

Installing YANK

- If you want to run YANK on Bridges, you need to install it in your directory
- For detailed instructions, see <http://getyank.org/latest/installation.html>
- In short, once you have anaconda/miniconda installed, you can type
 - `conda config --add channels omnia --add channels conda-forge`
 - `conda create --name yank yank`
 - `conda update --all`

Receptor setup

- The YANK workflow requests a PDB file for the receptor
- You have already done similar steps
- I prepared the receptor file by
 - submitting the structure 6LU7 to the PDB2PQR server
 - removing the ligand and water from the main result, 6LU7.pqr
 - results of these first steps are here: 3cl-pro/YANK/receptor
 - copying 6LU7.pqr to 6LU7.pdb to have a file name accepted by the YANK workflow