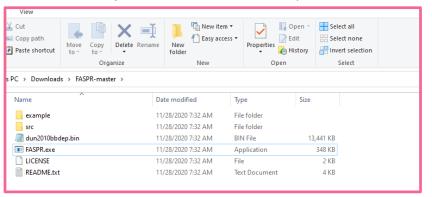
THIS INSTRUCTION APPLIES TO WINDOWS SUBSYSTEM FOR LINUX

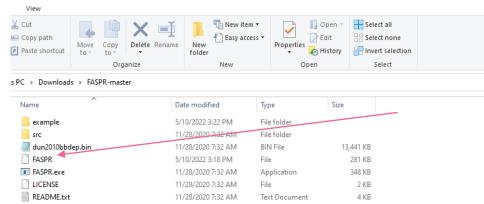




Download FASPR from https://github.com/tommyhuangthu/FASPR and extract to any folder as below. Then enter/open the folder



After successful compilation you will see FASPR as pointed towards



In FASPR.cpp, line 72, change string pdbin=(string)"example/1mol.pdb" to string pdbin=(string)"." In line 73, change string pdbout=(string)"example/1mol_FASPR.pdb" to string pdbout=(string)"."

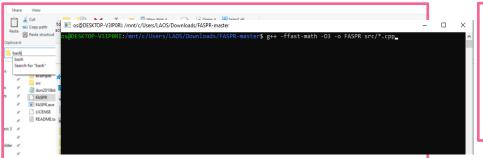
Without these changes, your input PDB and its path must correspond to example/1 mol.pdb





Type "bash" in the search window and hit enter to open lunix terminal as shown. Then, execute the code below to compile FASPR (it takes ≈ 1 min. to compile):

g++ -ffast-math -O3 -o FASPR src/*.cpp



Now install PRAS server (if not installed) using the command below

Install PRAS server which is added to path automatically with:

pip install Pras-Server==1.2.1

Now download 1aho.pdb and copy to the current folder and run the code in syntax.txt that invokes linux_examply.py in the folder containing this document to repair several missing atoms using chi from FASPR PDB output. It is a good practice to type commands in Linux than copy and paste as Linux is extremely sensitive.