# **Human fMRI project**

2022. 11. 08

설재민

### 1. Subject summary

- a. 16 subjects 사용 가능(10 good / 6 bad perform.)
- b. 나머지 17 subjects는 fMRI data가 없거나 부족

#### 2. Behavior data

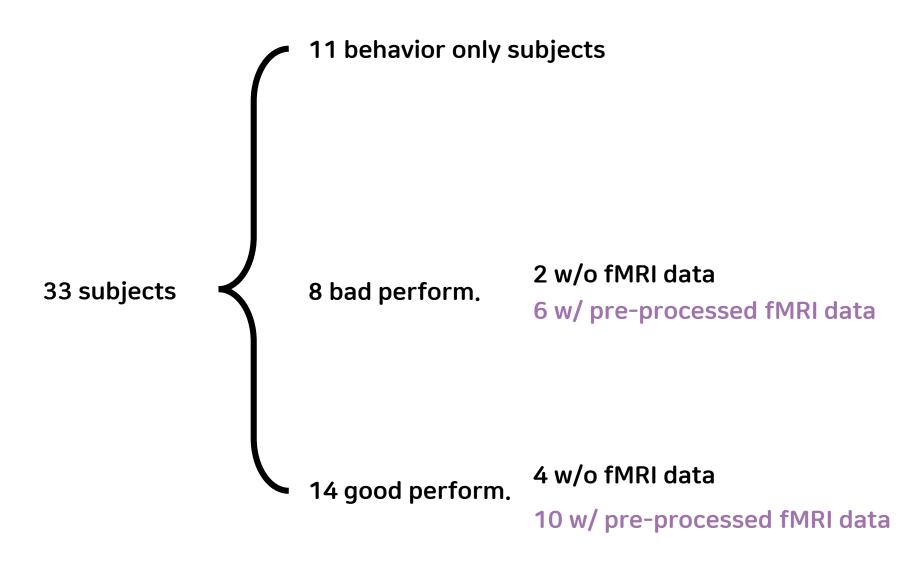
- a. Timestamp별 position head direction data 없어서, UDK log로부터 trial parsing 다시 해야 함
- b. MR signal timstamp는 있음

#### 3. MR data

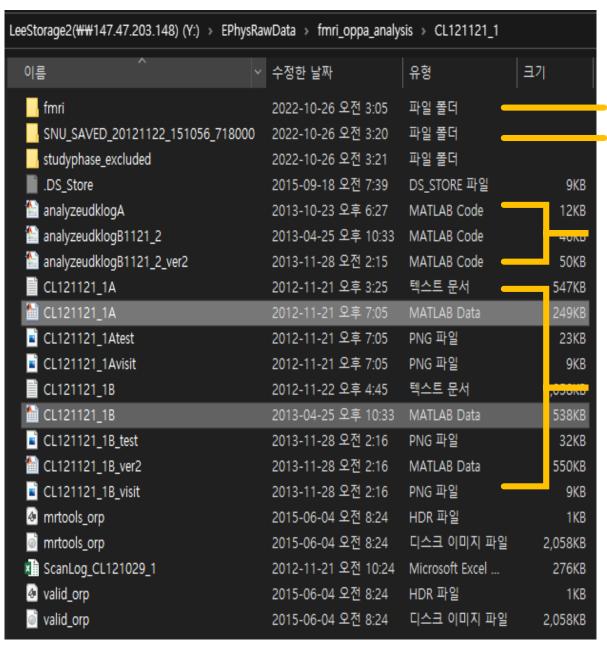
- a. SPM 사용하여 MR image processing 필요
- b. Hippocampus segmentation 및 preprocessing 완료됨

### **Subject summary**

총 16명 fMRI data를 사용할 수 있음



#### **Data structure**



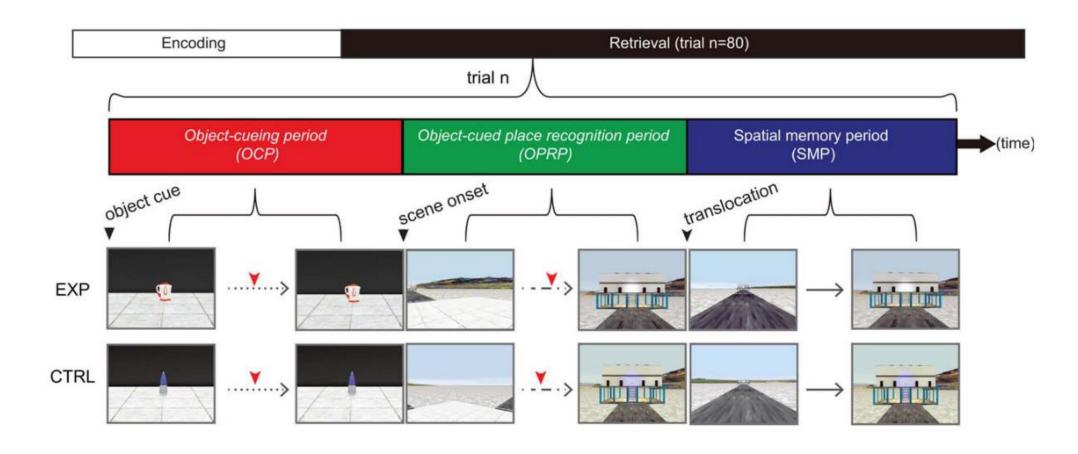
fMRI pre-processed datafMRI raw data

**UDK** log parsing program code

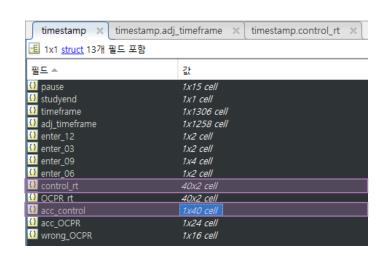
UDK log file (A=study session, B=test session)

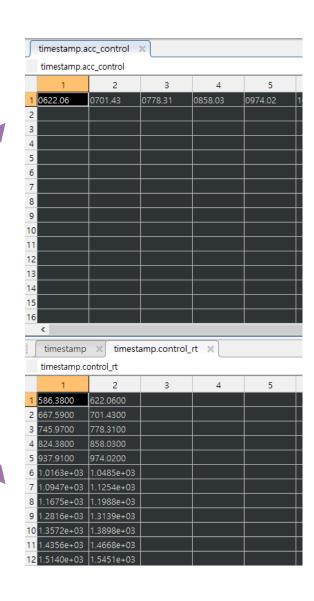
# Trial 구조 확인

: CTRL trial + EXP trial



# CTRL trials timestamp 확인

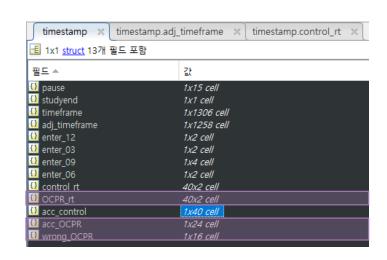


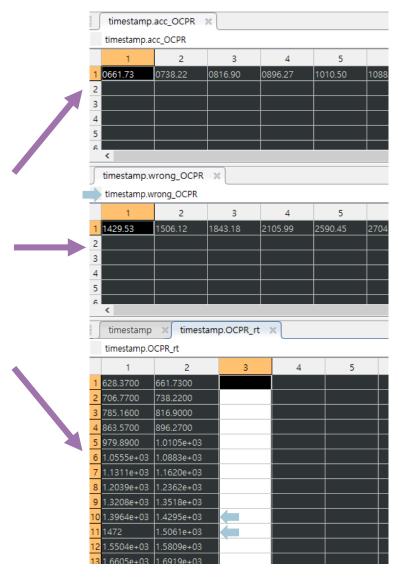


CTRL trials, Correct Trial end timestamp

CTRL trials, All
Trial start - Trial end

# EXP trials timestamp 확인



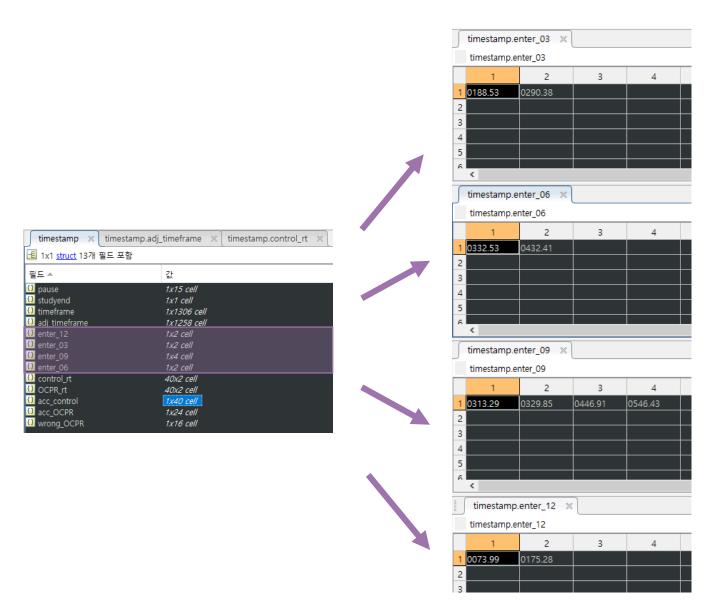


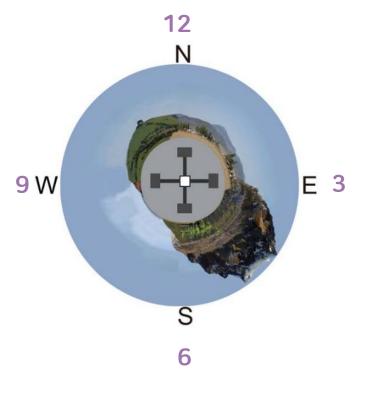
EXP trials, Correct Trial end timestamp

EXP trials, Wrong Trial end timestamp

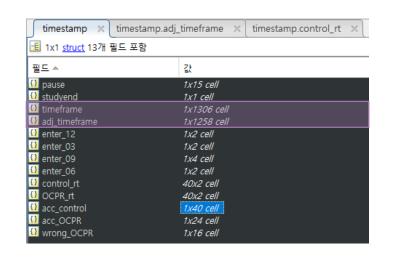
EXP trials, All Trial start - Trial end

# Encoding phase timestamp 확인

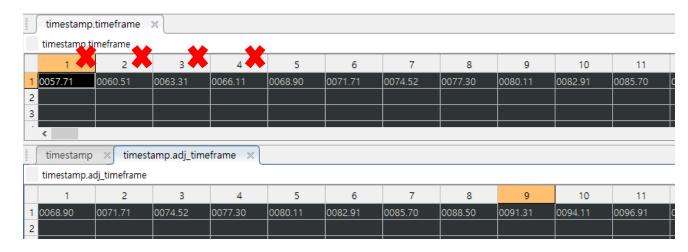




# MR timestamp 확인

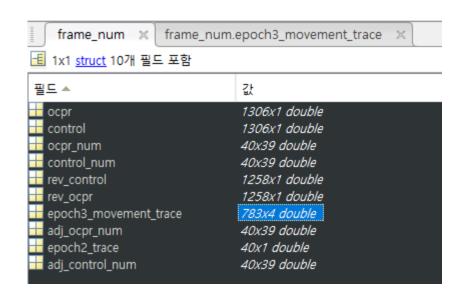


#### fMRI에서 2.8초마다 나오는 신호 + manual input(키보드 5)

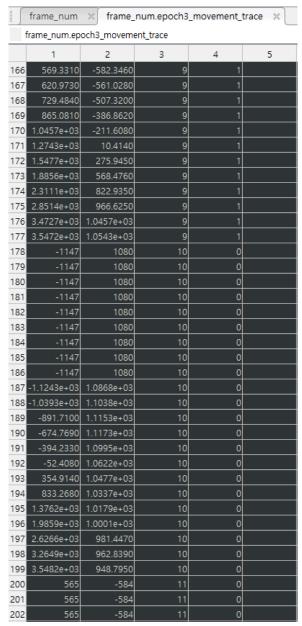


Manual input 제거하고 실제 MR image가 있는 timestamp만 모음

## Position info : raw data parsing 필요



- 각 position 값의 timestamp는?
- 각 timestamp에서 head direction은?
- 각 trial의 correct answer / player choice 정보?

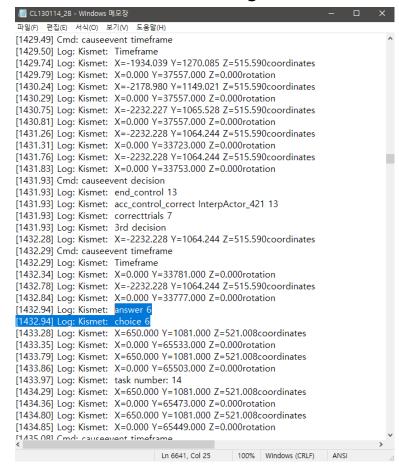


X? Y? EXP Correctness? Trial?

### Position info : raw data parsing에 필요한 raw data와 parsing code 모두 있음

- 각 timestamp의 position 및 head direction
- 각 trial의 correct answer / player choice

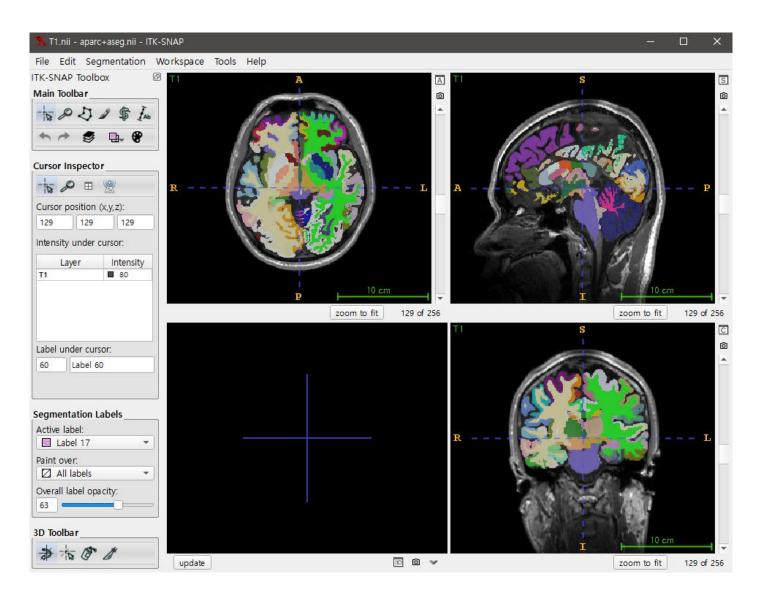
#### **Raw UDK log**



#### Log parsing code

```
analyzeudklogB0114_2_ver2.m × +
        close all;clear all;clc
        filename='CL130114_2B';
        fileToRead1=strcat(filename, '.log');
        DELIMITER = ' ';
11 -
        HEADERLINES = 22511;
12 -
        no_trials=80;
13
15 -
        rawData1 = importdata(fileToRead1, DELIMITER, HEADERLINES);
20 -
        [~,name] = fileparts(fileToRead1);
        log_data=rawData1;
22
24 -
        no_lines=max(size(log_data));
        count_pause=0;count_timeframe=0; count_12=0; count_03=0; count_06=0; count_09=0;
     ☐ for i=1:1:no_lines
28 -
           k=strfind(log_data{i}, 'Pause');
29 -
            if ~isemptv(k)
30 -
                count_pause=count_pause+1;
31 -
                line_indice_pause(count_pause)=i;
32 -
                string=log_data{i};
                timestamp.pause{count_pause}=string(2:8);
```

### MR data: SPM 사용한 trial parsing 필요



- Freesurfer 사용한 segmentation 완료
- SPM을 사용한 pre-processing 완료
- Mat file로 MR image가 변환된 것이 없어서, pre-processing 완료된 파일을 SPM으로 변환 필요 (timestamp \* voxel 형태)

## Analysis plan

~11/09

Behavior data UDK log parsing하여,

timestamp별 position, head direction, trial info 표시된 array 생성

~11/11

MR data SPM processing 진행,

timestamp\*voxel 형태의 MR BOLD activity array 생성

~11/15

Trial별 hippocampus activity average,

trial - correctness - reaction time - HPC activity로 구성된 array 생성 correct / wrong trial에 따른 hippocampus activity 분포 확인