



XChem Directory Structure & Linux commands

2024



Working directory



You will have a proposal number starting with lb, e.g.:

- lb13385
- **For each target/screen you will have a visit number, e.g.:**
 - lb13385-1
- **You will end up with visits assigned to both:**
 - Lab34: labxchem
 - The beamline: I04-1
- **Your first labxchem visit will be the working directory for the duration of the project**
 - Other lab visits will be created to trigger lab access
 - Other beamline visits will be created for data collection

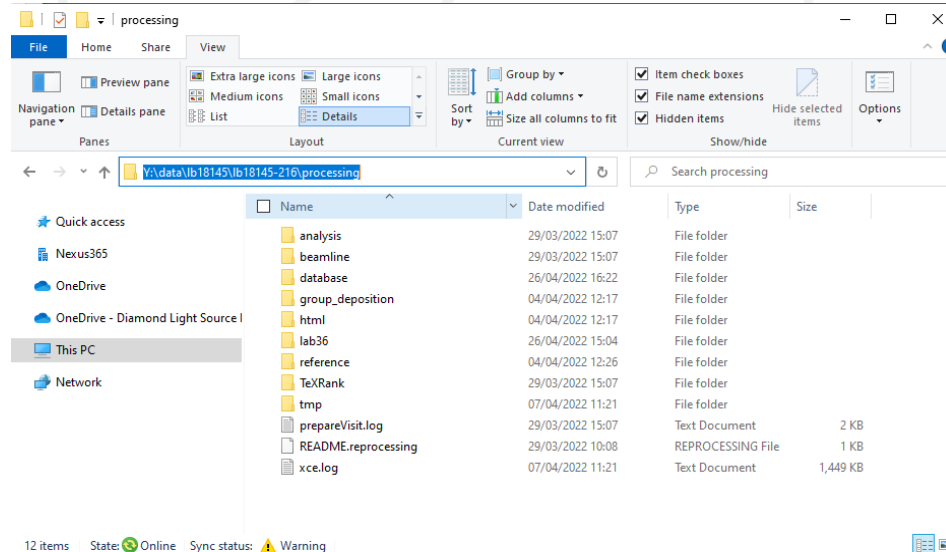
Linux and windows



- From linux, the location is accessed via:
 - `/dls/labxchem/data/proposal/visit/processing/`
- You (or your local contact) will create your subfolders in visit-1 by running:
 - `cd /dls/labxchem/data/proposal/visit/processing/`
 - `preparevisit`
- From windows (lab PCs mainly), you will have access to your visit -1 working directory via:
 - `Y:\data\proposal\visit\processing\`

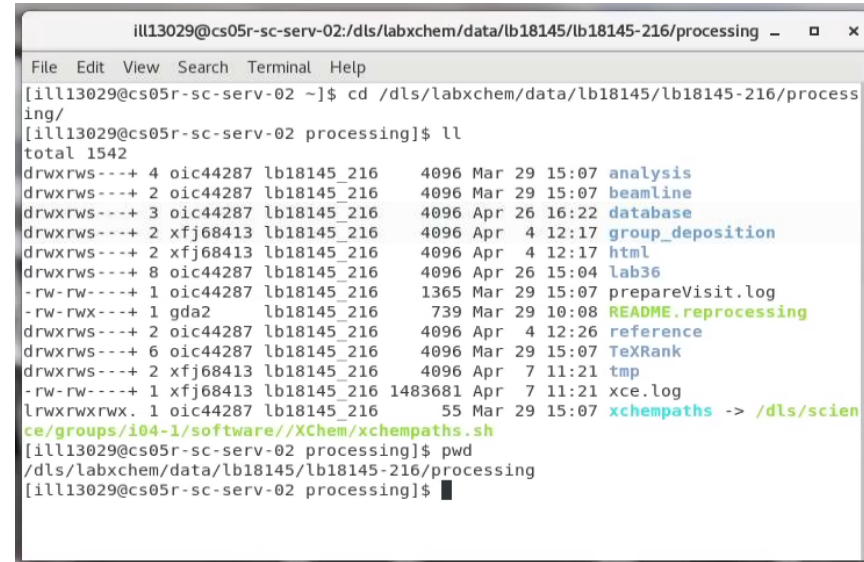
Windows world:

`Y:\data\lb18145\lb18145-216\processing`



Linux world:









`/dls/labxchem/data/lb18145/lb18145-216/processing`



Working directory structure

- analysis • Data collection results and pandda analysis
- beamline • Links to beamline visit directories (*obsolete – do not use*)
- database • SQLite datafile (and backups)
- group_deposition • Files for PDB deposition
- html
- lab36 • Directory for lab work (soakDB, echo, shifter)
- reference • pdb of a good reference model
- tmp
- prepareVisit.log
- xce.log

Lab 36 directory structure:

-  crystal-targets • Crystal target lists output from TeXRank
-  echo • csv files for echo soaking
-  ispyb • csv files for upload to ispyb
-  shifter • csv files for the shifter
-  soakDB.bat • OldsoakDB launcher (***obsolete – do not use***)
-  soakDB3.bat • Current soakDB launcher (use this one!)
-  sqlite3.dll
-  SQLite3_StdCall.dll

Useful linux commands



- Setup useful commands (**do this first**):
 - `cd /dls/labxchem/data/proposal/visit/processing/`
 - `source /dls/science/groups/i04-1/software/XChem/xchempaths.sh`
- `xchempaths.sh` will set paths for these commands:
 - `preparevisit` - to create the subfolders needed for XChem
 - `tserver` - to launch a windows remote desktop from linux
 - `xce` - to launch XChemExplorer
 - **Needs to be run under the 'processing' folder**
 - `csv2ispyb` - to automatically load the data collection information in iSPyB
 - **Needs to be run under the 'processing' folder**
- Checking the status of the cluster (type into terminal):
 - “`ssh wilson`” – connect to the DLS Slurm Cluster; must be done first
 - “`sacct`” – display jobs
 - “`watch sq.sh -u yourfedid -nf`” – watch jobs