# XTOLM use case - compute brain volumes from an MRI experiment and export data to spreadsheet

In this example, we will use XTOLM use case - compute brain volumes from an MRI experiment and export data to spreadsheet to compute whole brain volume using the Brain Extraction Tool for all MPRAGE scans in an MRI project in XNAT, and save the resulting statistics in a spreadsheet.

## Prerequisites

- FSL and dcm2nii on your path
- · We'll use data from surfmask\_smpl project on XNAT central so you'll need an XNAT Central account.

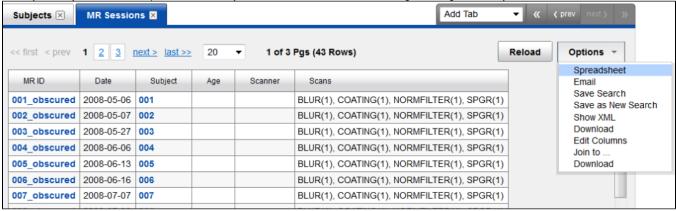
# Step by Step Instructions

1. Write a bash script that computes human brain volume from a T1-weighted MR image in DICOM format (assuming FSL v4+ is on the path).

```
#!/bin/bash
in=$1 #input DICOM directory
dcm2nii $in
cp $in/[0-9]*.nii head.nii
if [ ! -f "head.nii" ]; then exit -1; fi
bet head brain -m
vol=(`fslstats brain_mask -V`); vol=${vol[1]}
#report the volume in mm3 to console.
echo $vol
```

2. Create a CSV or XLSX spreadsheet listing all scans that you want to process.

In many cases, you'll have such spreadsheet; if not, you can start with the one created using the one generated by XNAT:



Now, modify the spreadsheet so that column names don't contain spaces, empty and irrelevant cells are removed, and set "Scans" to "SPGR", the scan type that we want to process for each session. If any fields have trailing zeroes, you'll need to modify the csv outside of Excel to add them.

```
MR_ID,Subject,Scans,
001_obscured,001,SPGR,
002_obscured,002,SPGR,
003_obscured,003,SPGR,
004_obscured,004,SPGR,
005_obscured,005,SPGR,
006_obscured,006,SPGR,
```

3. Adapt the brain\_volume.sh to process all sessions from your CSV file.

## brain\_volume.xt #!/bin/bash #this line will convert test.csv to a bash source test.params that assigns corresponding arrays. load\_data test.csv #as a result, the following arrays are assigned: #MR\_ID=("001\_obscured" "002\_obscured" ... #Subject=("001" "002" "003" ... #Scans=("SPGR" "SPGR" "SPGR" ... #now, iterate over all MR sessions. xt\_for Subject MR\_ID \$1 #load specified scan from XNAT. load type \$Scans #find the scan that was created by load\_type. in=`ls -d study\*` #compute brain volume, similarly to brain\_volume.sh dcm2nii \$in #select the unmodified NIFTI volume to process. cp \$in/[0-9]\*[0-9].nii head.nii if [ ! -f "head.nii" ]; then exit -1; fi bet head brain -m vol=(`fslstats brain\_mask -V`); vol=\${vol[1]} #now we use special command to save a variable with this offline session. save\_vars vol xt\_done #this will go through all sessions and generate a spreadsheet summarizing variables saved with "save\_vars" summary mpr\_vol MR\_ID

#### 4. Test the batch processing.

We will run the developed script on a single session to see if it works. For that, uncomment break statement on line 26. Now we are ready to run the script:

```
running brain_volume.xt

xtolm -sr https://central.xnat.org -u your_xnat_central_user -pr surfmask_smpl -o brain_volume.xt
cat mpr_vol.csv
```

Now we'll have the BET calculations saved in mpr\_vol.csv file ready for analysis (note that Excel removed trailing zeroes from the subject again).

	Α	В	С
1	session	subject	vol
2	001_obscured	1	1376148.75
3	002_obscured	2	1393418
4	003_obscured	3	1470579.75
5	004_obscured	4	1452396.75
6	005_obscured	5	1410483.875
7	006_obscured	6	1709883.75
8	007_obscured	7	1442823.25
9	008_obscured	8	1616453.5