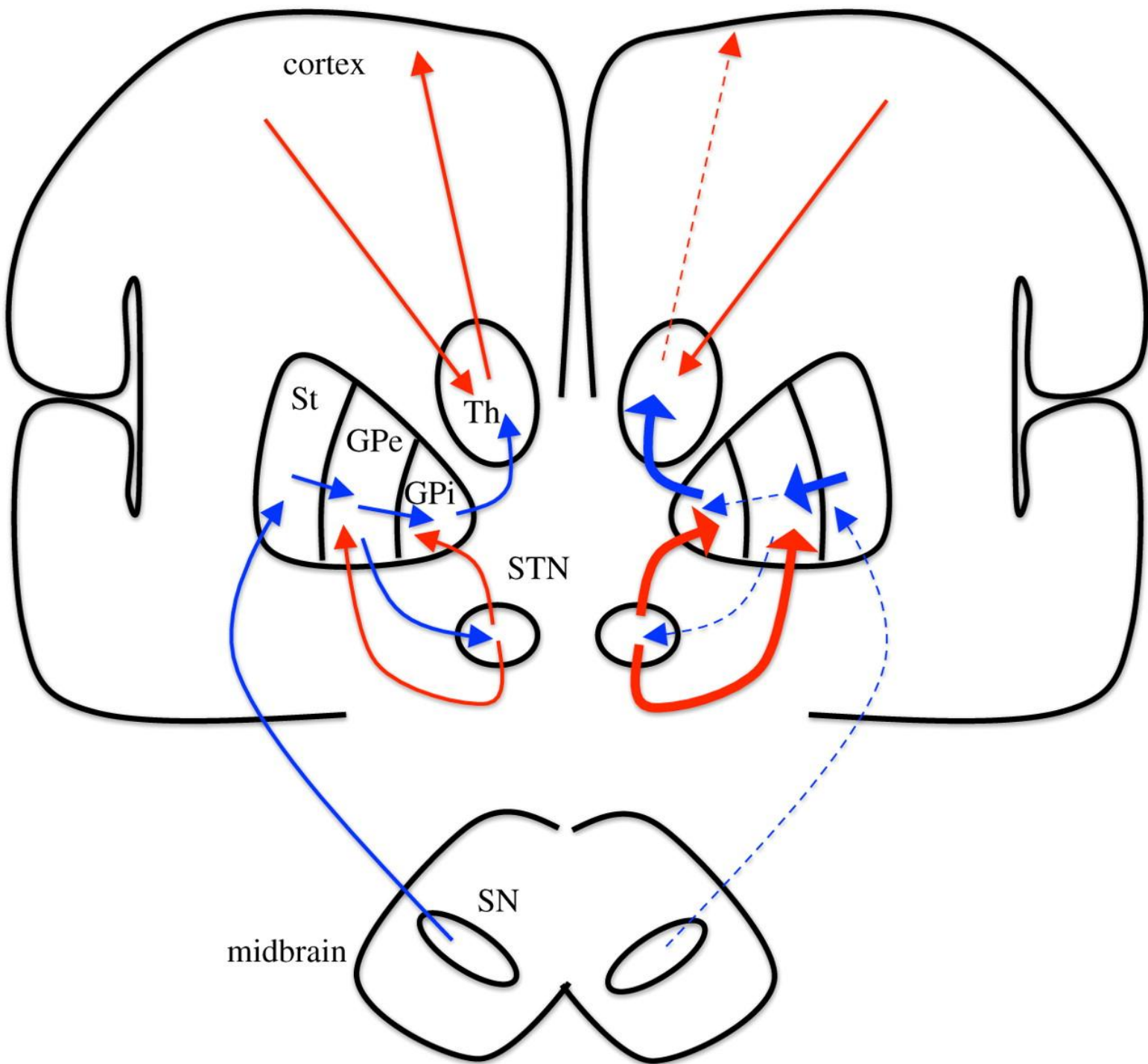


normal

Parkinson's disease



# **Toward a Dynamic Model of Parkinson's Disease**

Thoughts & Research Notes

By Wisam Reid & Iran Roman

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# 1 A Brief Overview

We are working toward a dynamic model of Parkinson's Disease (PD).

As a first step we are building a model of healthy coupled brain oscillation during movement coordination in healthy subjects and stroke victims, as observed by Takako Fujioka's MEG studies and double-blinded randomized clinical trials on stroke recovery. We then want to build up towards a PD model that will hold up against everything currently known about PD.

## 2 The Data

MEG data from healthy subjects and stroke victims during a tap based rhythm task. Collected by Takako Fujioka. [1] [2]

## 3 A Simple Baseline Conceptual Model

This is an attempt at replicating studies indicating that beta ERD occurs after hearing a beat, and at that point gamma ERS occurs. Below is a Matlab script and the plots that it generates.

### 3.1 Plots

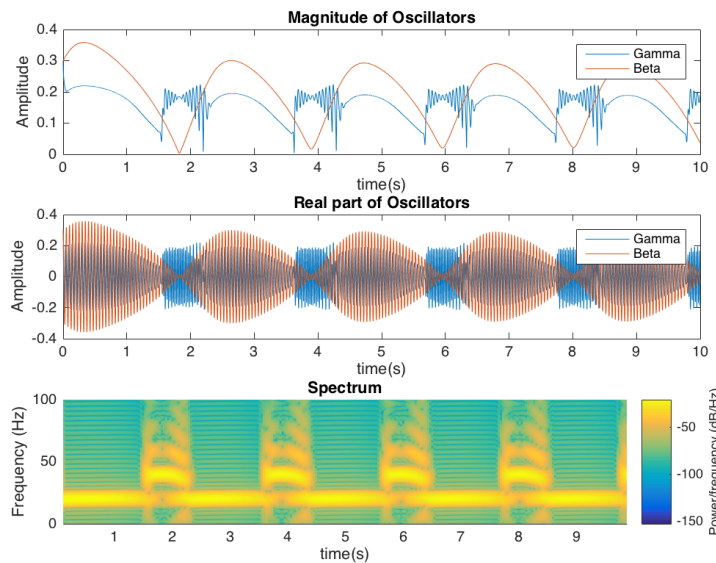


Figure 1: A "beta" oscillator driving a "gamma" oscillator. The "beta" oscillator is modulated by a 0.5 Hz envelope, which is supposed to emulate the result of stimulation by listening of an isochronous rhythm. The "gamma" oscillator is tuned to a limit cycle and oscillates at its frequency only when the beta oscillator has a low amplitude.

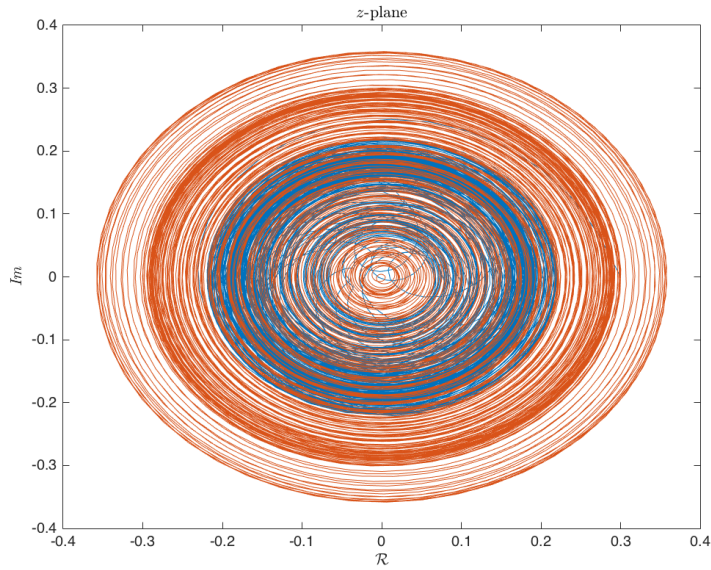


Figure 2: The "beta" and "gamma" oscillators in the  $z$ -plane

## 3.2 Matlab Code

```
%%%%%%%%%%%%%%%%%%%%%%%%%%%%%%%%%%%%%%%%%%%%%%%%%%%%%%%%%%%%%%%%%%%%%%%%%
%
% Authors: Iran Roman & Wisam Reid
%
% This script shows a limit cycle oscillator
% in the beta band (20Hz)
%
%%%%%%%%%%%%%%%%%%%%%%%%%%%%%%%%%%%%%%%%%%%%%%%%%%%%%%%%%%%%%%%%%%%%%%%%%
%% CLEAN AND CLEAR

clc;
clear;
close all;

%% Include the GrFNN toolbox
addpath(genpath(' /Documents/GrFNNToolbox/'))

%% the output of beta drives a gamma oscillator

dzdt = @(t,z,alpha,beta1,beta2,epsilon,F,omega0) ...
    [z(1)*(100*alpha + 1i*2*pi*40 + 200*beta1*abs(z(1))^2 + ...
    (epsilon*beta2*abs(z(1))^4)/(1-epsilon*abs(z(1))^2)) + 3.5*(z(2)*exp(pi
    → )); ...
    (z(2)*(alpha + 1i*2*pi*20 + beta1*abs(z(2))^2 + ... % beta layer
    (epsilon*beta2*abs(z(2))^4)/(1-epsilon*abs(z(2))^2)) + ...
    F*exp(1i*omega0*t))]; % input to beta

%% parameters

% for the limit cycle oscillator
z0 = 0.3;
```

```

alpha = 1;
beta1 = -15;
beta2 = -1;
epsilon = 1;
F = 0.5;
f0 = 20.5;
omega0 = 2*pi*f0;

% for time
fs = 1000;
dur = 10; % in seconds
T = 1/fs;
time = 0:T:dur;

%% integrate the ode to obtain the amplitude

[t,z] = ode45(@(t,z) dzdt(t,z,alpha,beta1,beta2,epsilon,F,omega0),time,[z0,
    ↪ z0]);

% extract information
r = abs(z);
dr2dt = diff(r(:,1));
dr1dt = diff(r(:,2));

%% Plot

figure(2)
plot(z)
xlabel(' $\mathcal{R}$ ', 'Interpreter','latex')
title(' $z$ - plane ', 'Interpreter','latex')
ylabel(' $\textit{Im}$ ', 'Interpreter','latex')

figure(3)
subplot(3,1,1)
plot(time,r)
title(' Magnitude of Oscillators ')
ylabel(' Amplitude ')
xlabel(' time(s) ')
legend('Gamma','Beta')
subplot(3,1,2)
plot(time,real(z))
title(' Real part of Oscillators ')
legend('Gamma','Beta')
ylabel(' Amplitude ')
xlabel(' time(s) ')
subplot(3,1,3)
spectrogram((z(:,1)),kaiser(256,5),220,20000,fs,'yaxis')
title(' Spectrum ')
ylim([0 100])
xlabel(' time(s) ')

```

### 3.3 To Do

After Iran's meeting with Takako, we know that this needs some refinement. Since, in healthy subjects gamma does not spontaneously oscillate. Beta does, and Gamma must

be evoked by sound. Beta is always oscillating and gamma gives exogenous or endogenous input to affect beta. High gamma gives rise to the desynchronization of beta [3].

In contrast, with the advance of PD, dopamine is depleted in the basal ganglia, and experimental evidence suggests that this depletion has an effect on the synaptic weights of the subthalamic nucleus-external globus pallidus (STN-GPe) network. This gives rise to constant beta oscillation and obstructs motor coordination. In deep brain stimulation (DBS) patients, 20 Hz stimulation is ineffective, with 40 Hz gamma nothing happens (no beta desynchronization), and with high gamma 80 Hz there is improvement.

**We plan to modify this baseline model to accommodate the correct healthy behavior.**

## 4 Proposed Method

We want to move toward something much more interesting.

Said quickly, we want to get away from “engineering” bifurcation parameters. We want to marry ONNs that fit their parameters directly from brain data, with dynamic causal analysis.

### 4.1 Some Philosophical Alignment

As you already know, Iran and I are both interested in Oscillatory/Spiking Neural Networks (O/SNN) and with that comes a lot of the interesting questions. We want to model the brain in the framework of Bifurcation Theory, but modeling at what scale is appropriate or most beneficial? Without going on a tangent here, I would say that those answers depend on what you are measuring and what you are trying to model. Here are some papers that are currently inspiring this thought process [4] [5] [6].

### 4.2 Fitting Bifurcation Parameters to Data

My literature review started with some published bifurcation models [7] [8] but this quickly lead me to a particularly interesting model implemented by Pavlides et al. Their model fixes some of its parameters from known physiology and fits others (e.g. neuronal coupling weights) with a cost function directly from data. The cost function constrains the solution including penalizing for losses of known functionality when simulating a variety of lesions in the Basal Ganglia [9] [10].

If you are interested, the Pavlides et al. paper does an amazing job of explaining how Tachibana et al. physiological work on macaques, which involved blocking pathways between brain regions implicated in PD, has invalidated several formally proposed PD models. Their model claims to uphold appropriate behavior when simulating a variety of lesions in the Basal Ganglia. This is an important property I would like to capture in our model as well.

## 4.3 Causality Analysis

I am also inspired by PD models that leverage causality analysis. There are many recent models using Granger Causality (GC) [11] [12] or a more recent extension, Dynamic Causal Modeling (DCM) [13]. I just recently ran into Karl J. Friston’s impressive body of work on DCM and I am now staring down the barrel of yet another needed deep literature review [14] [15] [16] [17].

I think causality analysis needs to be incorporated into models like Pavlides et al. While understanding how neural populations are connected is powerful, DCM could help us understand the directional flow of information between neural populations. The difference between functional and effective connectivity.

### 4.3.1 Granger Causality Analysis

In a nutshell... with GC, one examines how to best predict the future of a neuron: using either the entire ensemble or the entire ensemble except a certain target neuron. The GC test is a statistical hypothesis test for determining whether one time series is useful in forecasting another, disambiguating correlations from causal relationships.

As extra motivation for this analysis, MEG is especially equipped to overcome the common challenges/pitfalls of time-series causality analysis such as [18]:

1. common reference
2. volume conduction

*Note:* There is much greater detail on this in my [literature review slides](#).

### 4.3.2 Dynamic Causal Modeling

Given that we want to model the network dynamics, DCM is a logical extension of GC that overcomes many of the shortcomings of GC in this setting.

Added benefits of DCM:

1. Unlike Bayesian Networks, the graphs used in DCM can be cyclic
2. Unlike Structural Equation modelling and Granger causality, DCM does not depend on the theory of Martingales, i.e., it does not assume that random fluctuations’ are serially uncorrelated.
3. DCM Bayesian predictions could overcome artifactual GC modulations induced by averaging event-related responses, and GC issues with phase coherence.

In contrast to many causal models, DCM does not look for statistical dependencies among measured time-series directly. Instead, it combines a biophysical model of the hidden (latent) dynamics with a forward model that translates hidden states into predicted measurements.



This provides a general framework, allowing us to plug in interchangeable models for the underlying neural activity given measured (hemodynamic or electromagnetic) responses over the sensors considered. The hidden states cover any neurophysiological or biophysical variables needed to form the observations and bayesian inversion furnishes the marginal likelihood (evidence) of the model and the posterior distribution of its parameters (e.g., neuronal coupling strengths).

## 5 Appendix: Extra Material

A verbose compilation of my thoughts from my literature review can be accessed [here](#).

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