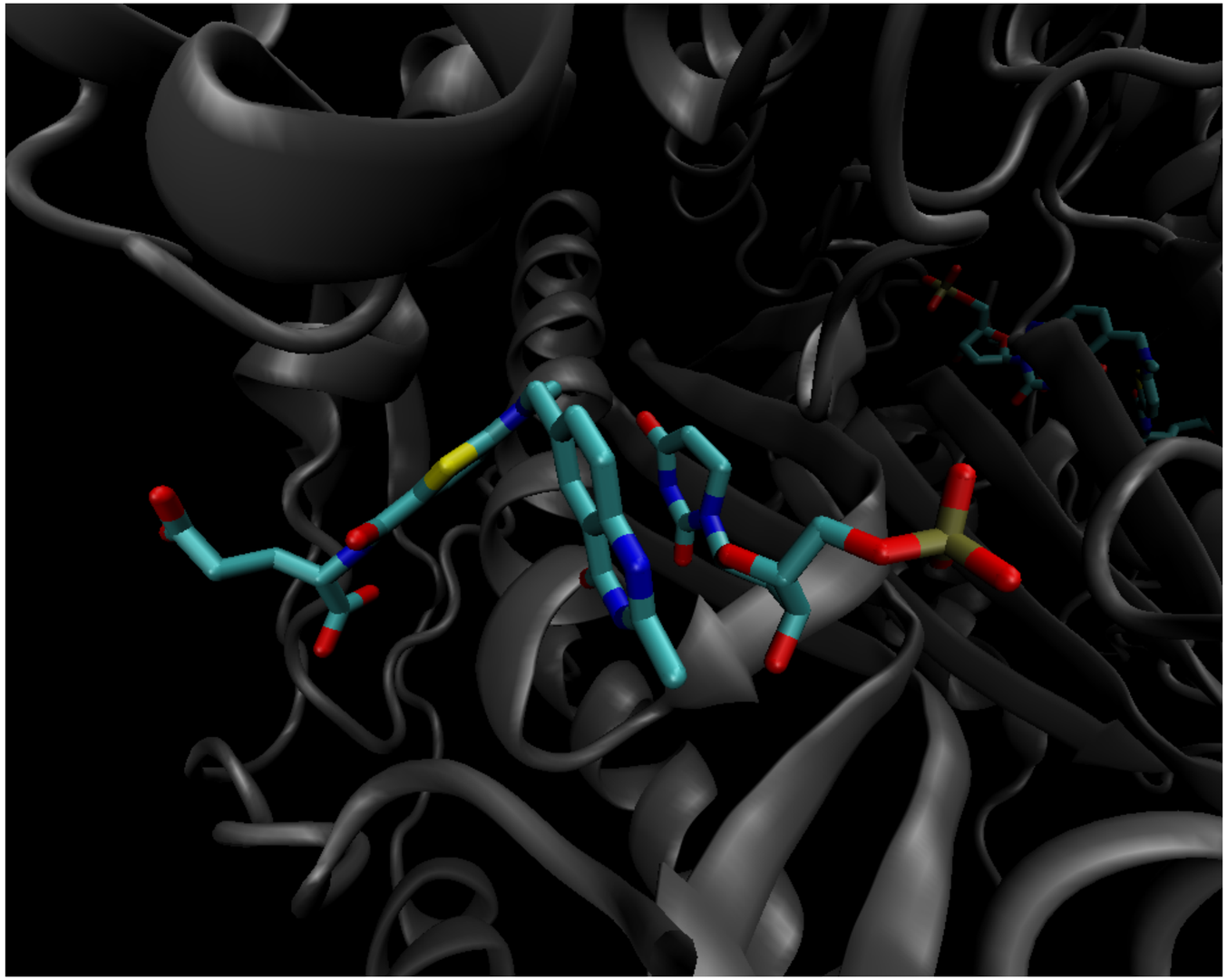
- The location of the active site is a lot more evident if you create a representation with Selected Atoms as “resname D16 or resname UMP”, Color Method “Element”, and Drawing Method “Licorice”



- Now let's actually load the electrostatic potential
- In the "VMD Main" window, right click on a molecule and select "Load Data Into Molecule"
- Browse to the .dx file in the same directory as the molecule of interest and load it
- It won't look like anything has happened except there will be a line under "Volumetric Datasets"
- Let's focus on human and hide the other structures by double-clinking on D in the VMD Main window