# Functional Connectivity using iRSFC toolbox

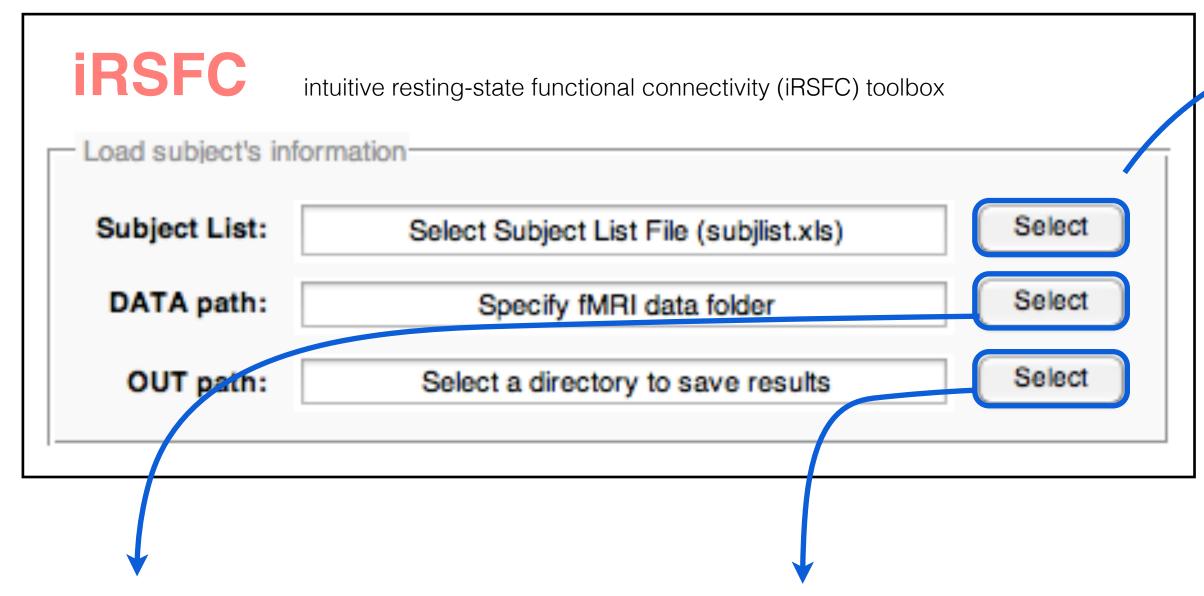
Sunghyon Kyeong, PhD.

Clinical Scientist @ Philips Korea

#### **IRSFC**

intuitive resting-state functional connectivity (iRSFC) toolbox

#### Step 1 - Dataset and Directory



#### **DATA** path

데이터의 위치를 지정한다. 데이터는 다음과 같이 구조로 저장되어 있어야 함.

(DATA path)/NOR001/rest (DATA path)/NOR002/rest

#### **OUT** path

분석결과 및 ROI mask가 저장될 위치를 지정.

#### Subject List

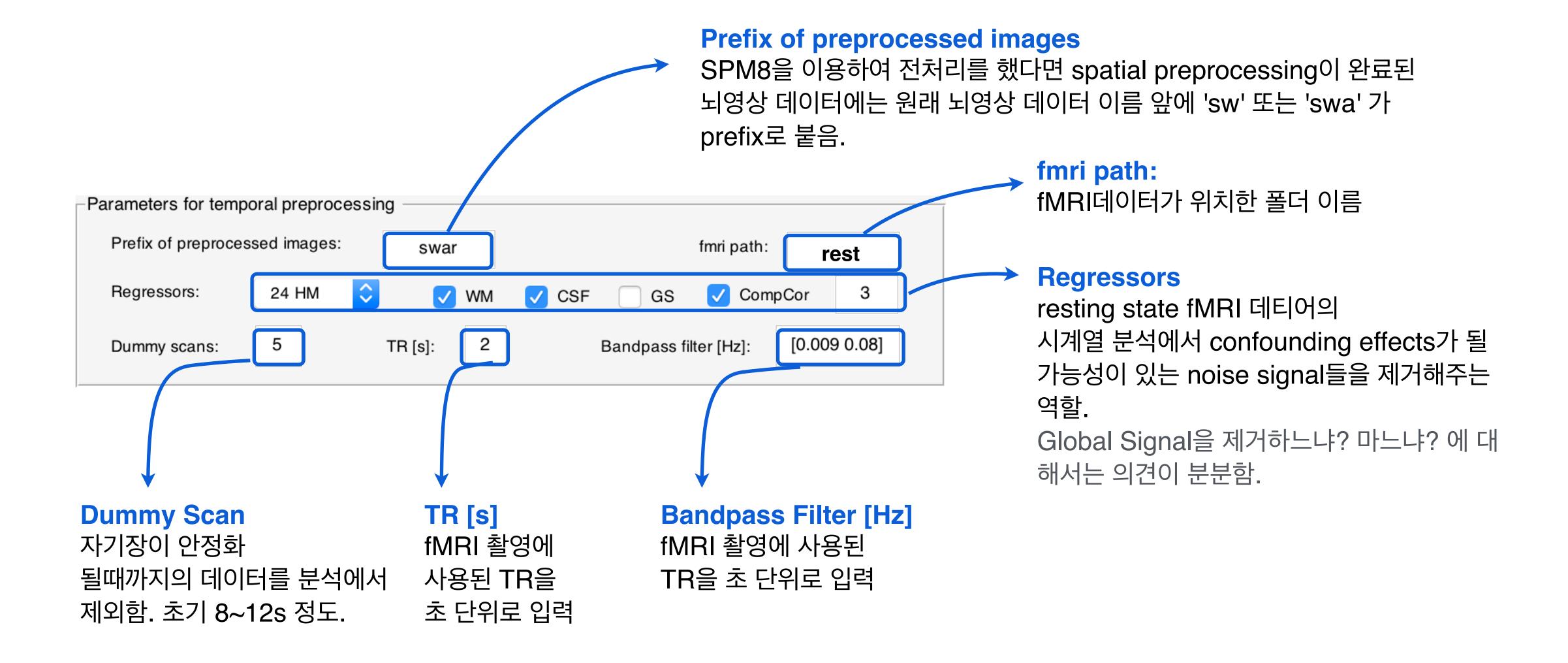
피험자 리스트가 기록되어 있는 엑셀 데이터를 선택.

헤더가 subject 인 컬럼에 폴더명에 해당되는 피험자 이름을 기록.

subjname	Dx	Age	Sex
NOR001	1	20	1
NOR002	1	24	2
NOR003	1	28	1
NOR004	1	22	1
SPR001	2	27	2
SPR002	2	26	2
SPR003	2	20	1
SPR004	2	21	2

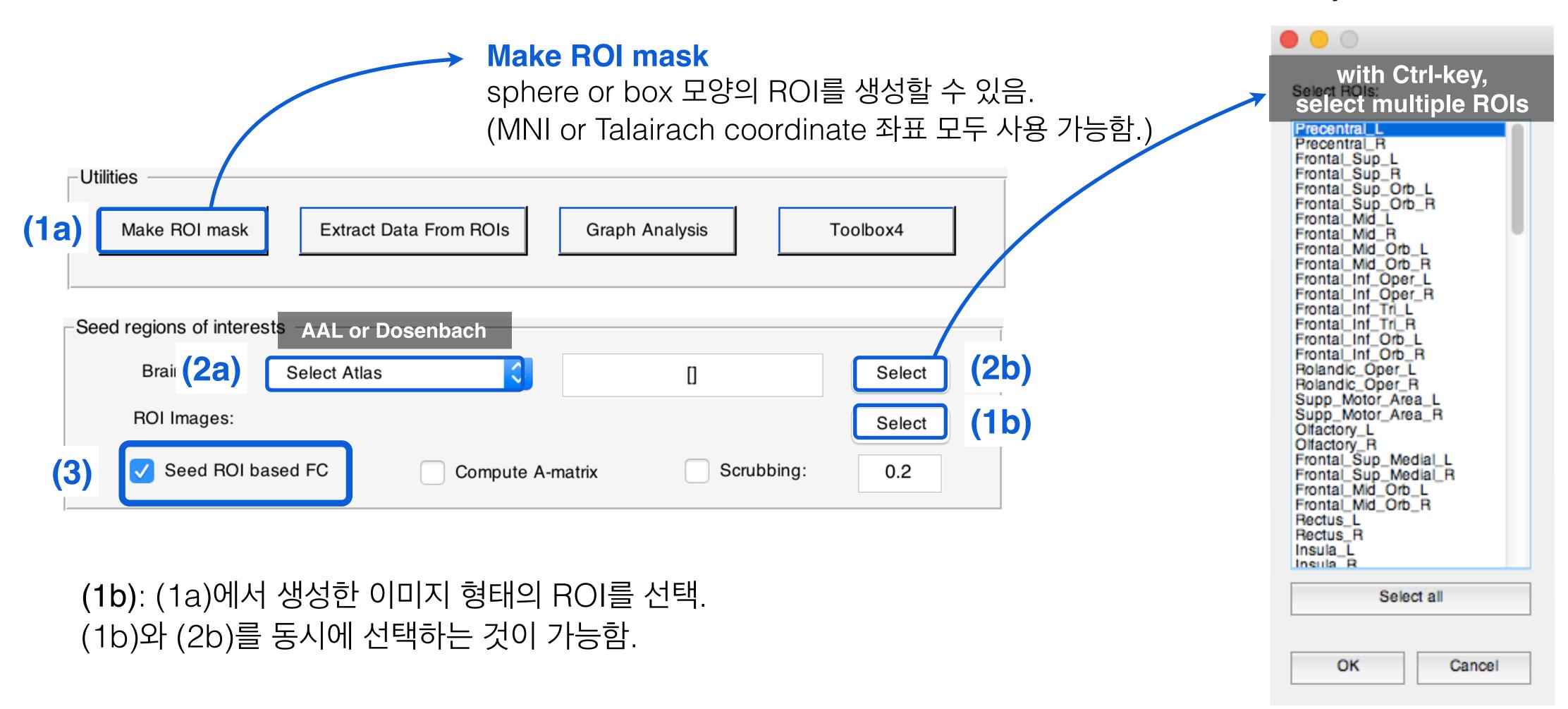
<Subject List 엑셀 파일 예시>

### Step 2 - Filter & Regressors

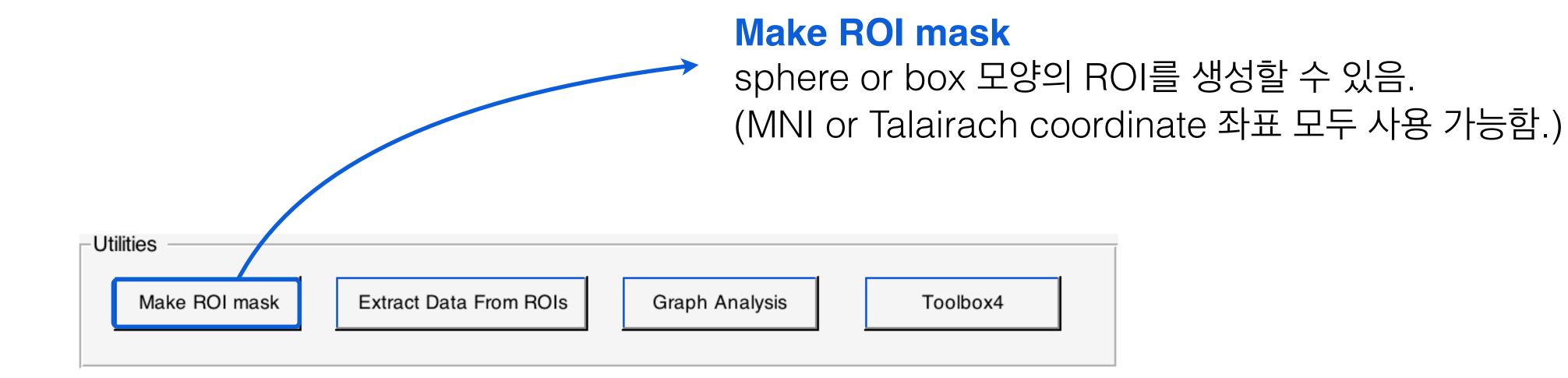


### Step 3 - Select ROIs

: Both atlas based and user defined ROIs can be used for the seed-based FC analysis.



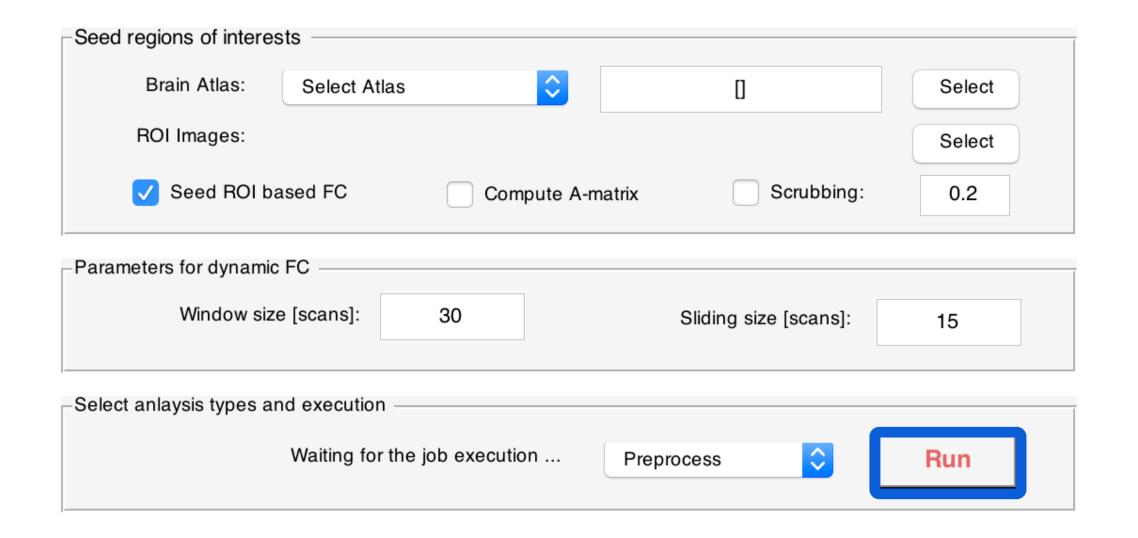
#### Create ROIs



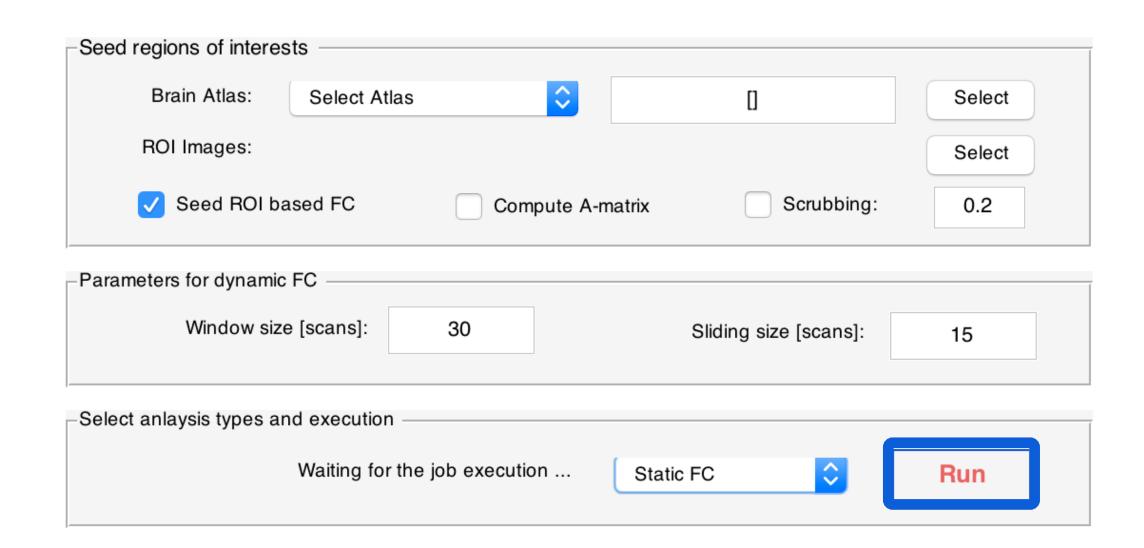
go to slide 19 page

### Step 4 - Run Analysis

#### Temporal preprocessing

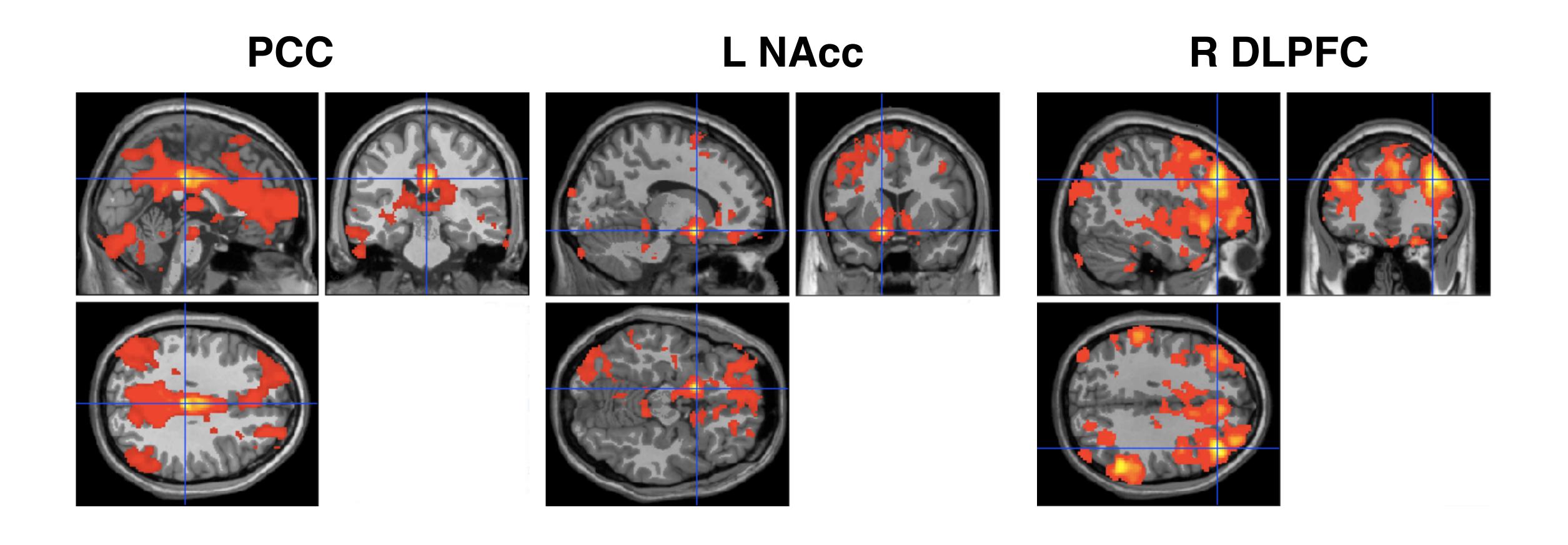


#### Functional connectivity



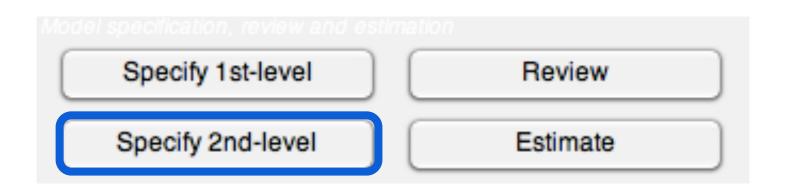
### Now, we have z-maps

for each seed ROI and each subject

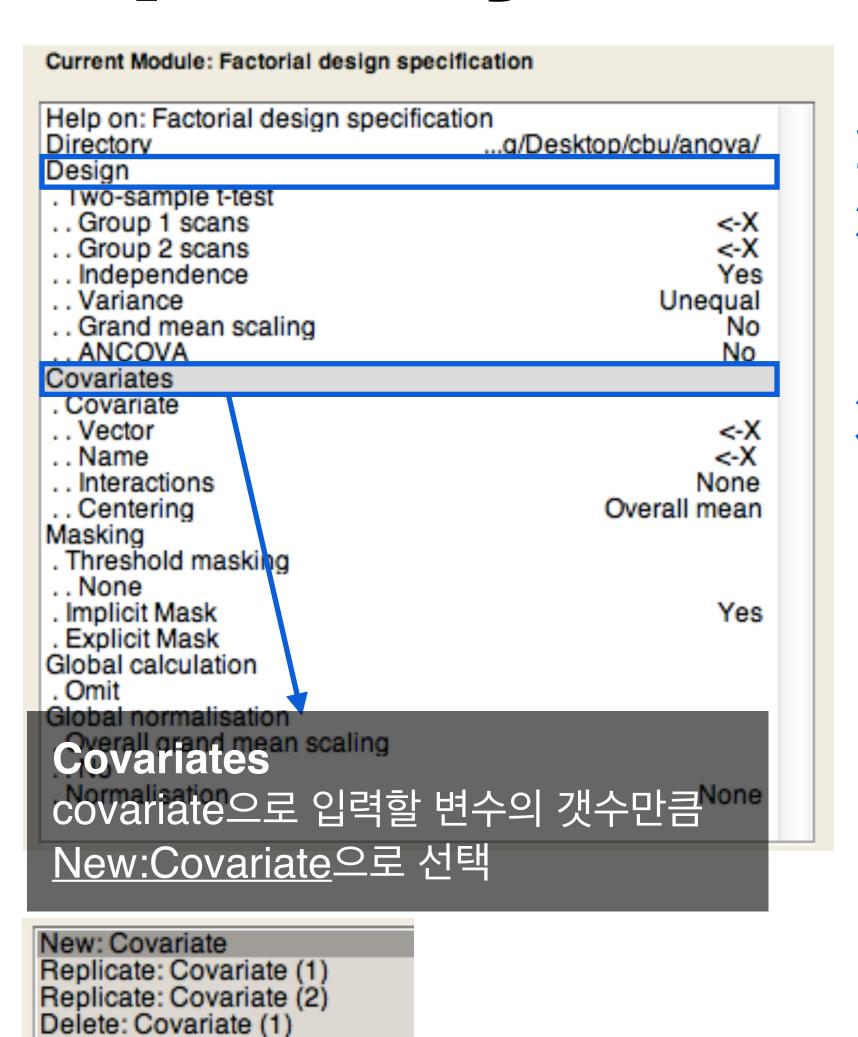


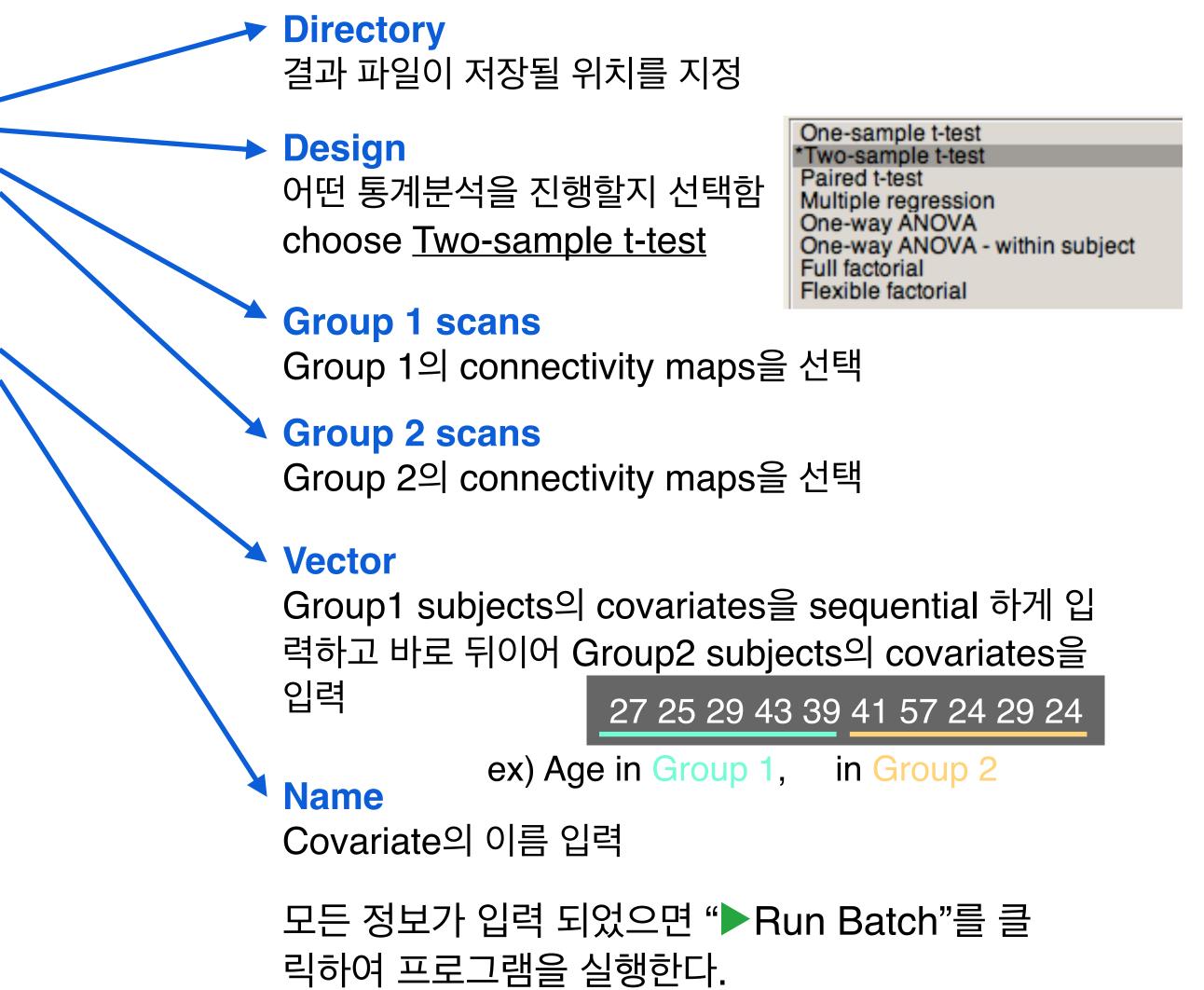
### Two-sample t-test

with covariates



### Specify 2nd-level

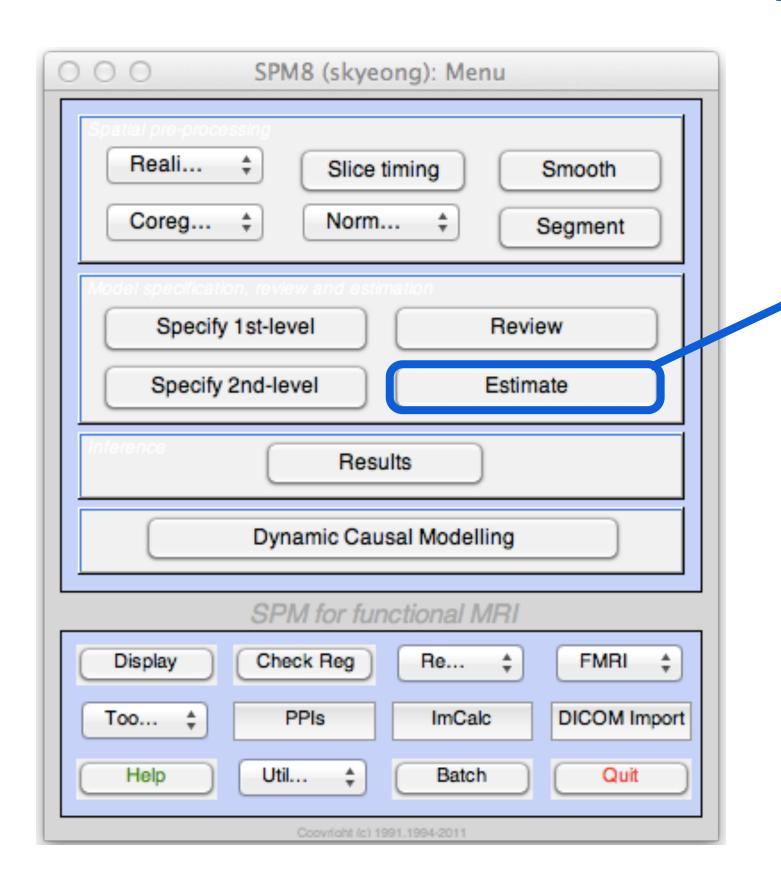


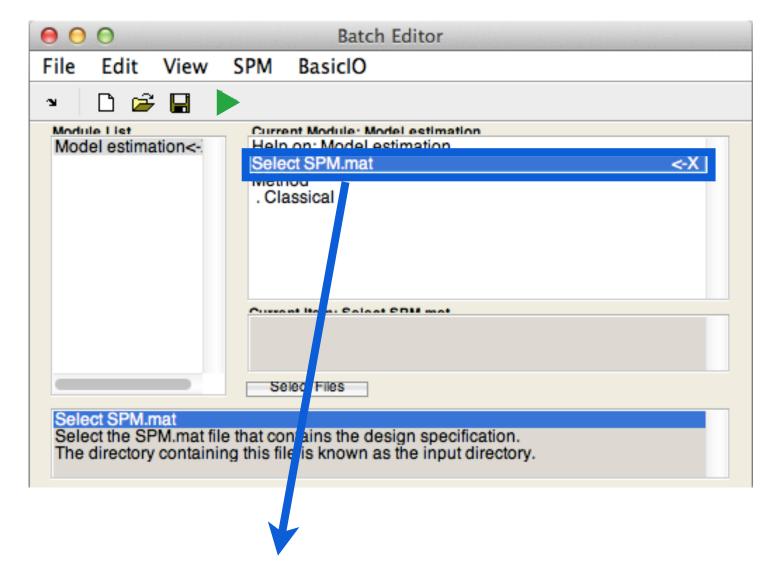


Delete: Covariate (2)

#### Estimation of Parameters

Specify 2nd-level 을 통해서 구성한 General Linear Model의 parameter를 추정(계산)함.



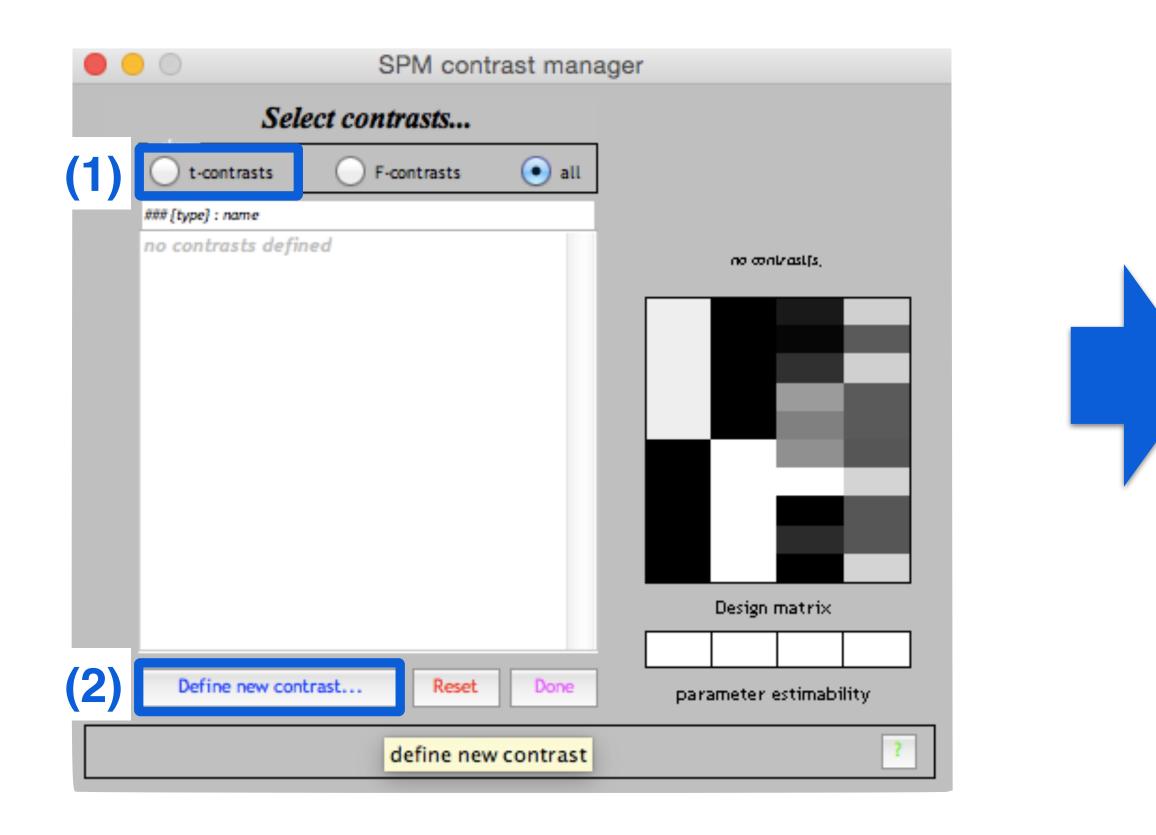


#### **Select SPM.mat**

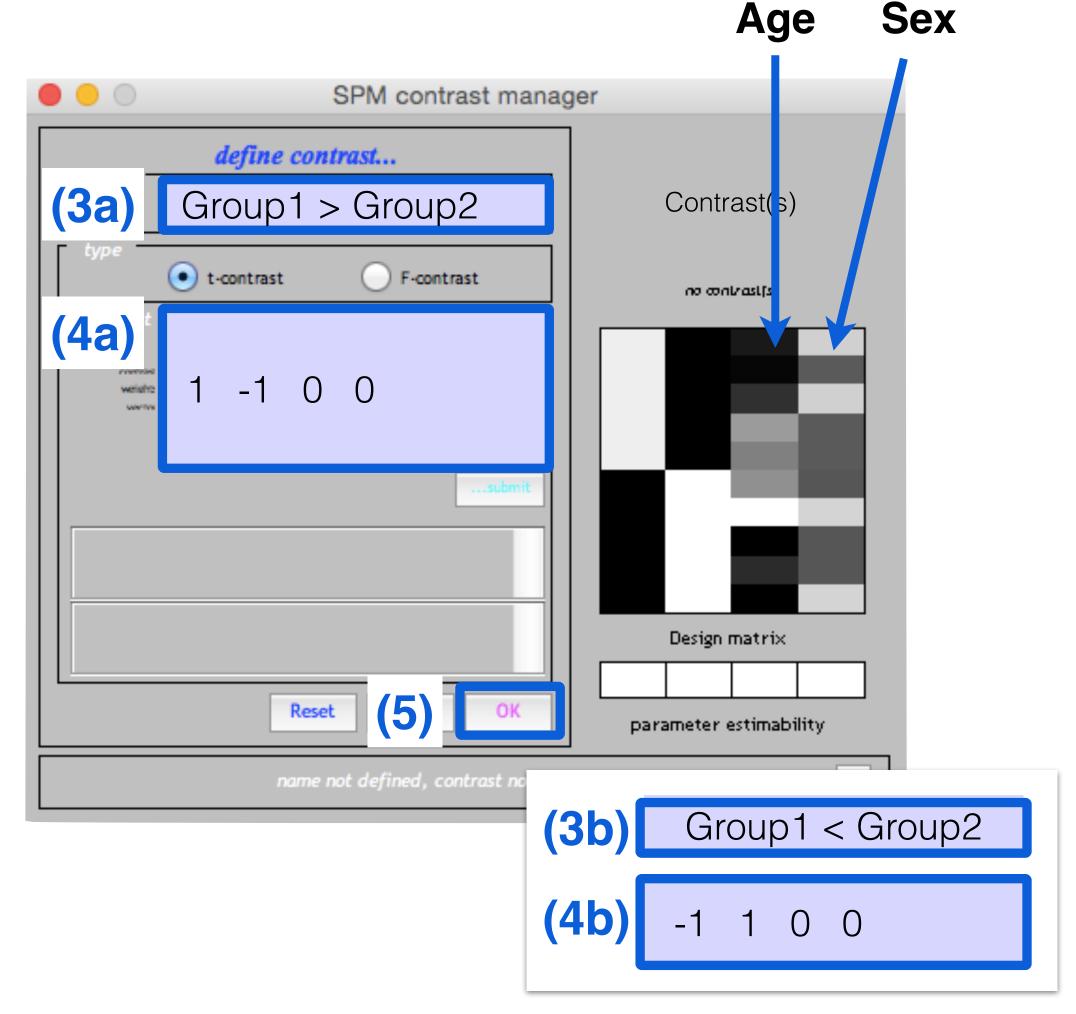
Slide의 8페이지에서 지정했던 Directory에 생성된 SPM.mat 파일을 선택함.

Run Batch를 클릭하여 프로그램을 실행.

### Contrast manager

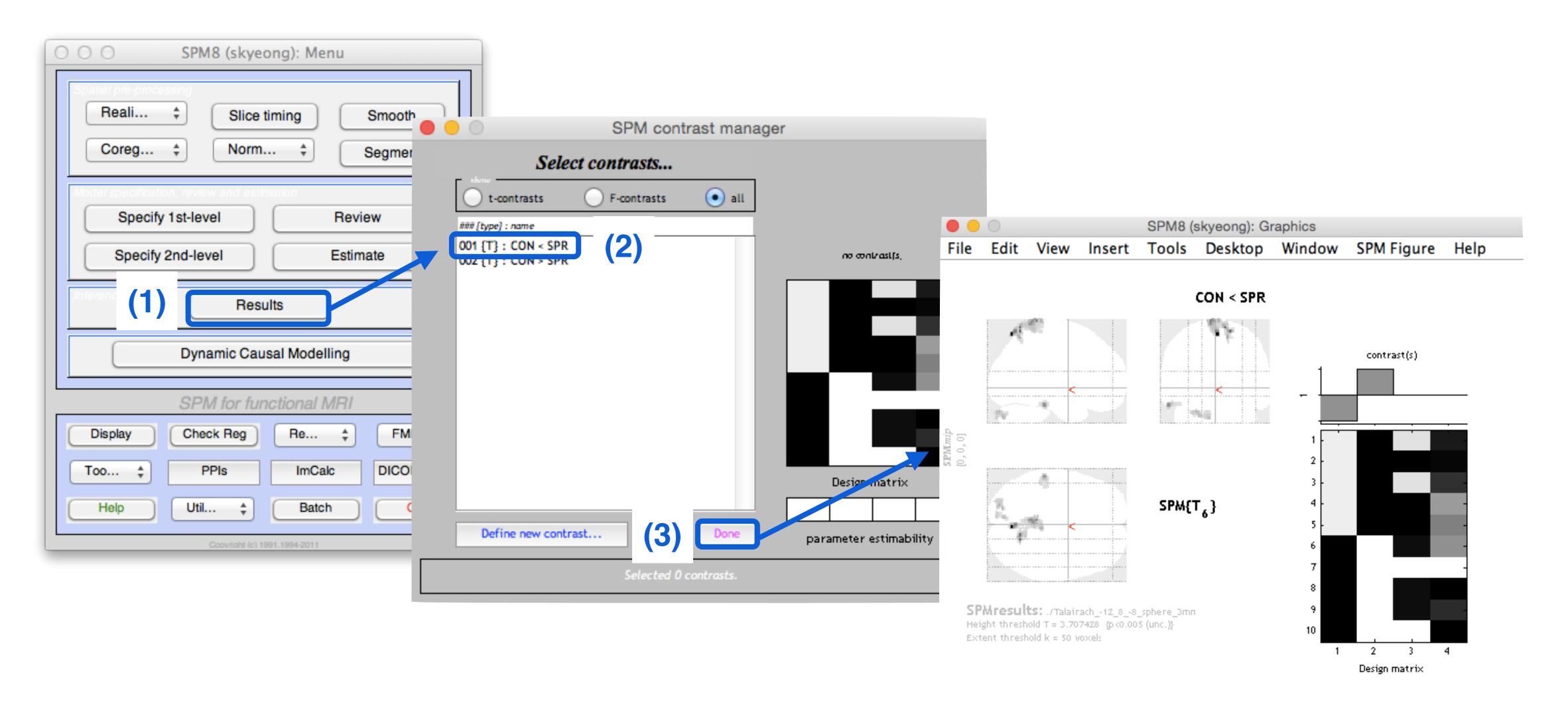


Group1>Group2 contrast는 Group1에서 증가된 FC or Group2에서 감소된 FC를 의미함.

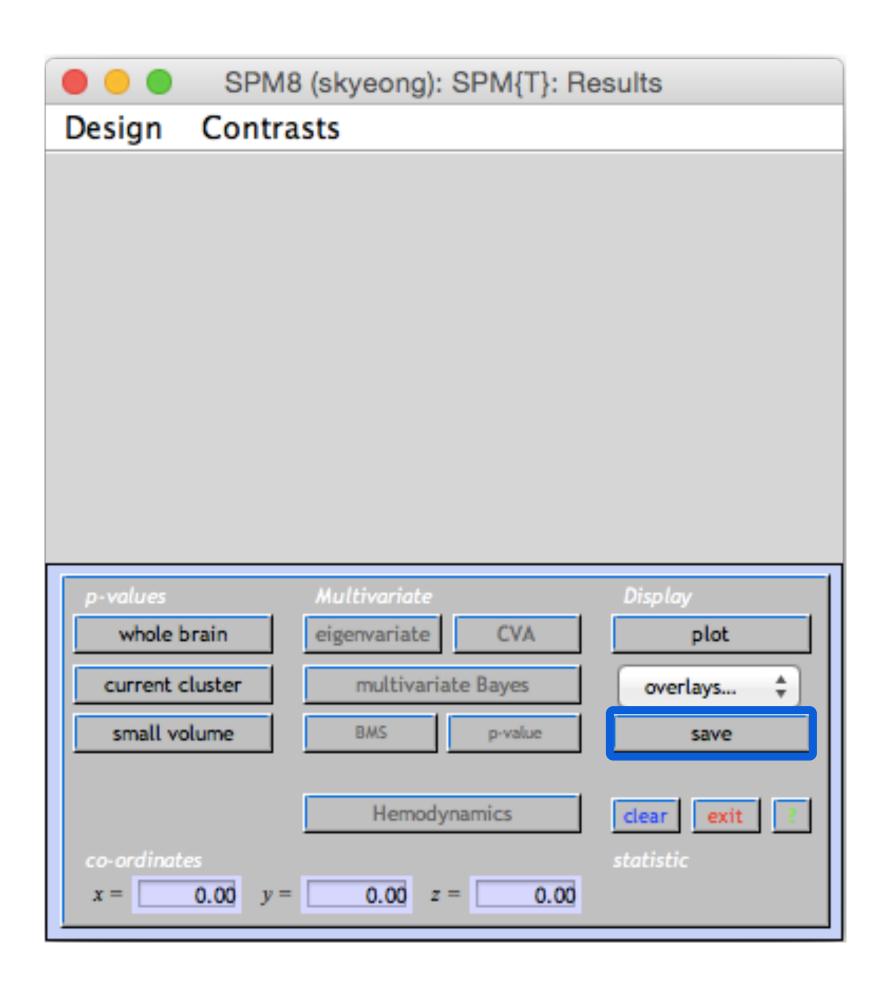


## Correlation analysis between FC and clinical scales

### in Results

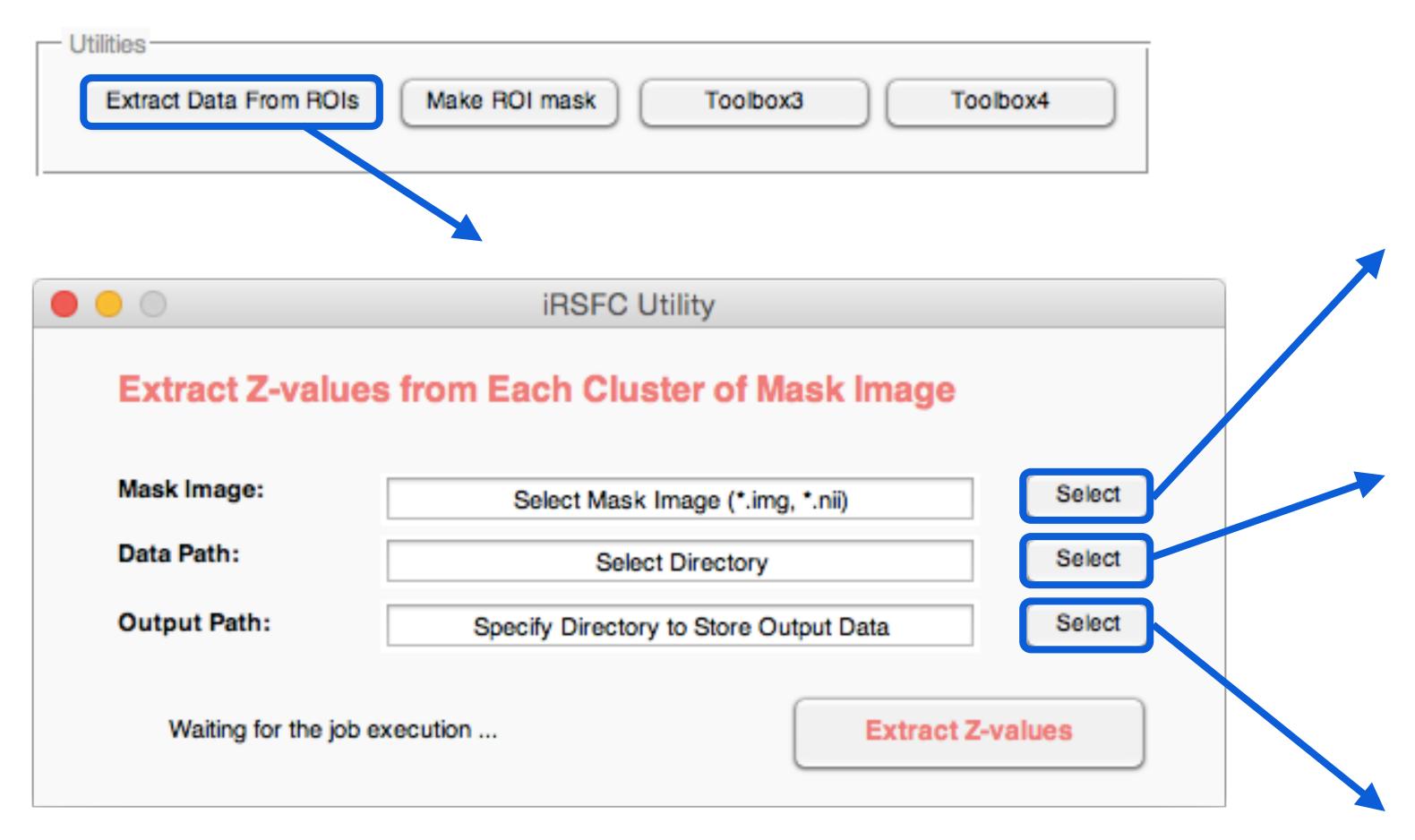


#### Save Results as MASK



To extract connectivity value in each cluster, regional clusters should be saved as mask image.

### Extract FC using iRSFC



page 14에서 저장했던 mask image 를 선택함. 또는 Maks ROI mask를 통해서 직접 그린 ROI를 선택할 수도 있음.

추출하고 싶은 데이터가 위치한 경로를 입력함. 데이터 구조는 다음과 같이 이루어져 있어야 함.

(Data path)\Lt\_Amygdala\subj001.img (Data path)\Lt\_Amygdala\subj002.img (Data path)\Lt\_Amygdala\subj003.img (Data path)\Lt\_Amygdala\subj004.img

결과 파일이 저장될 폴더를 선택함.

### Example of extracted FC

open tables.csv file which was saved in (Output path)

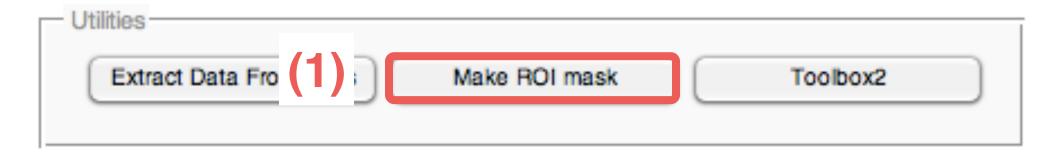
filename	cl01	cl02	cl03	cl04
zscore_Talairach12_88_sphere_3mm_CON1	-0.8	-1.3	-0.61	-0.84
zscore_Talairach12_88_sphere_3mm_CON2	-2.48	-3.64	-1.13	-0.79
zscore_Talairach12_88_sphere_3mm_CON3	-1.1	-1.6	0.36	-0.34
zscore_Talairach12_88_sphere_3mm_CON4	-0.66	-2.83	-2.14	-1.6
zscore_Talairach12_88_sphere_3mm_CON5	-1.05	-2.2	-1.24	-1.81
zscore_Talairach12_88_sphere_3mm_PAT1	1.36	0.82	0.93	2.09
zscore_Talairach12_88_sphere_3mm_PAT2	3.53	2.07	1.37	1.01
zscore_Talairach12_88_sphere_3mm_PAT3	-0.79	0.23	1.55	1.4
zscore_Talairach12_88_sphere_3mm_PAT4	1.51	0.25	2.2	2.45
zscore_Talairach12_88_sphere_3mm_PAT5	1.85	0.28	5.19	2.75

#### ROI connectivity Analysis

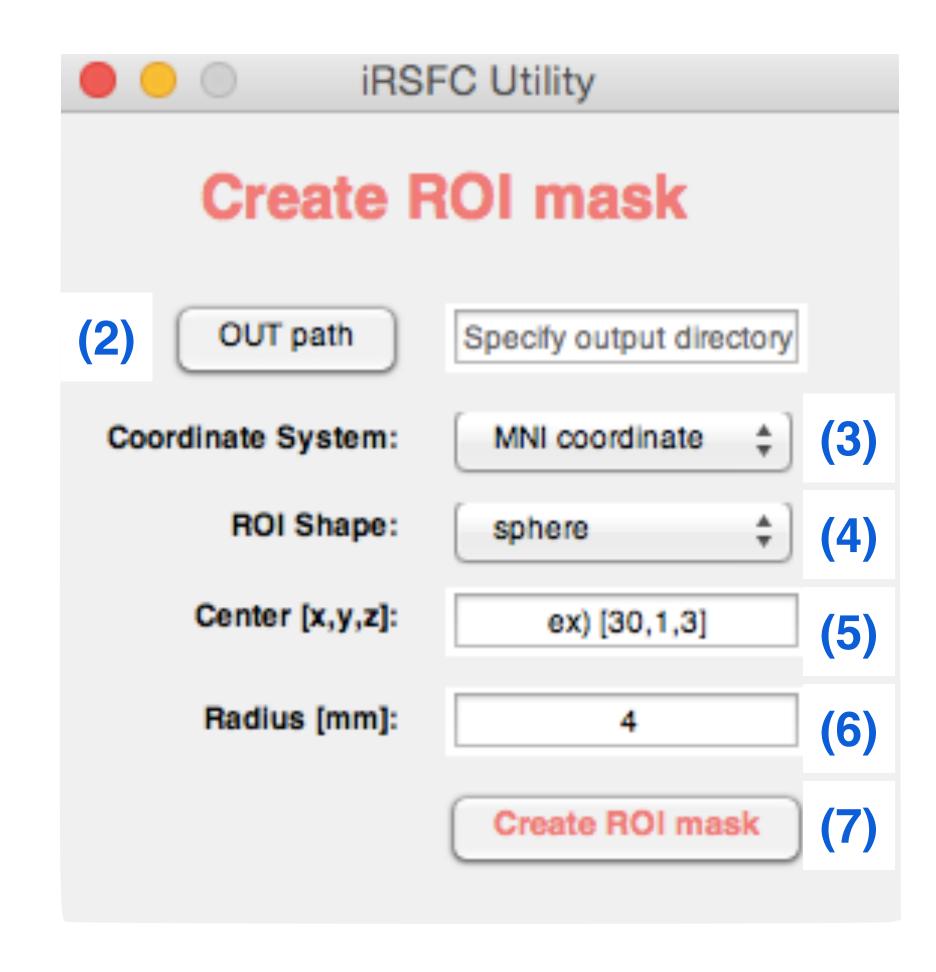
#### Create ROIs

#### Make ROI mask

sphere or box 모양의 ROI를 생성할 수 있음. (MNI or Talairach coordinate 좌표 모두 사용 가능함.)

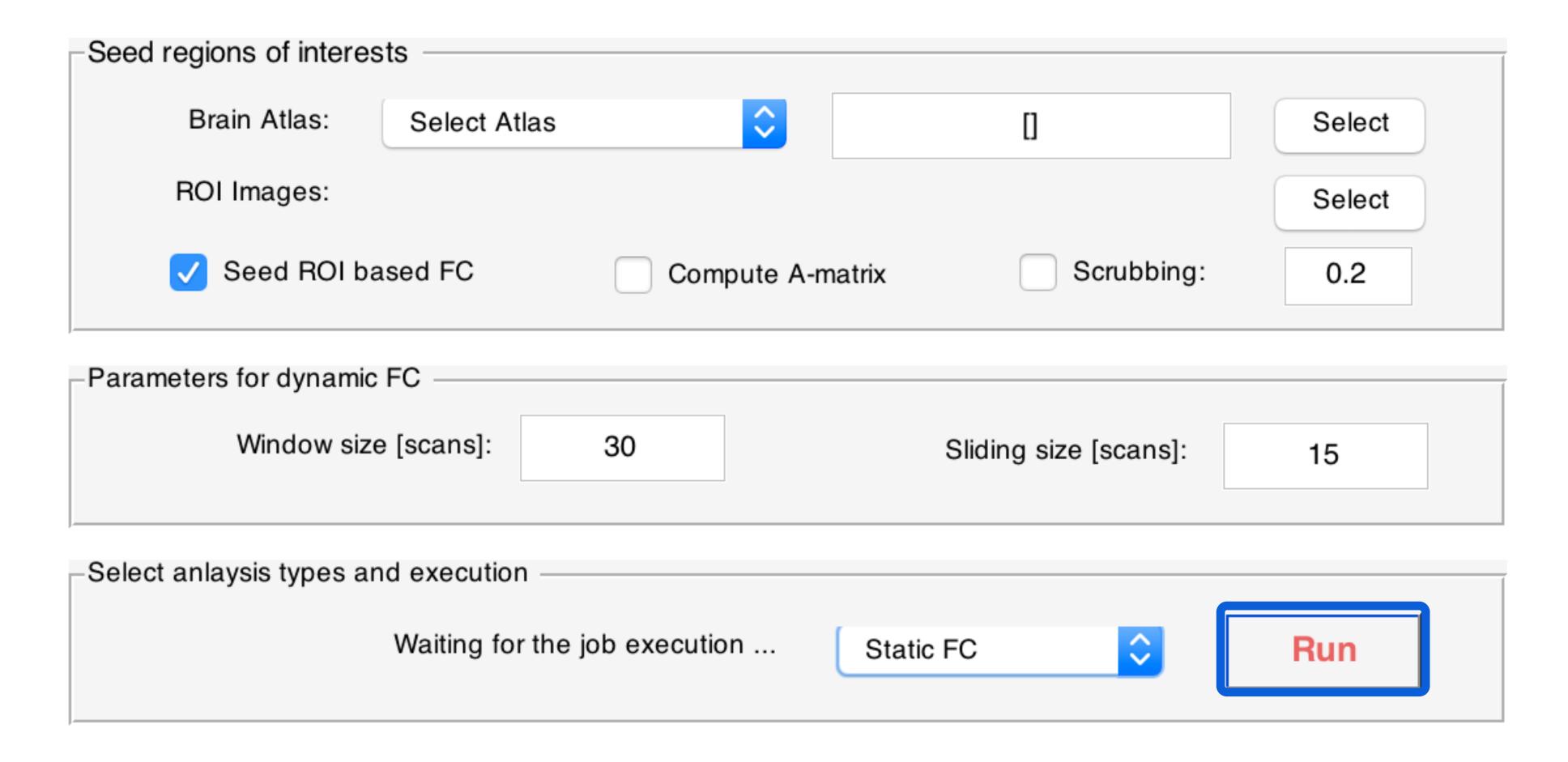


Region of Interest	X	у	Z
Left TPJ	-50	-31	23
Left ACC	-7	21	36
Left Broca's Area	-41	12	14
Left Amygdala	-24	-5	-18
Left Insula	-36	8	4
Right TPJ	50	-31	23
Right ACC	7	21	36
Right Homotopy of Broca's Area	41	12	14
Right Amygdala	24	-5	-18
Right Insula	36	8	4



Reference for ROIs, A. Vercammen et al. BIOL PSYCHIATRY 2010 67:912-918, http://neuro.imm.dtu.dk/services/jerne/ninf/voi.html

### Step 4 - Run Analysis



#### Check OUTPUT

node number	ROI file name		
1	Lt_ACC		
2	Lt_Amyg		
3	Lt_Broca		
4	Lt_Insula		
5	Lt_TPJ		
6	Rt_ACC		
7	Rt_Amyg		
8	Rt_Broca		
9	Rt_Insula		
10	Rt_TPJ		

subjname	ROI_1-2	ROI_1-3	ROI_1-4	ROI_1-5	ROI_1-6	ROI_1-7
CON1-20130825-KSK	0.068	0.408	0.818	0.123	0.957	-0.048
CON2-20130825-KKM	-0.164	0.577	0.62	0.051	0.753	-0.05
CON3-20130901-KMY	-0.28	0.081	0.202	-0.006	0.684	-0.392
CON4-20130929-KSB	0.02	0.146	0.739	0.095	0.371	-0.131
CON5-20131006-HTH	-0.245	0.039	0.1	0.154	0.203	-0.156
SPR1-20130901-CCS	0.058	0.229	0.514	0.301	0.577	-0.006
SPR2-20130901-OKS	0.218	-0.236	-0.262	0.128	0.264	-0.187
SPR3-20130901-BY	-0.017	0.033	0.021	-0.215	0.342	-0.081
SPR4-20131013-SSH	0.069	0.115	0.372	0.207	0.625	0.131
SPR5-20130929-YMJ	0.225	0.275	0.663	0.046	0.624	0.01

#### References for Atlas

#### • **AAL Atlas** (*n*=116)

N. Tzourio-Mazoyer *et al.* (2002), Automated Anatomical Labeling of Activations in SPM Using a Macroscopic Anatomical Parcellation of the MNI MRI Single-Subject Brain, NeuroImage 15, 273-289 (2002)

#### • Dosenbach Atlas (n=160)

Nico U. F. Dosenbach *et al.* (2010), Prediction of Individual Brain Maturity Using fMRI, *Science* 329:5997 pp.1358-1361.

n=160, sphere shape, radius =5 mm, minimum distances between ROI center = 10 mm, no overlap among ROIs