

(M) Top PSID-1: optimal_pair_stuermer

Partial R² of confounder(s) with the outcome

0.5

0.4

0.3

0.2

0.1

0.0

0.0

0.1

0.2

0.3

0.4

Partial R² of confounder(s) with the treatment

3x re75
(-1511.087)

2x re75
(-1557.722)

1x re75
(-1604.352)

Unadjusted
(-1650.976)

4500

4000

3500

3000

2500

2000

1500

1000

500

-500

-1000

-1500

