

(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)

-1500 -1000 -500 0

