Predicting functional connectivity(FC) using dynamic mean field model for a given structural connectivity(SC) matrix

The file *DMF_lesioned.m* implements dynamic mean field model described in Deco et al., 2014. Use the below syntax to call this function.

[fc, maxfrN] = DMF_lesioned(scPath, scName, fic, ffi, simTime, dt, G, noiseAmp, lesionAreas, ficWeights, setFICWeights);

Input parameters

scPath : Path to SC matrix file

scName : SC matrix path in the given hdf5 file

fic : true or false, set it to true to use feedback inhibtion control(FIC)

algorithm as proposed in Deco et al., 2014 and false otherwise

ffi : true or false, set it to true to enable feedforward inhibition

simTime : simulation time in milliseconds dt : Step size for numerical integration

G : Global coupling strength

noiseAmp : noise amplitude

lesionAreas : Index of the areas which are lesioned in the SC ficWeights : local inhibitory synaptic coupling initialization values

setFICWeights : [True/False] if set true ficWeights will be used for initialization

Return Values

fc : Predicted functional connectivity.

maxfrN : maximum firing rate of excitatory population in each area

throughout the simulation.

Example 1: Simulation for predicting FC of a healthy subject.

This example assumes SC matrix is stored in /home/data/Human_68.hdf5 under the name 'C'. Simulates 8 minutes of activity.

```
scPath = '/home/data/Human_68.hdf5';
scName = '/C';
fic = true;
ffi = false;
simTime = 8*60*1000;
dt = 0.1;
G = 0.5;
noiseAmp = 0.001;
lesionAreas = [];
ficWeights = [];
setFICWeights = false;
[fc, maxfrN] = DMF_lesioned(scPath, scName, fic, ffi, simTime, dt, G, noiseAmp, lesionAreas, ficWeights, setFICWeights);
```

Generating virtually lesioned SC from the SC of healthy subjects.

The file gen_lesioned_sc_nn.m implements a function to generate virtually lesiond SC matrix from SC of healthy subjects.

Input Parameters

scPath : Path to SC matrix file

centresPath : Path to file(hdf5 formate) with spatial coordinates of centres of

each area.

scName : SC matrix path in the given hdf5 file

centresName : Path to centres matrix in the hdf5 file located at centresPath

lesionIdx : Index of the area that should act as lesion centre

lesionPercent : Percentage of total areas that would be lesioned along with lesion

centre. This many nearest neighbours to lesion centre would be

also lesioned

saveDir : Directory where lesioned SC matrix should be saved

Return Values

ISC : A virtually lesioned SC matrix with lesion centre given by

lesionIdx

Example2: Generating a virtually lesioned SC with lesion centre as ROI number 25. Along with lesion centre, 5 nearest neighbours of lesion centre in the same hemisphere are also lesioned.

Assuming there are a total of 68 ROIs (34 per each hemisphere) and 5 nearest neighbours are lesioned along with

```
scPath = '/home/data/Human_68.hdf5'
centresPath = '/home/data/ROI_centres.hdf5'
scName = '/C';
centreName = '/Centres';
lesionIdx = 25;
lesionPercent = (5.0/68)*100;
saveDir = '/home/results/';
ISC = gen_lesioned_sc_nn(scPath, centresPath, scName, centresName, lesionIdx, lesionPercent, saveDir);
```

Predicting FC using DMF model and lesioned SC.

Lesioned structural connectivity from example 2 will be stored as /home/results/sc_l25_nn5.hdf5. Predicting FC with such lesioned SC is similar to example 1 except now the SC is a lesioned one. Use the below syntax for predicting FC of a lesioned SC. Changes from example 1 are highlighted in yellow.

```
ScPath = '/home/results/sc_l25_nn5.hdf5';
scName = '/C';
fic = true;
ffi = false;
simTime = 8*60*1000;
dt = 0.1;
G = 0.5;
noiseAmp = 0.001;
lesionAreas = [25];
ficWeights = false;
[fc, maxfrN] = DMF_lesioned(scPath, scName, fic, ffi, simTime, dt, G, noiseAmp, lesionAreas, ficWeights, setFICWeights);
```