Automatic archiving occurs upon completion of zg. Usually the archived dataset (that on castor) will have raw
data (fid or ser) only, and no spectrum exists on castor. IconNMR does process prior to archiving.

last update: 30 Apr 2014 (cgf)

- → Intermediate processing using: **tr** → **efp** → will create a processed spectrum of the data, but with only the number of scans from the **tr** command. This often poor spectrum (e.g., showing just NS=32 of the 640 actually completed) will be copied during auto-archiving. <u>Process the **fid** or **ser** file, not the spectrum file (1r or 2rr)!</u>
- → Some acquisition types will not auto-archive at all: **multizg** and **multi\_zgvd** are two; others may exist.
- $\rightarrow$  To use processing done on the *spectrometer* after **zg** completion, copy the data folder from *spect* to castor.
- On the spectrometer, open the file browser [in the Linux menu: **Applications** → **System Tools** → **File Browser**] and make 2 bookmarks:

spect pointing to /home/username/persephone
castor pointing to /nmrdata/username/public\_html/persephone

