

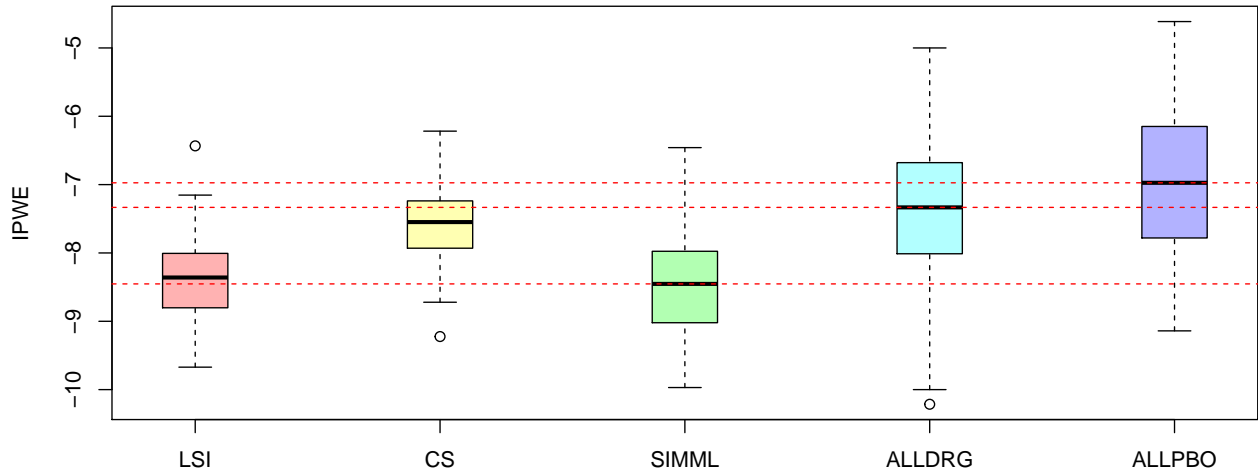
backward

2020-05-22

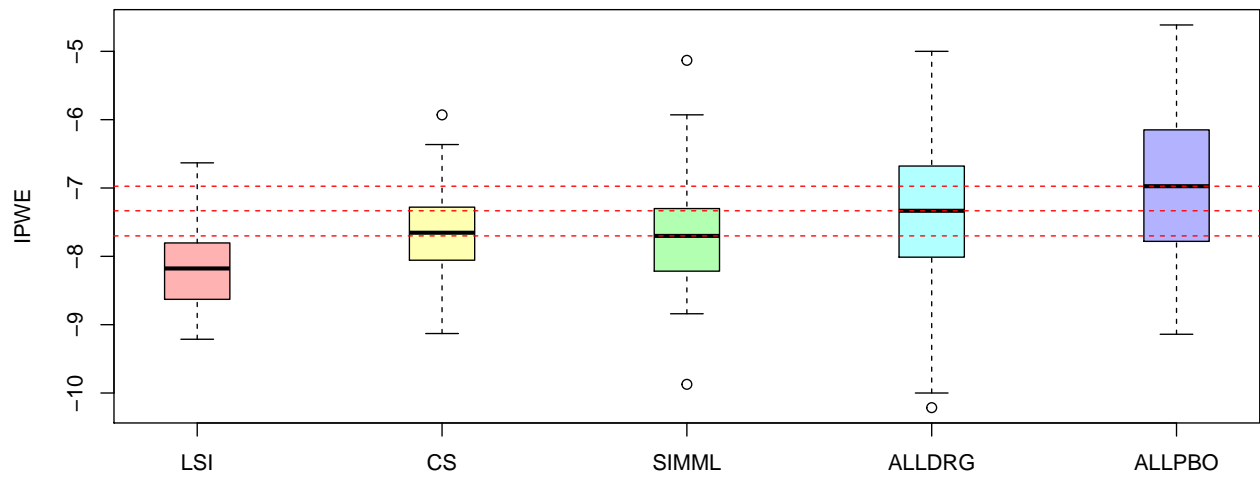
Procedures to pick up n biosignatures:

- 1. Let M_0 denote the full model, which contains all covariates
- 2. For $k = 0, \dots, p - 1$ ($p = 31$ covariates):
 - (a) consider all $p - k$ models that delete the covariates in M_k with one additional covariate.
 - (b) choose the worst among these $p - k$ models and call it M_{k+1} .
 - the worst is defined as whether:
 - * smallest purity
 - * largest IPWE
- 3. For the selected covariates, conduct 10 fold CV with methods (longitudinal single index, linear change score method, SIMML). This procedure is repeated for 100 times.

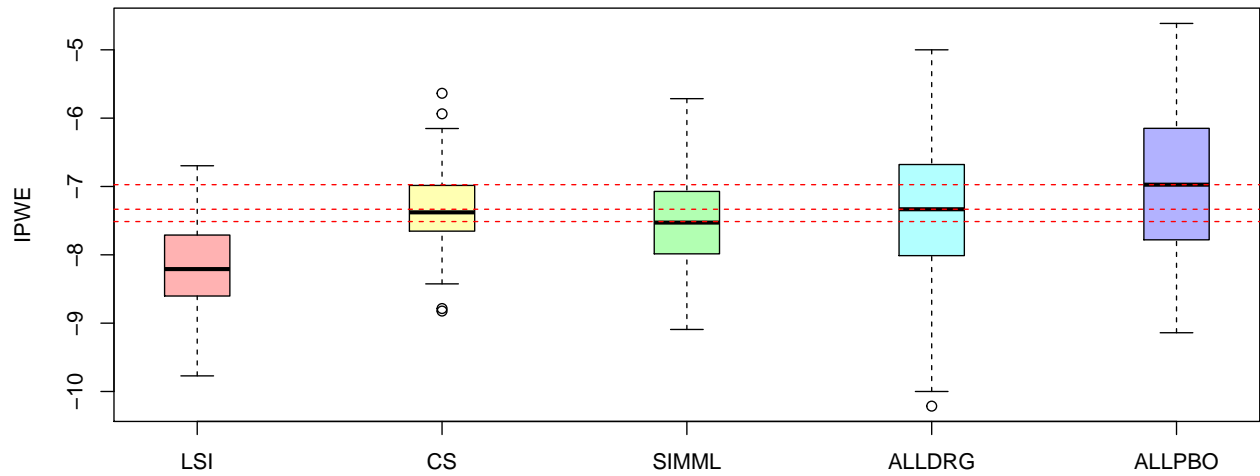
backward selection: 13 covariates



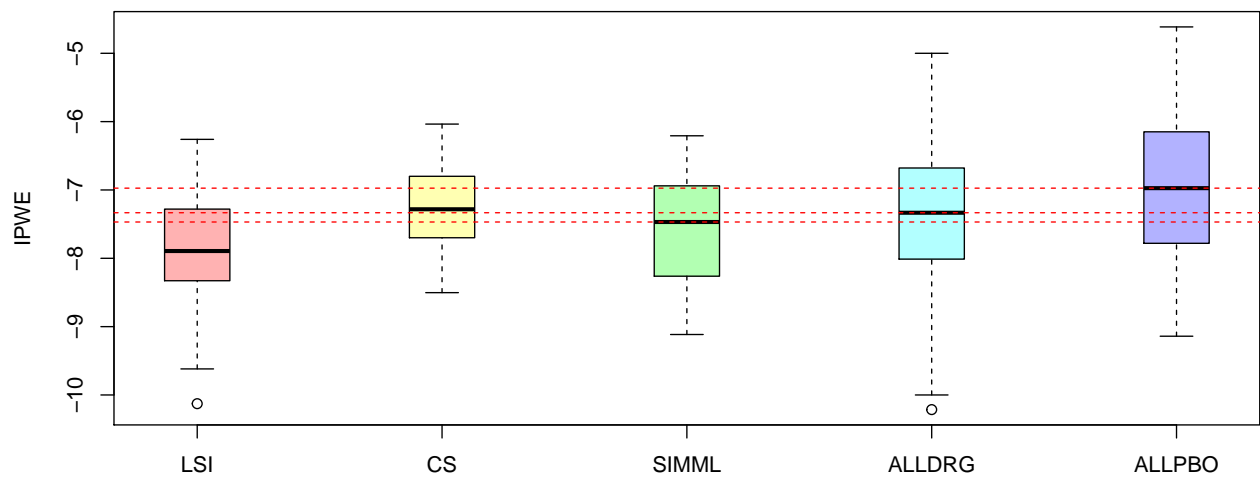
backward selection: 14 covariates



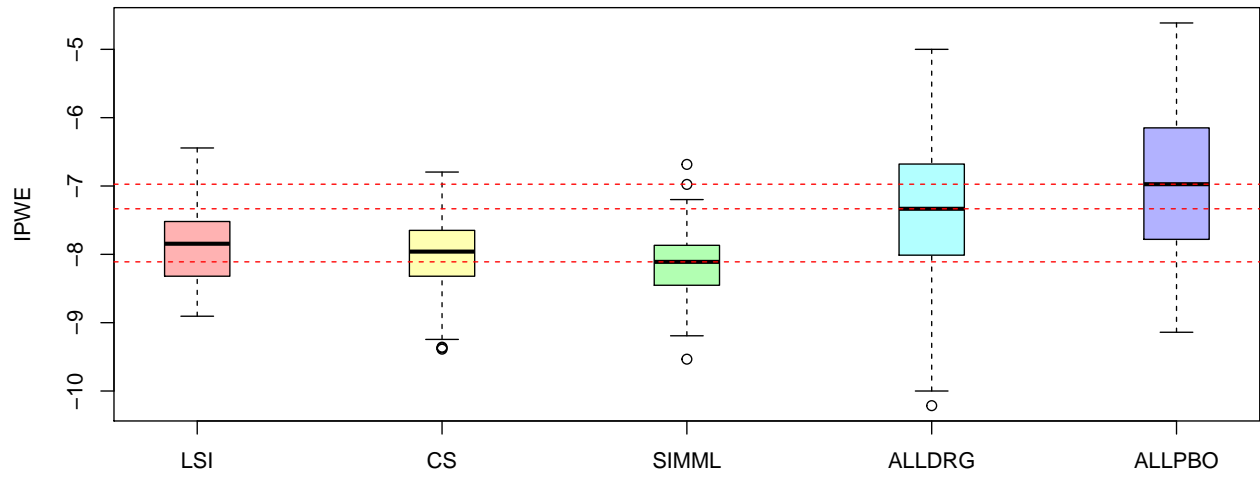
backward selection: 16 covariates



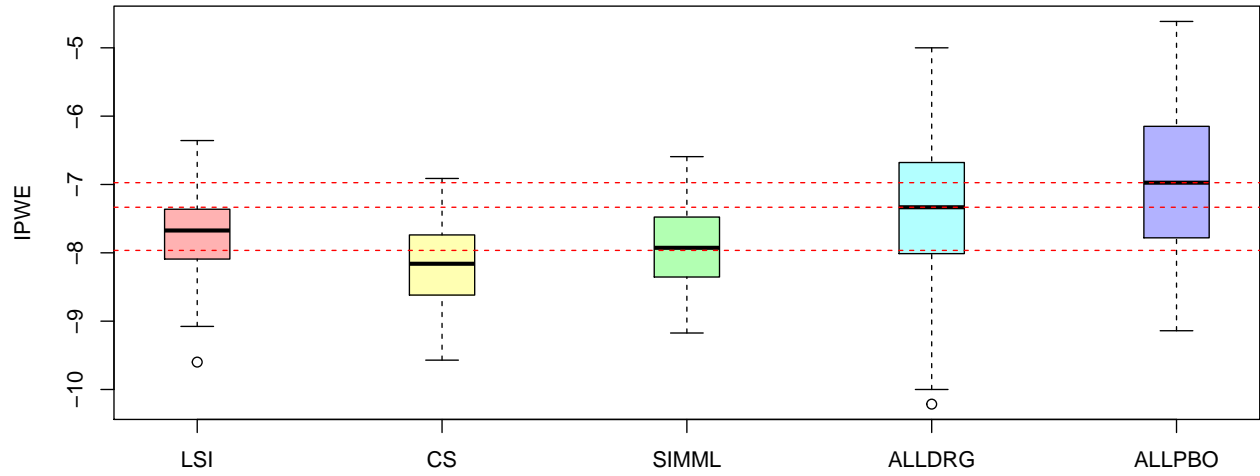
backward selection: 18 covariates



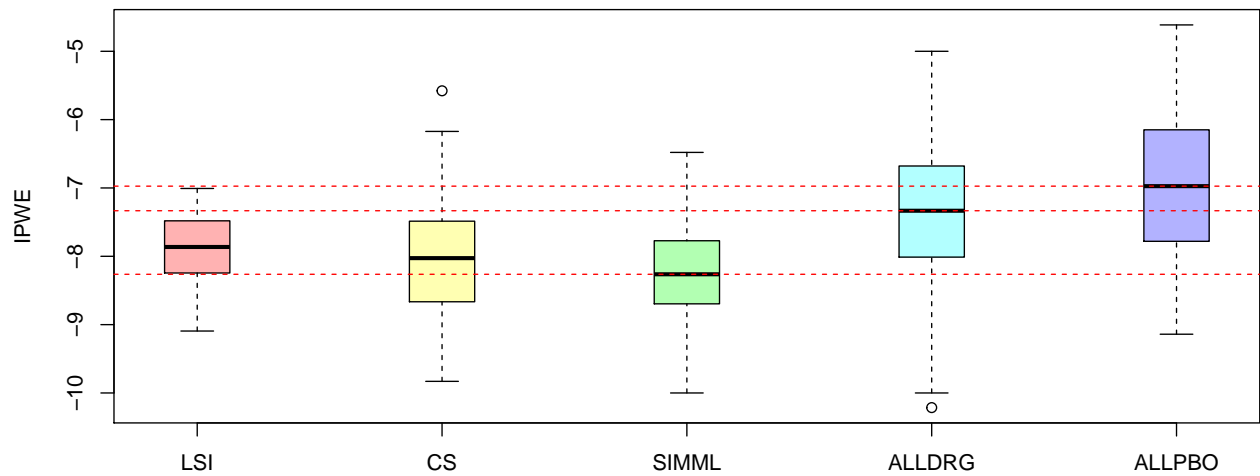
backward selection: 20 covariates



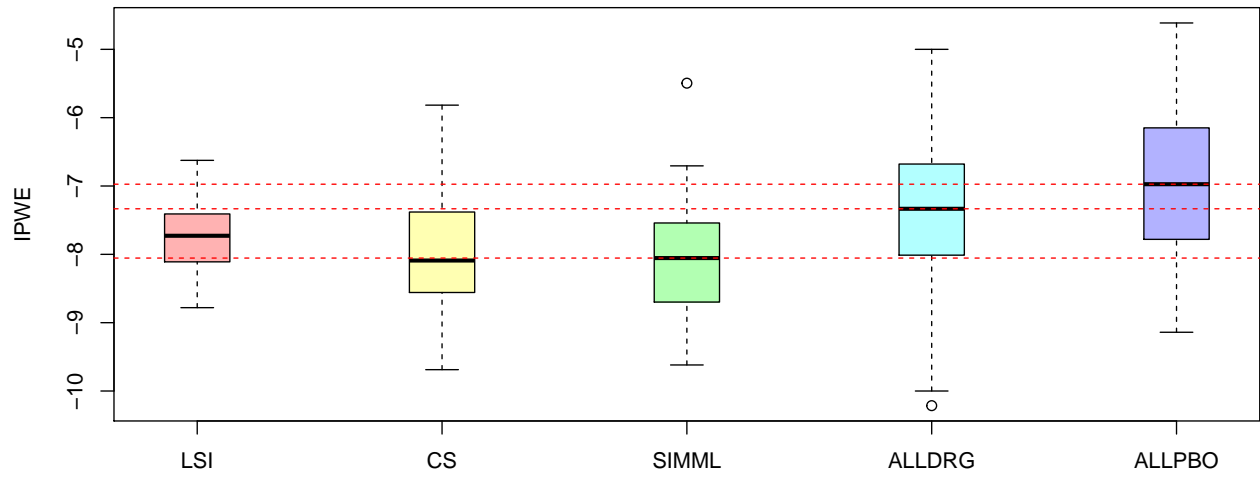
backward selection: 21 covariates



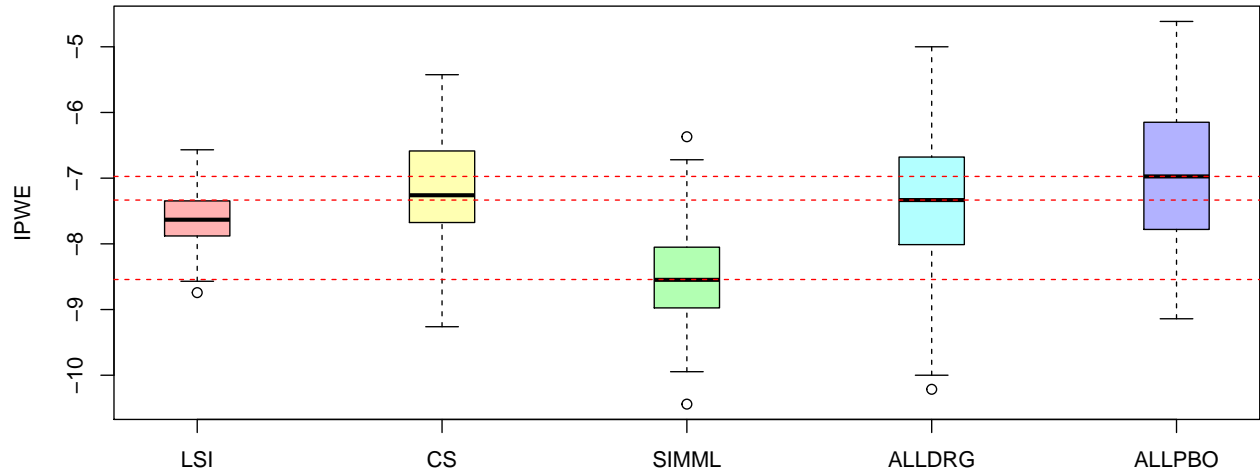
backward selection: 24 covariates



backward selection: 25 covariates



backward selection: 28 covariates



backward selection: 30 covariates

