4. I'd like to see more information regarding the the propensity score estimates, since this is where you are getting differing results from the paper. Again, you could create a 3D plot of the "true" propensity score and compare it to your linear and non-parametric estimate. If you need, you could see if you can get similar results with a single covariate. At which point it becomes a simple 2D plots.

Up to now

Two more things to do:

1. Run the naïve bootstrap
2. Run BootstrapDGP0 which has only one covariate and use log p/1-p both for kernel and liner specification
3. Estimation error of linear specification