**EI protocol**

**1. File-> New protocol**

(Default anatomy->No, use individual anatomy)

(Default channel file-> no, use one channel file per acquisition run)



**2. New subject**

(Default anatomy->No, use individual anatomy)

(Default channel file-> no, use one channel file per acquisition run)



**3. Pre-implantation MRI**

subject-> import MRI



yes



Click here to compute MNI normalization



maff8



**4. post-implantation MRI**

subject-> import MRI (DICOM or nii)



Yes



ignore



Yes



**5. Generate SPM canonical surfaces**

Run->import->import anatomy->Generate SPM canonical surfaces-> run



Check the results



**6. Link the recordings**

Right-click on the subject folder > Review raw file



**7. Import sEEG locations**

Right-click on the subject folder > add eeg positions-> import from file



yes



**8. Event marking**

Event->add group (onset, baseline)



Right-click on the data-> Add event



**9. export from the data (onset, baseline)**

Right-click on the subject folder > import in database



Right-click on the subject folder > import in database



**10. elipeptogenecity maps**

Process2-> files A: baseline, file B: onset



Run-> epilepsy-> elipeptogenecity maps



Result



When using tutorial data (in Same protocol, only difference is sEEG and location files)



