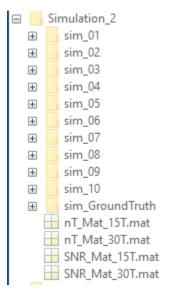
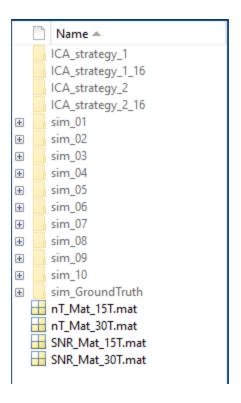
```
%% 1.5T simulation
        % rng('default');
        D_SNR = randi([40,110],100,1)+randn(100,1);
 8 - for a = 1:5
10 -
            D_SNR_a = D_SNR(a); %%% SNR levels, this can be a CNR value as well Randomly generate number from 40 to 45
            SNR Mat(a, :) = D_SNR_a; %%% save the SNR is the loop
12
13 -
          nT_a = randi([50,100],1,1); %%% number of subject Randomly generate number from 30 to 100
14 -
            nT Mat(a,:) = nT a;
15 -
            nV = 300; %%% size of the image would be nV*nV
            comp_ID = [3 8 9 11 16 17 19 20 21 22 23 25 26 29 30];%%% select 15 components
16 -
17 -
            verbose_display = 0; %%% if you want to see which components you've selected, reset this value to 1
18
19 -
            mkdir(['C:\Simulation_Data\Simulation_l\sim_',sprintf('%02d',a)]); %%% make a dictionary to save the simulated data
            out_path = [['C:\Simulation_Data\Simulation_l\sim_',sprintf('%02d',a)]]; %%you need to change the out path of your owr sP = simulation_sMRI_simTB_ryg_20200717(nT_a,D_SNR_a,out_path,nV,comp_ID,verbose_display);
20 -
21 -
            simtb_main_ryg_sMRI(sP);
22 -
23
24 -
25 -
26 -
        cd('C:\Simulation_Data\Simulation_1');
        save SNR Mat 15T SNR Mat;
27 -
        save nT_Mat_15T nT_Mat;
29 -
        cd('C:\Simulation_Data\Simulation_1');
```

1. In line 13, change the range of the subjects from 50-100 to 30-100 for 1.5T, do the same for 3.0T, as well. Since this is the second time simulation, change all output direction to a new folder. Then you should have following files in your new folder. (code file: demo_sim10sites.m)



2. Before you go change the Strategy code, remember to add a few new folders as the output direction for the ICA results, your simulation data file should look like this.



3. Now since we generated new data, the input and output direction in both strategy code need to be changed, and locate the iceMask.nii file which is contained in the githubGitHub file, as well.— (code file-: batchICA strategy1.m and batchICA strategy2.m)

```
% Input data file pattern for data-sets must be in a cell array. The no. of rows of cell array correspond to no. of subje
 61
       % and columns correspond to sessions. In the below example, there are 3
       % subjects and 1 session. If you have multiple sessions, please see
 62
        % Input data subjects 2.m file.
       input_data_file_patterns = {'C:\Simulation_Data\Simulation_1\sim_01\Sim*.nii';
 65
             C:\Simulation_Data\Simulation_1\sim_02\Sim*.nii';
 66
           'C:\Simulation_Data\Simulation_l\sim_03\Sim*.nii';
           'C:\Simulation_Data\Simulation_1\sim_04\Sim*.nii';
 67
           'C:\Simulation_Data\Simulation_1\sim_05\Sim*.nii';
 68
           'C:\Simulation_Data\Simulation_l\sim_06\Sim*.nii';
 70
           'C:\Simulation_Data\Simulation_1\sim_07\Sim*.nii';
 71
           'C:\Simulation_Data\Simulation_1\sim_08\Sim*.nii';
 72
           'C:\Simulation Data\Simulation 1\sim 09\Sim*.nii';
 73
           'C:\Simulation_Data\Simulation_1\sim_10\Sim*.nii';};
85
       %% Enter directory to put results of analysis
86 -
      outputDir = 'C:\Simulation_Data\Simulation_1\ICA_strategy1_16';
87
       %% Enter Name (Prefix) Of Output Files
88
89 -
      prefix = 'ica';
90
91
       %% Enter location (full file path) of the image file to use as mask
       % or use Default mask which is []
 %% Enter location (full file path) of the image file to use as mask
 % or use Default mask which is []
 maskFile = ['C:\Simulation Data\MRI-ICA-remake\MRI-ICA-master\src\icaMask.nii'];
```

4. If you want your result display contains 16 or 15 diagrams in 1 picture, change line142-143.

```
%% Number of pc to reduce each subject down to at each reduction step
% The number of independent components the will be extracted is the same as
% the number of principal components after the final data reduction step.

142 - numOfPC1 = 16;
143 - numOfPC2 = 16;
144 % numOfPC3 = 25;
145
```

5. After you finished changing code in both strategy, open file run_ICAbatch.m in matlab, and make sure the code finds the path of strategy code. (code file-: run_ICAbatch.m)

```
4 - inputFiles = ['C:\Simulation_Data\MRI-ICA-remake\MRI-ICA-master\src\batchICA_strategyl.m'];
5 - icatb_batch_file_run(inputFiles);
6 - close all;clear;
7
8 - inputFiles = ['C:\Simulation_Data\MRI-ICA-remake\MRI-ICA-master\src\batchICA_strategy2.m'];
9 - icatb_batch_file_run(inputFiles);
10 - close all;clear;
```

6. Now run 'run_ICAbatch.m', it might take a couple minutes. It's done when the command window shows

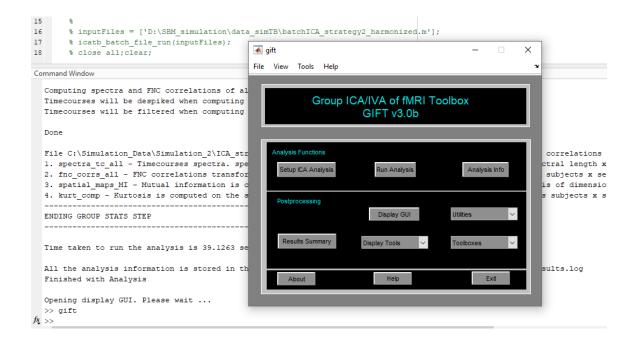
```
Time taken to run the analysis is 39.1263 seconds

All the analysis information is stored in the file C:\Simulation_Data\Simulation_2\ICA_strategy_2_16\ica_results.log
Finished with Analysis

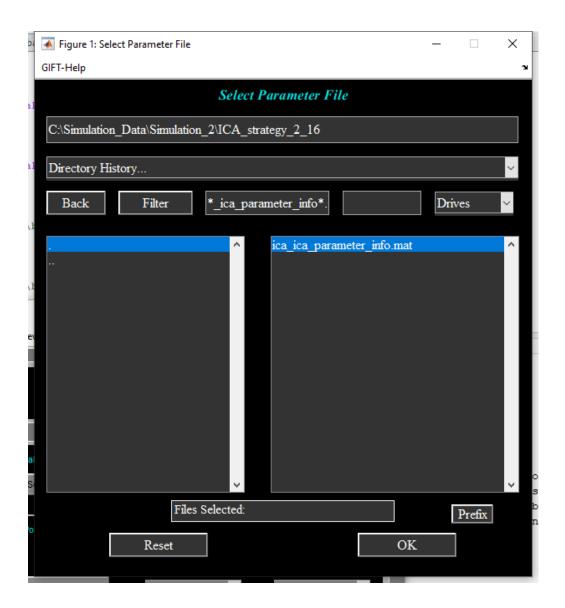
Opening display GUI. Please wait ...

at the very end.
```

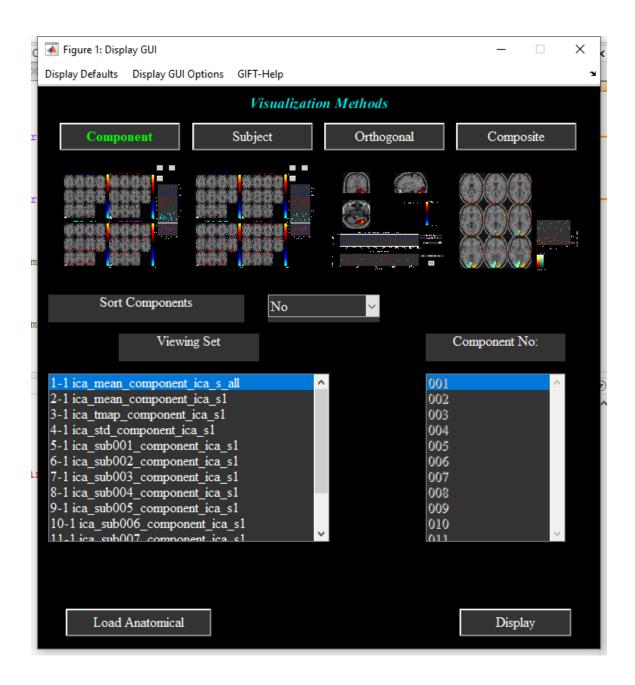
7. Now type 'gift' in command window and press Enter, a toolbox like below will show up.



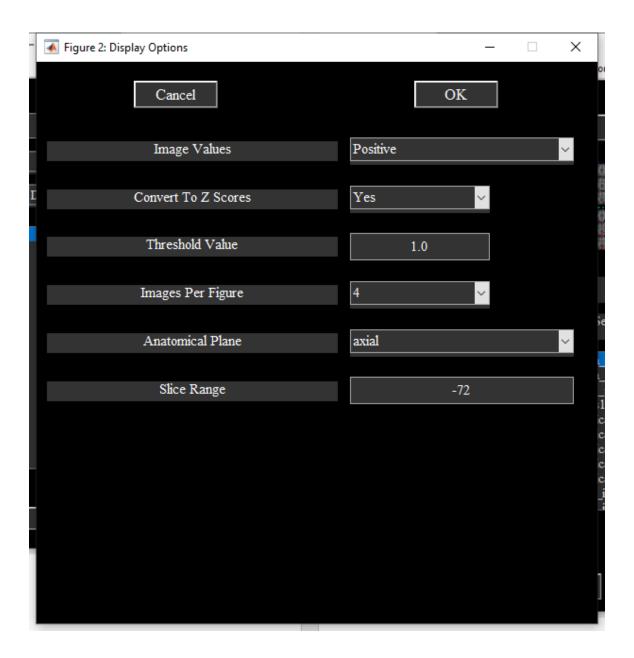
8. Then click 'Display GUI', this should appear, and you can change folders from the left side and choose file from right side



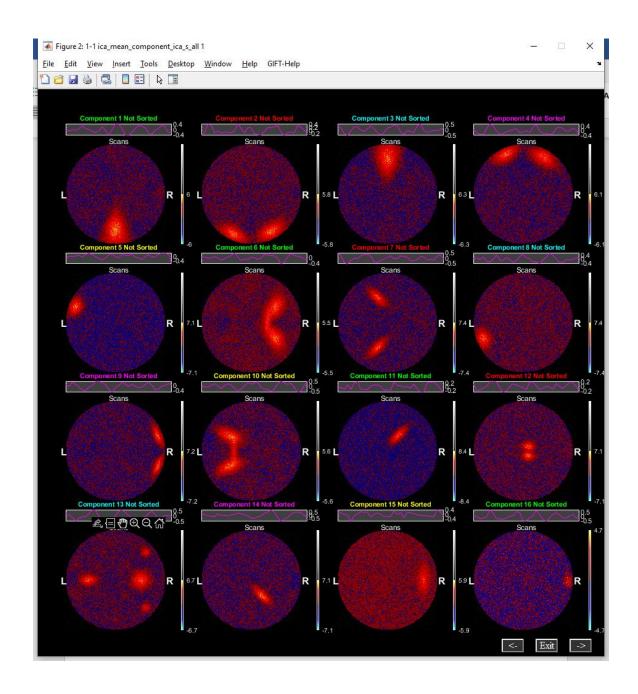
9. Click 'OK'. This would be shown on your screen.



10. Now before you hit 'Display' at the bottom right, click 'Display Defaults' on the top left.



11. Now change the 'Image values' to 'Positive and Negative', 'Threshold Value' to 0, 'Image Per Figure' to 16, Click 'OK'. Now click 'Display', it might take a few second then the results should be shown in form like



End of the Manuel Manual