(j+1)})' (\hat{\Sigma}_x + \hat{\mu}_x \hat{\mu}'_x) \\ \frac{\partial \frac{\lambda}{2} (\alpha^{(j+1)})' \Psi \alpha^{(j+1)}}{\partial \alpha^{(j+1)}} &= \lambda (\alpha^{(j+1)})' \Psi \\ \Psi &= \operatorname{diag}\{\frac{1}{\alpha_1^{(j)}}, \dots, \frac{1}{\alpha_p^{(j)}}\} \end{aligned}$$

Therefore,

$$\begin{aligned} A_1 \hat{\mu}'_x + A_2 (\alpha^{(j+1)})' (\hat{\Sigma}_x + \hat{\mu}_x \hat{\mu}'_x) - \lambda (\alpha^{(j+1)})' \Psi &\equiv 0 \\ \Rightarrow \alpha^{(j+1)} &= A_1 (\lambda \Psi - A_2 (\hat{\Sigma}_x + \hat{\mu}_x \hat{\mu}'_x))^{-1} \hat{\mu}_x \end{aligned}$$

## Attempt 2

1. Initial  $\alpha^{(0)}$  and  $\lambda$
2. Fit LME, and estimate  $\beta_1^{(0)}, \beta_2^{(0)}, \Gamma_1^{(0)}, \Gamma_2^{(0)}, D_1^{(0)}, D_2^{(0)}$
3. Calculate  $A_1^{(0)}, A_2^{(0)}, B_1^{(0)}, B_2^{(0)}, B^{(0)}$
4. Update  $\alpha$ : for the  $i$ th iteration ( $i = 0, 1, 2, \dots$ ),
  - for each  $j \in \{1, 2, \dots, p\}$ , calculate  $\alpha_j^{(i+1)}$  and update  $\alpha_j^{(i+1)}$
5. Calculate  $w = \alpha_j^{(i+1)} x$  and fit new LME, update  $\beta_1^{(i+1)}, \beta_2^{(i+1)}, \Gamma_1^{(i+1)}, \Gamma_2^{(i+1)}, D_1^{(i+1)}, D_2^{(i+1)}$
6. Repeat 3-5 until convergence.

$$\hat{\alpha}_j^{(i+1)} = \begin{cases} -\frac{1}{(B_j^{(i)})' B_j^{(i)}} (B_j^{(i)})' B_{-j}^{(i)} \alpha_{-j}^{(i)} - \frac{B_{1,j}^{(i)}}{2(B_j^{(i)})' B_j^{(i)}} + \frac{\lambda}{2(B_j^{(i)})' B_j^{(i)}} & (B_j^{(i)})' B_{-j}^{(i)} \alpha_{-j}^{(i)} + \frac{B_{1,j}^{(i)}}{2} > \frac{\lambda}{2} \\ 0 & |(B_j^{(i)})' B_{-j}^{(i)} \alpha_{-j}^{(i)} + \frac{B_{1,j}^{(i)}}{2}| \leq \frac{\lambda}{2} \\ -\frac{1}{(B_j^{(i)})' B_j^{(i)}} (B_j^{(i)})' B_{-j}^{(i)} \alpha_{-j}^{(i)} - \frac{B_{1,j}^{(i)}}{2(B_j^{(i)})' B_j^{(i)}} - \frac{\lambda}{2(B_j^{(i)})' B_j^{(i)}} & (B_j^{(i)})' B_{-j}^{(i)} \alpha_{-j}^{(i)} + \frac{B_{1,j}^{(i)}}{2} < -\frac{\lambda}{2} \end{cases}$$

Parameter tuning

- For a given sequence of  $\lambda \in \{0.001, 0.01, 0.1, 1, 10, 100\}$ , find the  $\hat{\alpha}$  that maximizes the penalized purity function.
- Do 10 fold cross validation. Use 9 of the part to train the data, fit LME with  $\hat{\alpha}'x$  and calculate  $\hat{\beta}_1, \hat{\beta}_2, \hat{\Gamma}_1, \hat{\Gamma}_2, \hat{D}_1, \hat{D}_2$ .
- Calculate the IPWE for each of the 10 fold cross validation
- Repeat IPWE calculation for 100 times
- Choose the  $\lambda$  that achieve the max IPWE.

Update  $\alpha$  calculation

Maximum value of

$$G_p(\alpha) = A_0 + A_1 \hat{\mu}_x' \alpha + \frac{A_2}{2} [\alpha' \hat{\Sigma}_x \alpha + \alpha' \hat{\mu}_x \hat{\mu}_x' \alpha] - \lambda \|\alpha\|_1, \quad \lambda > 0$$

Or Minimum value of

$$G_p(\alpha) = -\alpha' B_2 \alpha - B_1 \alpha + \lambda \|\alpha\|_1$$

where

- $B_1 = A_1 \mu_x'$
- $B_2 = \frac{A_2}{2} (\hat{\Sigma}_x + \hat{\mu}_x \hat{\mu}_x')$ . And let  $B_2 = B' B$

Then

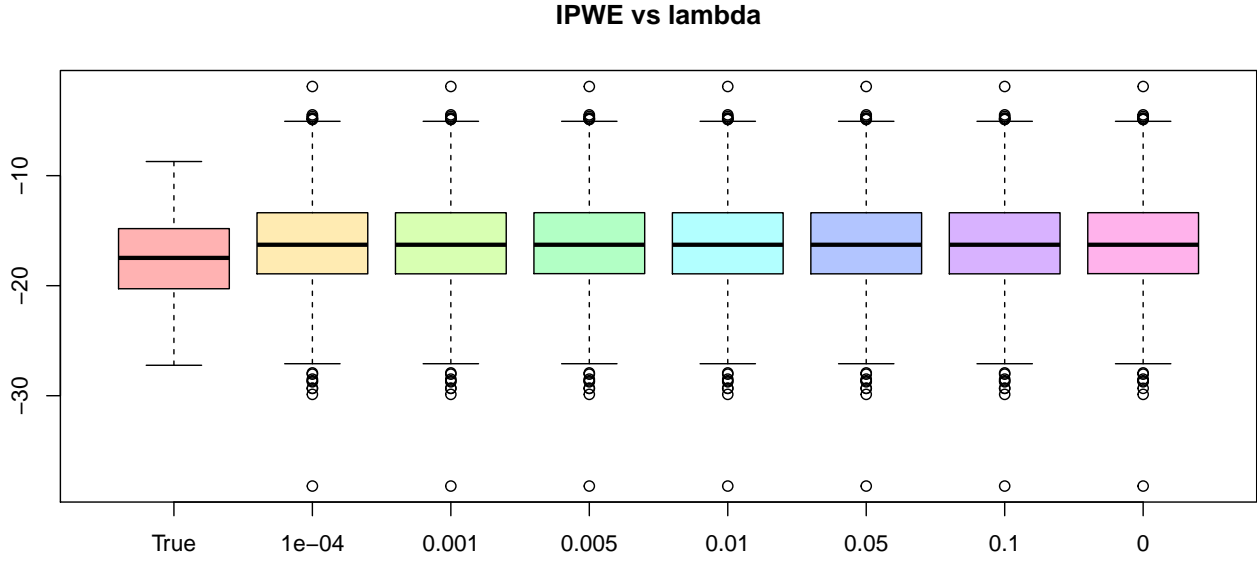
$$\frac{\partial G_p(\alpha)}{\partial \alpha_j} = -2B_j' B_j \alpha_j - 2B_j' B_{-j} \alpha_{-j} - B_{1,j} + \lambda \text{sign}(\hat{\alpha}_j) \equiv 0$$

$$\alpha_j = -\frac{1}{B_j' B_j} B_j' B_{-j} \alpha_{-j} - \frac{B_{1,j}}{2B_j' B_j} + \frac{\lambda}{2B_j' B_j} \text{sign}(\hat{\alpha}_j)$$

$\Rightarrow$

$$\hat{\alpha}_j = \begin{cases} -\frac{1}{B_j' B_j} B_j' B_{-j} \alpha_{-j} - \frac{B_{1,j}}{2B_j' B_j} + \frac{\lambda}{2B_j' B_j} & B_j' B_{-j} \alpha_{-j} + \frac{B_{1,j}}{2} > \frac{\lambda}{2} \\ 0 & |B_j' B_{-j} \alpha_{-j} + \frac{B_{1,j}}{2}| \leq \frac{\lambda}{2} \\ -\frac{1}{B_j' B_j} B_j' B_{-j} \alpha_{-j} - \frac{B_{1,j}}{2B_j' B_j} - \frac{\lambda}{2B_j' B_j} & B_j' B_{-j} \alpha_{-j} + \frac{B_{1,j}}{2} < -\frac{\lambda}{2} \end{cases}$$

## Boxplot of IPWE estimation vs different $\lambda$ values



## IPWE boxplot, with 100 covariates

If we input all the 100 covariates in the three models:

## IPWE boxplot for EMBARC dataset

For example, the  $\hat{\alpha}$  value

```
## [1] 0.082 -0.085 0.023 -0.104 0.000 0.000 -0.047 -0.103 -0.054 -0.035
## [11] -0.028 0.000 -0.074 -0.119 0.000 0.005 -0.126 -0.134 -0.139 -0.020
## [21] -0.028 -0.068 -0.060 -0.061 -0.135 -0.124 -0.032 -0.095 -0.126 -0.059
## [31] -0.004 -0.022 -0.049 -0.097 0.000 -0.103 0.000 0.000 -0.013 -0.119
## [41] -0.043 -0.040 -0.028 -0.074 0.018 -0.063 -0.136 0.000 -0.011 -0.102
## [51] -0.106 -0.131 -0.023 0.006 -0.086 -0.060 -0.067 -0.041 -0.137 -0.134
## [61] -0.030 -0.116 -0.101 -0.048 -0.030 -0.031 -0.052 -0.089 -0.021 0.000
## [71] -0.101 -0.090 -0.110 -0.031 -0.060 -0.064 -0.019 0.000 -0.105 -0.080
## [81] -0.128 -0.127 -0.123 -0.033 -0.028 -0.061 0.003 -0.010 0.000 0.000
## [91] -0.078 -0.072 -0.114 -0.024 0.005 -0.061 -0.055 -0.025 0.000 -0.024
## [101] -0.088 -0.030 -0.076 -0.021 -0.112 -0.066 -0.098 -0.065 0.023 -0.040
## [111] 0.000 0.000 -0.025 -0.076 -0.017 -0.084 -0.035 -0.003 -0.127 -0.094
## [121] -0.074 -0.033 -0.045 -0.032 -0.030 -0.055 -0.043 0.000 -0.059 -0.014
## [131] -0.049 -0.009 0.020 -0.062 -0.103 -0.077 -0.060 -0.046 -0.062 -0.113
## [141] -0.021 -0.084 -0.020 0.026 -0.074 -0.103 -0.104 -0.063 -0.073 -0.072
## [151] -0.007 0.000 -0.070 -0.093 -0.108 -0.130 -0.125 -0.023 -0.029 -0.068
## [161] 0.004 0.016 0.000 0.000 -0.042 -0.111 -0.096 -0.027 0.021 -0.063
## [171] -0.040 0.020 0.000 -0.034 -0.102 -0.045 -0.079 -0.015 -0.135 -0.060
## [181] -0.045 -0.069 0.000 -0.011 0.015 0.000 -0.045 -0.103 -0.060 -0.081
## [191] 0.005 -0.040 -0.108 -0.098 -0.081 -0.035 -0.037 -0.018 -0.021 -0.068
## [201] -0.067 -0.016 -0.066 -0.014 -0.093 -0.038 -0.003 0.003 -0.085 -0.040
## [211] -0.031 -0.041 -0.065 -0.124 -0.030 0.057 0.022 0.000 0.000 0.000
## [221] 0.009 0.000 0.000 -0.015 -0.005 -0.024 -0.013 0.007 0.015 0.000
## [231] 0.000 0.000 0.000
```

