Hello Reader,

For figure2D. The .r file must be run on the RStudio Application. A different R code processing software may be substituted for RStudio instead.

There is one .pdb files that must be downloaded along with its respective script for Figure 4 titled:

lig.pdb (inhibitor docked to apo 7LV7)

For the scripts to run, ensure the .pdb file is in the same directory as the script