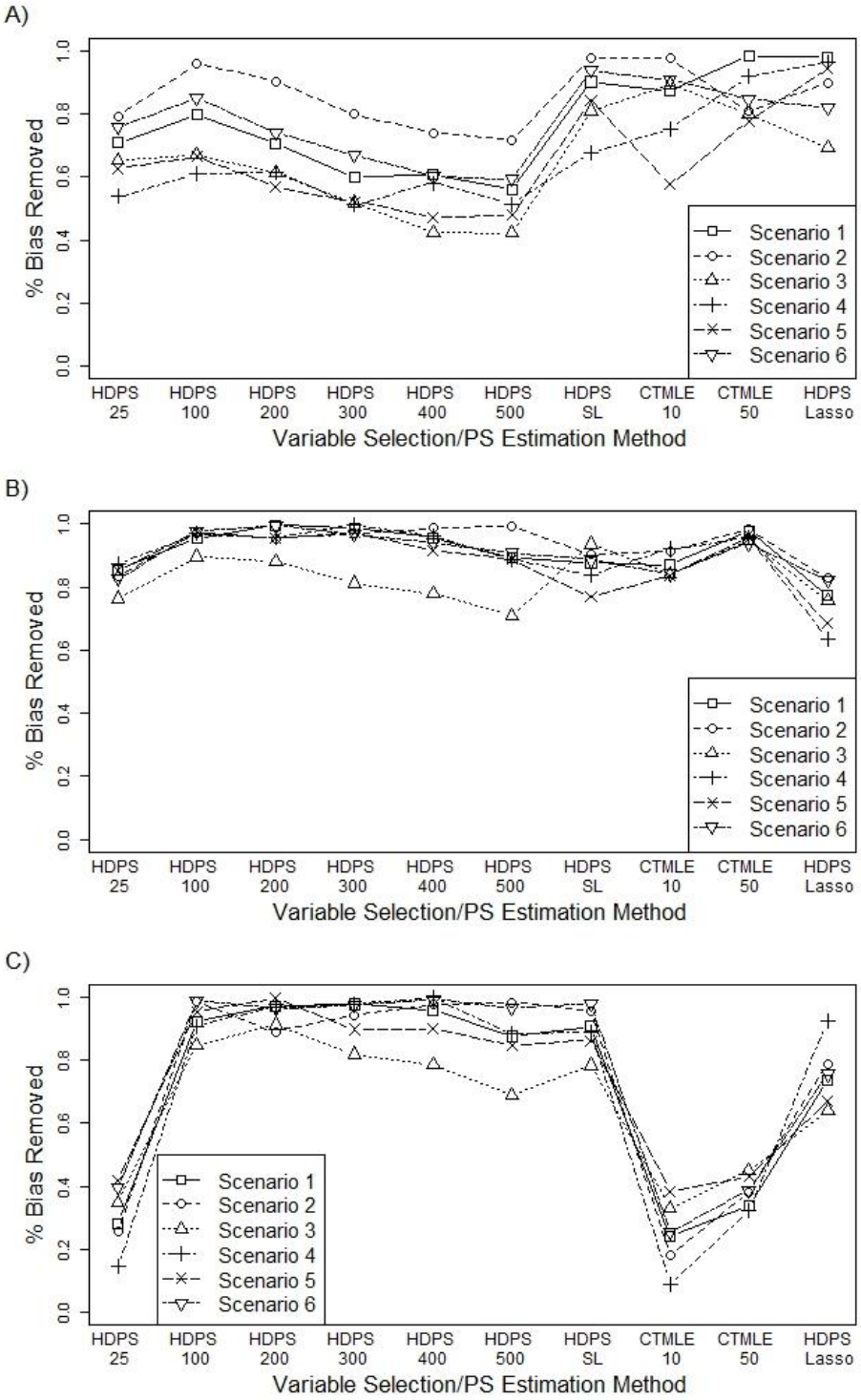
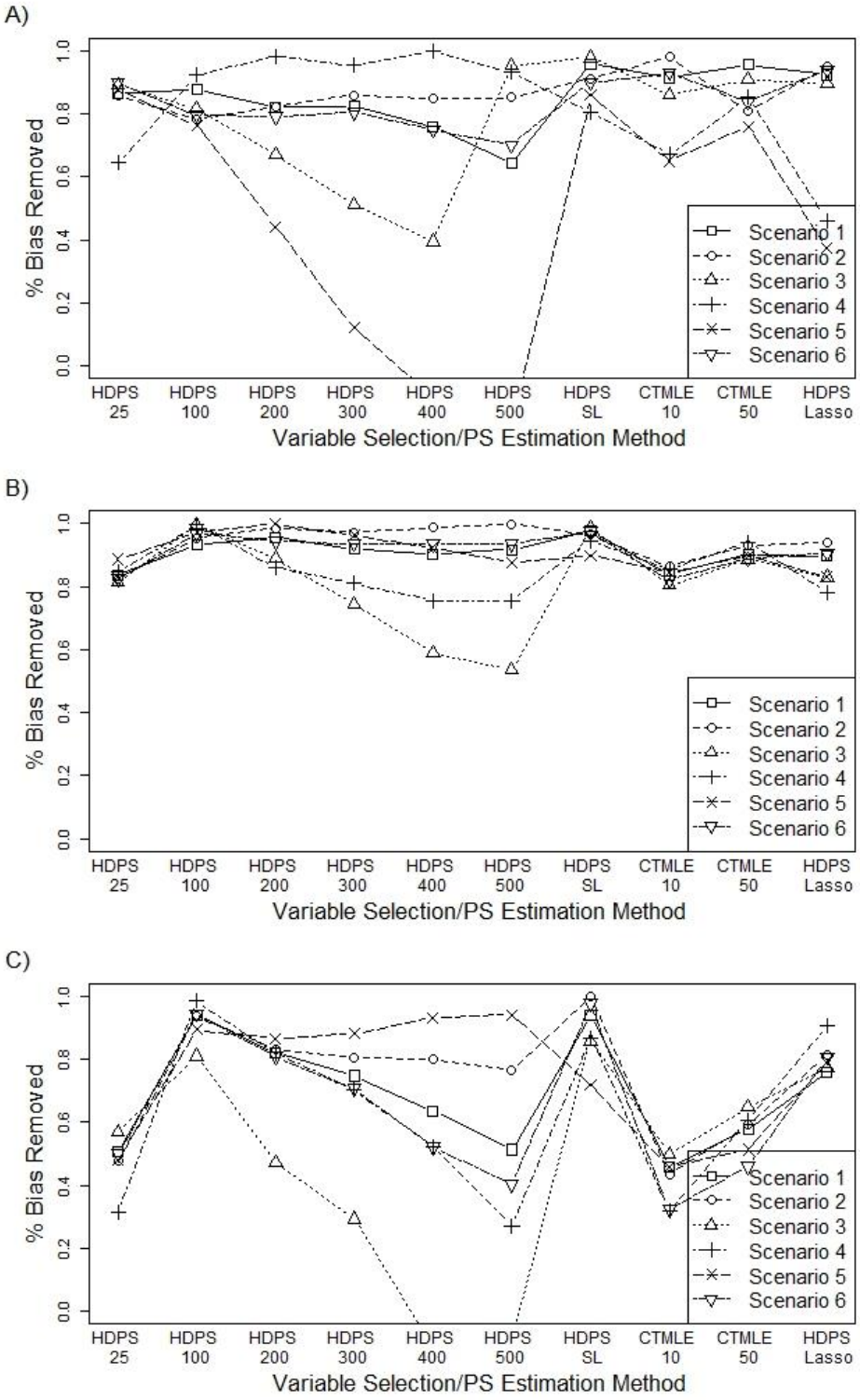
Appendix 4. Supplemental Figures

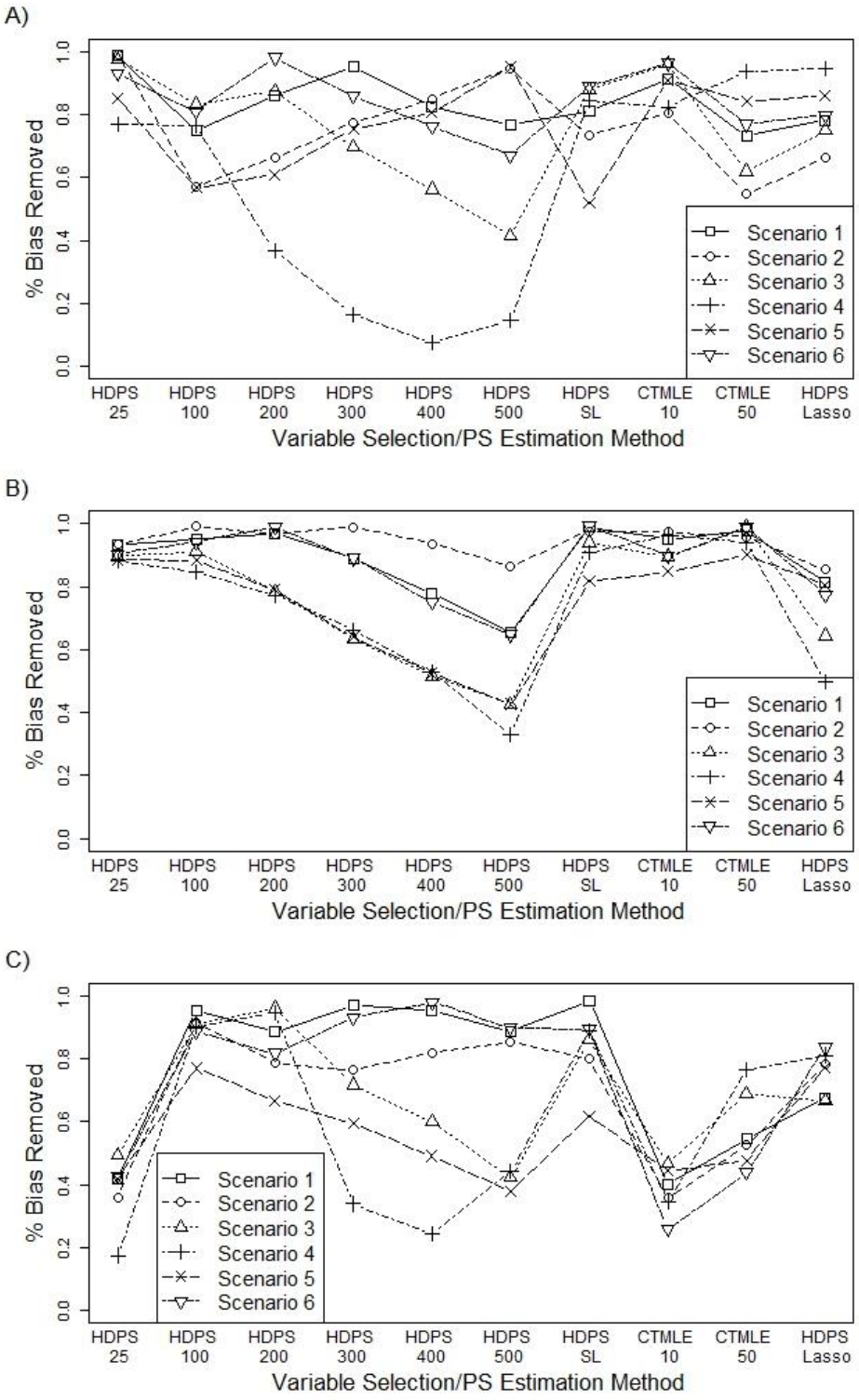
Appendix 4: Supplemental Figures



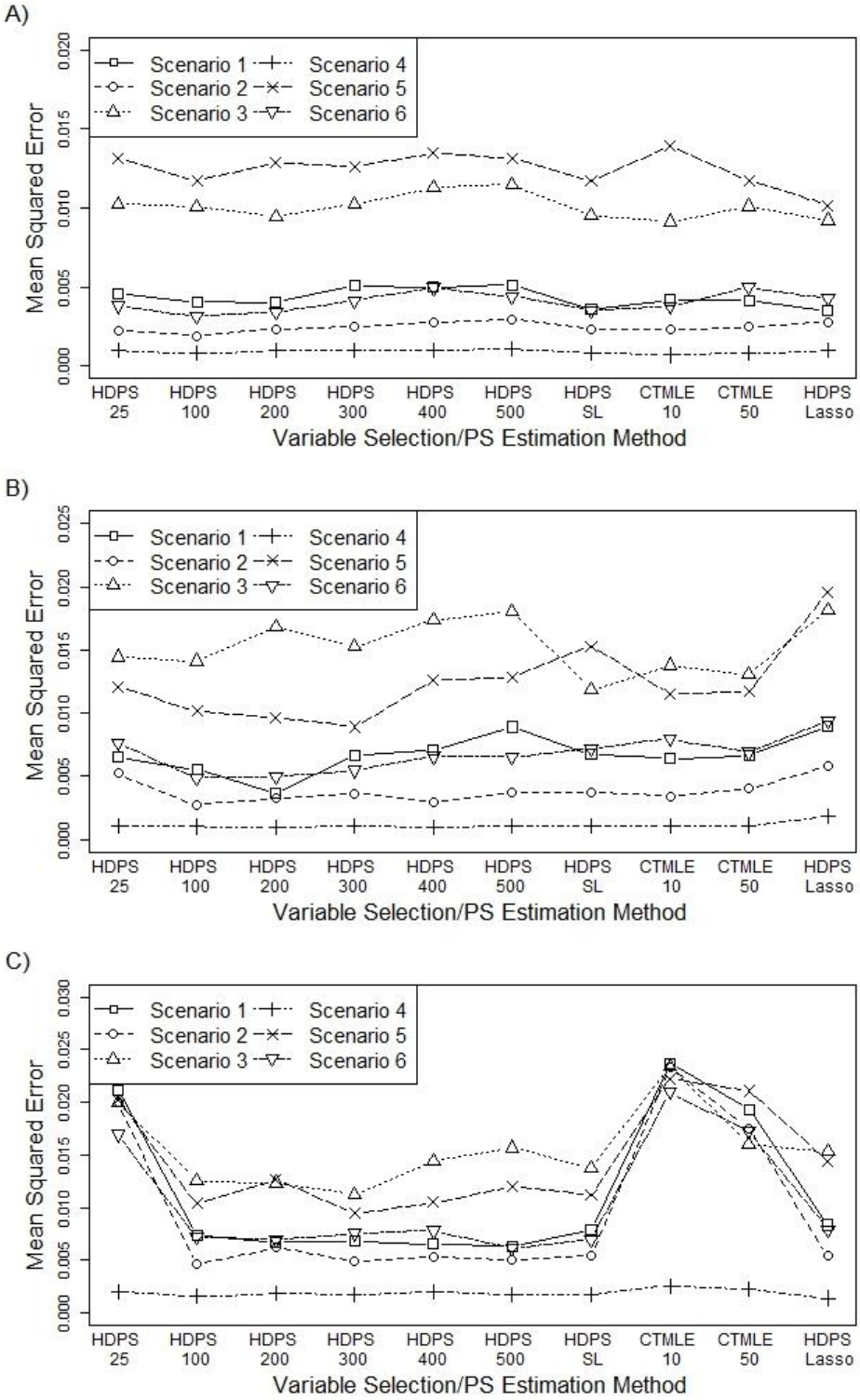
Appendix Figure 1. Percent bias removed for each scenario and variable selection method when matching on the estimated propensity scores. Plots A, B, and C show results for plasmode simulations based on the NSAID, NOAC, and Statin datasets, respectively.



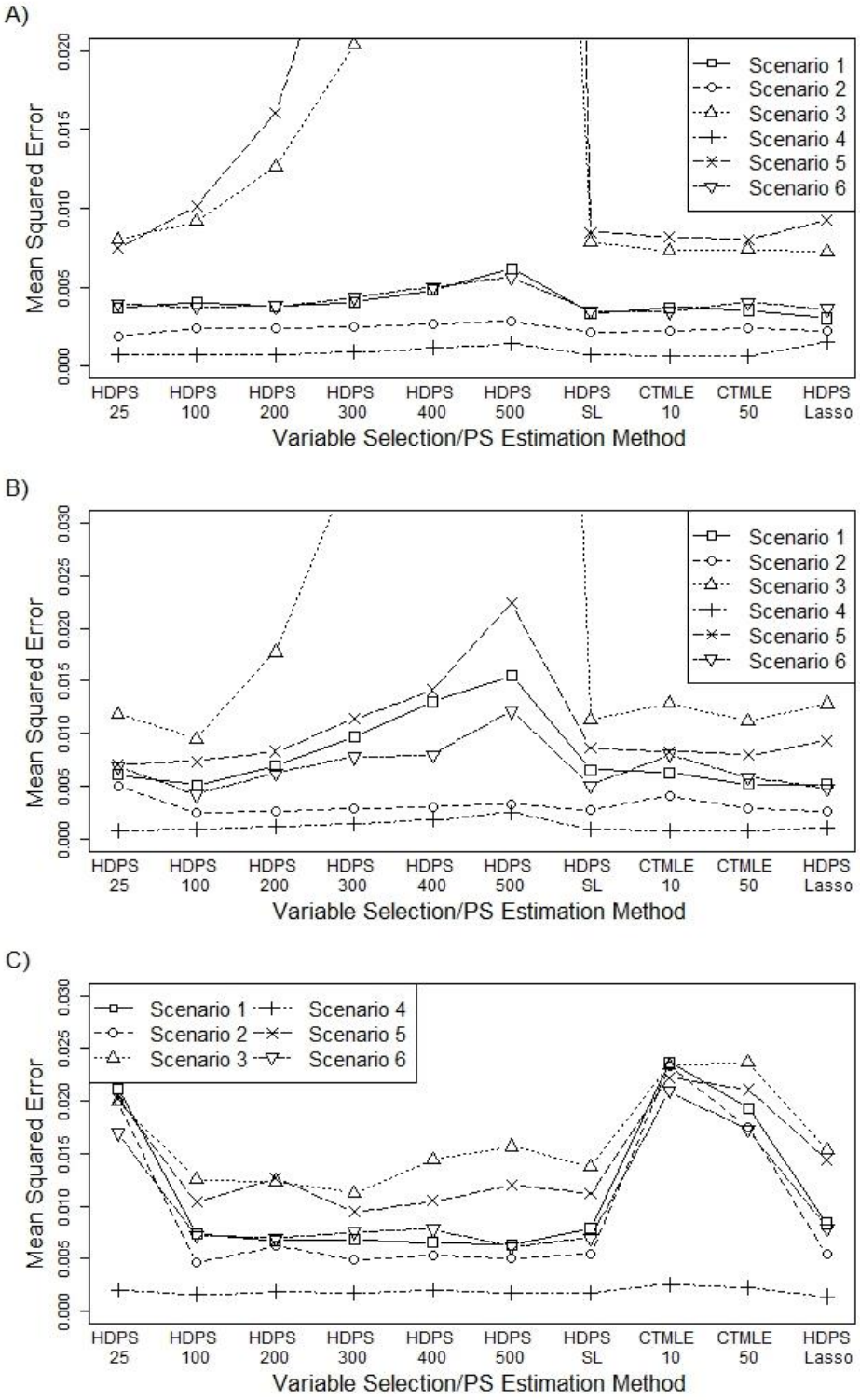
Appendix Figure 2. Percent bias removed for each scenario and variable selection method when using IPTW to implement the estimated propensity scores. Plots A, B, and C show results for plasmode simulations based on the NSAID, NOAC, and Statin datasets, respectively.



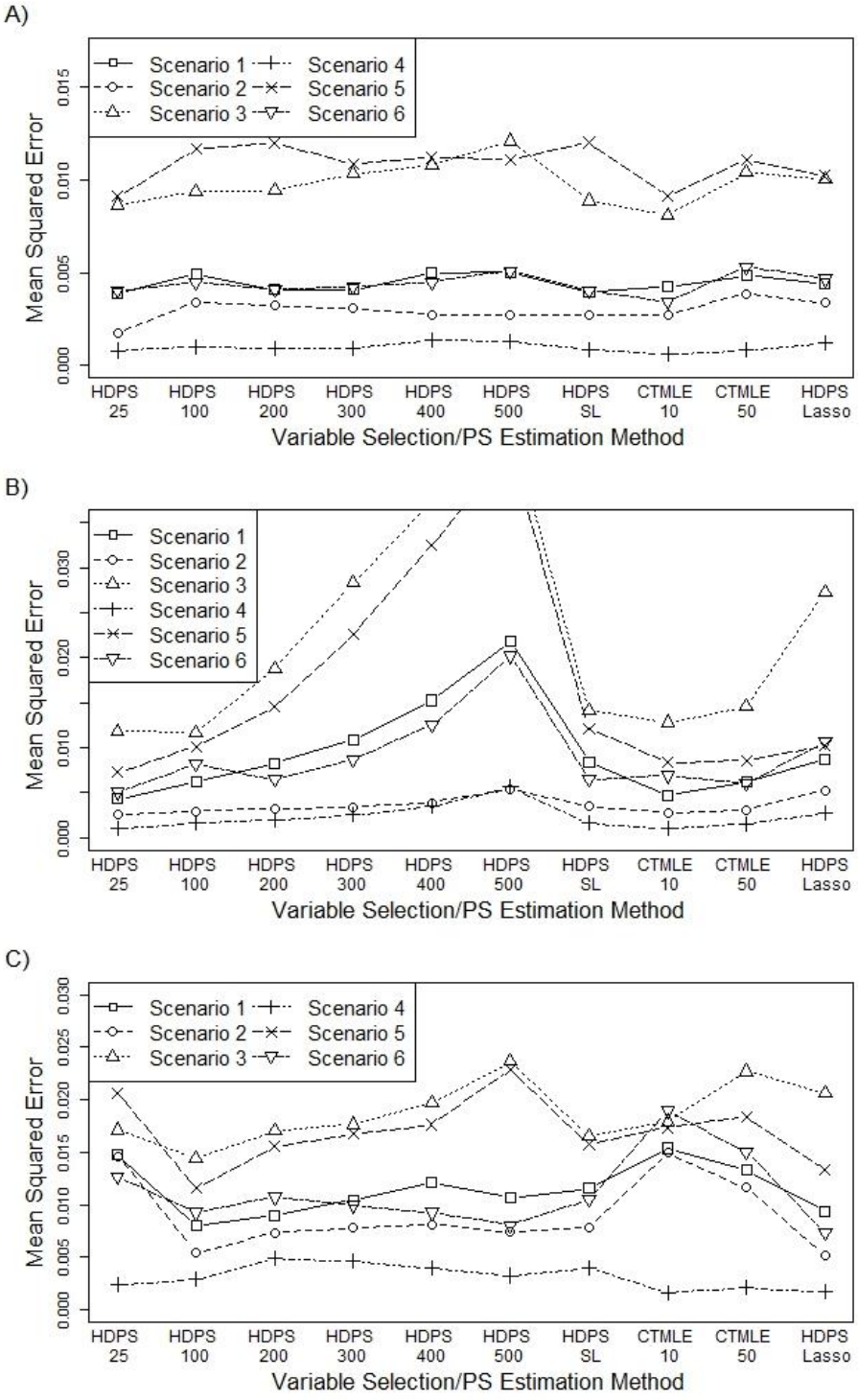
Appendix Figure 3. Percent bias removed for each scenario and variable selection method when using TMLE to implement the estimated propensity scores. Plots A, B, and C show results for plasmode simulations based on the NSAID, NOAC, and Statin datasets, respectively.



Appendix Figure 4. Mean squared error (MSE) for each scenario and variable selection method when matching on the estimated propensity scores. Plots A, B, and C show results for plasmode simulations based on the NSAID, NOAC, and Vytorin datasets, respectively.



Appendix Figure 5. Mean squared error (MSE) for each scenario and variable selection method when using IPTW to implement the estimated propensity scores. Plots A, B, and C show results for plasmode simulations based on the NSAID, NOAC, and Vytorin datasets, respectively.



Appendix Figure 6. Mean squared error (MSE) for each scenario and variable selection method when using TMLE to implement the estimated propensity scores. Plots A, B, and C show results for plasmode simulations based on the NSAID, NOAC, and Vytorin datasets, respectively.