

# ptraj scripts

Common ptraj scripts:

- [hbond.in](#) - A script used with RNA (and modified RNA residues) for hydrogen bond analysis.
- [grid.in](#) - A script to create a grid of average water density in X-Plor format (viewable in Chimera, VMD) with the correct offset for the respective avg.pdb file.
- [combined-clustering.in](#) - Example of combined clustering commands with multiple trajectories (same topology)