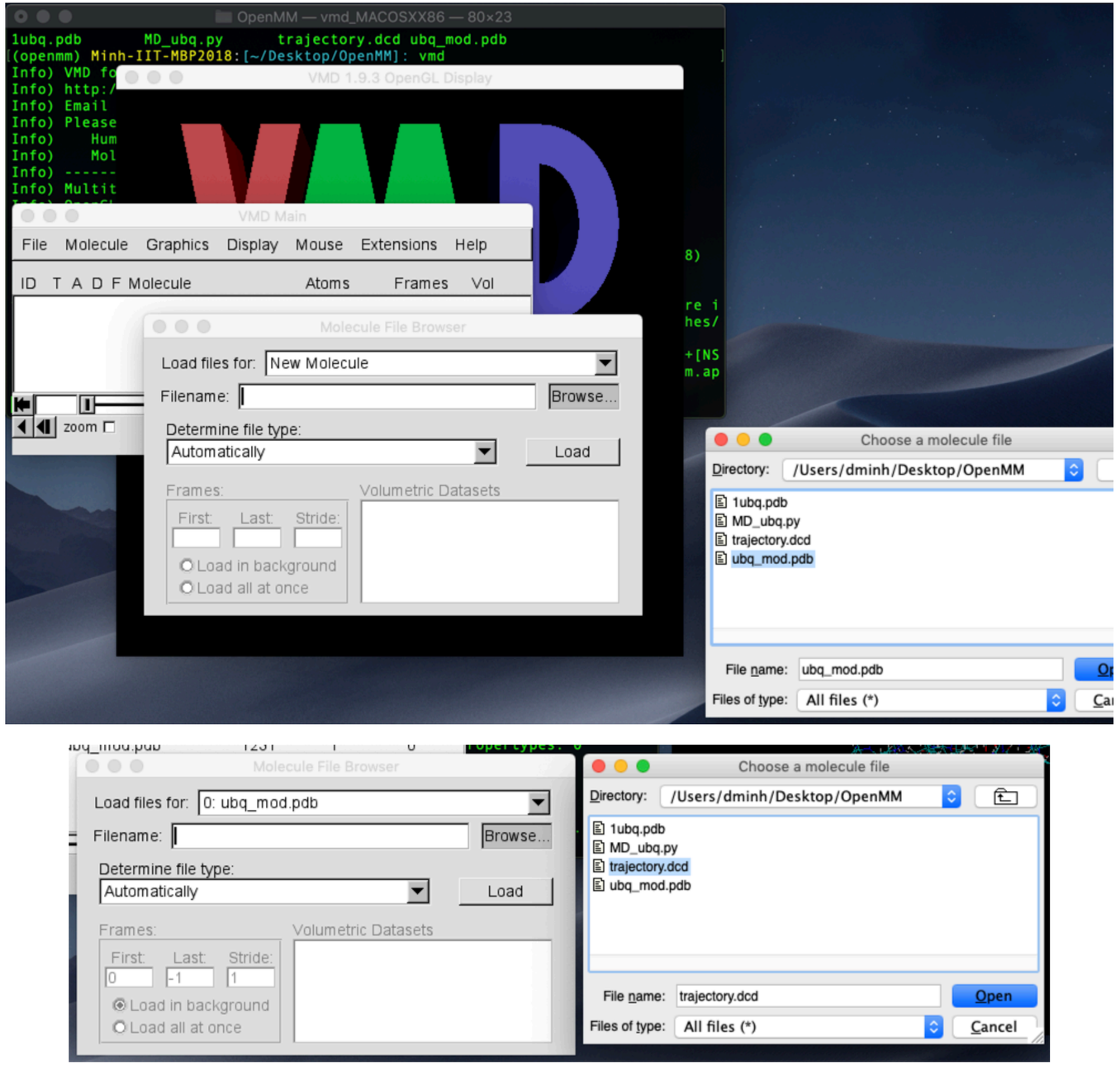
- To visualize the molecular dynamics trajectory
  - open vmd
  - create a new molecule based on “ubq_mod.pdb”
  - load files “trajectory.dcd” as a file for the new molecule



- Now look at the “VMD Main” window
  - “Frames” reads 11 instead of 1
  - You can use the bottom of the window to start a movie and control where we are
- There probably won’t be much going on in your movie, but all of molecular dynamics is just like this, just rinse and repeat