

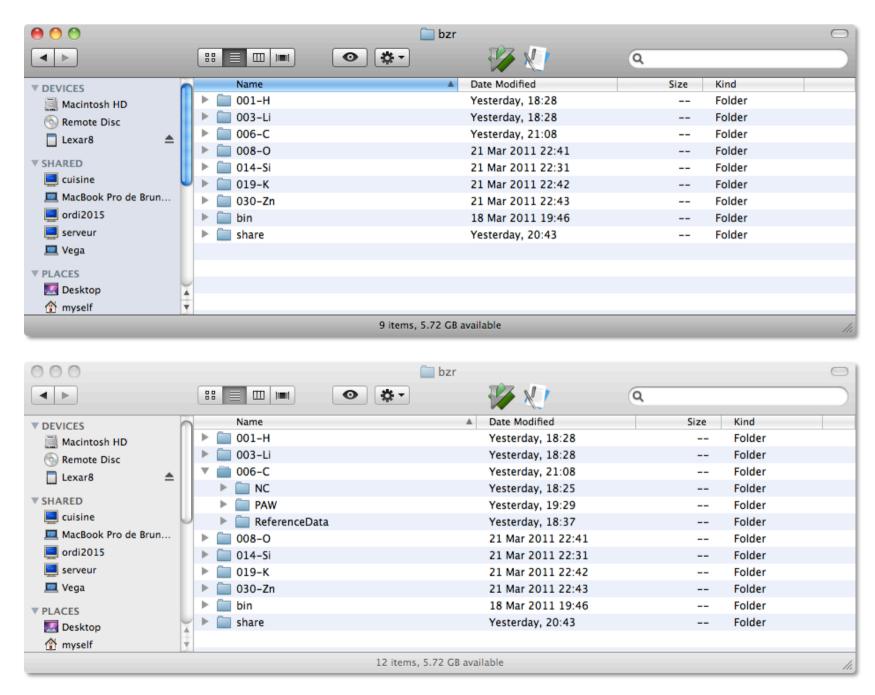
abinit/psps project

Access to the repositories ... the usual location:

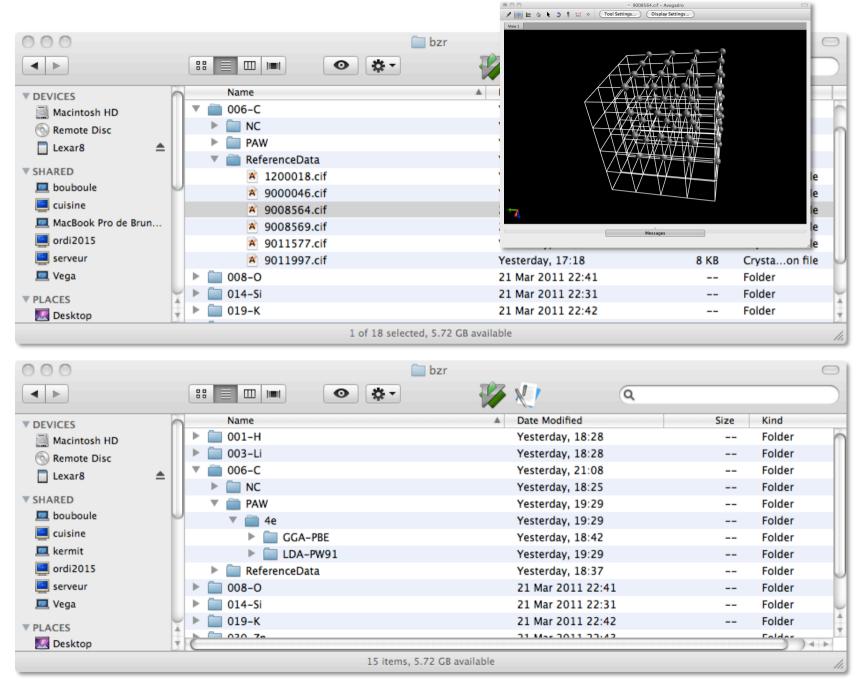
bzr+ssh://psps@archives.abinit.org/forge/psps/xxx-AtomName

xxx is 001-H to 112-Cn; one project per atom repositories have one username – psps but use existing Abinit ssh keys

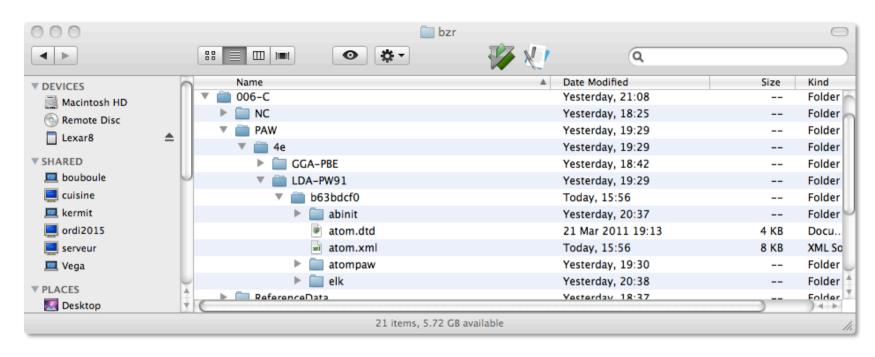












UUID

Two-liner python code

>>> import uuid

>>> uuid.uuid1()

Conflicts between contributions

bzr merge before bzr push



```
atom.xml
<?xml version="1.0" encoding="UTF-8"?>
<atom version="1.0">
<dict>
       <key>DatasetName</key>
       <string>b63bdcf0-6391-11e0-b8af-5855caf38db7</string>
       <key>AtomicNumber</key>
       <integer>6</integer>
       <key>AtomName</key>
       <string>C</string>
       <key>XCFunctional</key>
       <dict>
              <key>ID</key>
              <integer>7</integer>
              <key>Name</key>
              <string>LDA-PW</string>
       </dict>
       <key>ScalarRelativistic</key>
       <true/>
       <key>GridSize</key>
       <integer>801</integer>
```



```
<key>CoreElectrons</key>
<integer>2</integer>
<key>ValenceElectrons</key>
<integer>4</integer>
<key>ElectronicConfiguration</key>
<dict>
      <key>1s</key>
      <dict>
              <key>State</key>
              <array>
                     <integer>1</integer>
                     <integer>0</integer>
              </array>
              <key>Occupation</key>
              <real>2.0</real>
              <key>Valence</key>
              <false/>
      </dict>
```

•••



```
<key>PartialWaveBasis</key>
<dict>
      <key>MaximumLQuantumNumber</key>
      <integer>1</integer>
      <key>PartialWave</key>
      <dict>
              <key>State</key>
              <integer>0</integer>
              <key>ReferenceEnergy</key>
              <array>
              </array>
              <key>Unit</key>
              <string>Rydberg</string>
      </dict>
      <key>PartialWave</key>
      <dict>
              <key>State</key>
              <integer>1</integer>
              <key>ReferenceEnergy</key>
              <array>
              </array>
              <key>Unit</key>
              <string>Rydberg</string>
```

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```
<key>DataGenerator</key>
<dict>
      <key>Name</key>
      <string>atompaw</string>
      <key>Version</key>
      <string>3.0.1</string>
</dict>
<key>DataConvertor</key>
<dict>
      <key>Name</key>
      <string>AtomPAW2Abinit</string>
      <key>Version</key>
      <string></string>
</dict>
<key>DataValidator 2</key>
<dict>
      <key>Name</key>
      <string>elk</string>
      <key>Version</key>
      <string>1.2.15</string>
</dict>
```





Working directories atompaw – abinit – elk

Naming conventions:

input files i.e. atompaw.in, abinit.in, elk.in output files i.e. atompaw.log, abinit,log

Validation of the PAW pseudopotentials follows guidelines of PAW3 tutorial, comparison between *Abinit* and an all -electron FP-LAPW code – Elk on standardized systems: elemental solid, dimer, oxides, GS and GW

Open questions

- how to have a systematic approach on the whole table?
- robust submission engine how to check for completeness?
- efficient validation routine how to rebuild the database?