#### Presurgical MRI data processing

## 1. Open the following location on NEF:

sftp://nef-devel/data/athena/user/monno/connectc/

## **2. Create a directory for the new Patient on NEF**. We will call it Patient 1:

sftp://nef-devel/data/athena/user/monno/connectc/patient01

Note that the directory name can be composed of letters, digits, or the '-' character for a separation. Don't use other characters.

Then, create a subdirectory for storing DICOM files:

sftp://nef-devel/data/athena/user/monno/connectc/patient01/dicom

# **3. Copy DICOM images from CD disks to NEF**, respecting the structure of directories as in the example below:

```
sftp://nef-devel/data/athena/user/monno/connectc/patient01/dicom/cd1/IMAGES/sftp://nef-devel/data/athena/user/monno/connectc/patient01/dicom/cd2/IMAGES/sftp://nef-devel/data/athena/user/monno/connectc/patient01/dicom/cd3/IMAGES/
```

Note that the "IMAGES" directory appears in the root directory of every disk recorded by the MRI lab at CHU Nice. It is the only directory that we need.

## **4.** Copy all .sh files from the 'scripts' directory located here:

sftp://nef-devel/data/athena/user/monno/connectc/scripts/

to our new patient's directory:

sftp://nef-devel/data/athena/user/monno/connectc/patient01/

#### 5. Login to NEF from the terminal:

ssh nef-devel

#### 6. Go to our patient's directory:

cd /data/athena/user/monno/connectc/patient01



./run convert.sh

Running time is 1-2 days.

**8. Use MI-Brain to virtually dissect the interesting fibers.** Save them in the .TCK format and place in the following location on NEF:

sftp://nef-devel/data/athena/user/monno/connectc/patient01/tracts

## 9. Run the script to create the DICOM images for the neuronavigation system:

./run\_neuronav.sh

Running time is about 15 minutes.

**10.** Copy the "neuronav" directory to the USB key and load it on the neuronavigation system. The directory can be found on NEF here:

sftp://nef-devel/data/athena/user/monno/connectc/patient01/neuronav