

LAB ROTATION 02

1st Week Assignment

Şeyma Bayrak, Advisor: Philipp Hövel

11.09.2013

1 Nonlinear Dynamics of Neural Networks

1.1 Simple FitzHugh-Nagumo Model

$$\varepsilon \dot{x} = x - \frac{x^3}{3} - y \quad (1)$$

$$\dot{y} = x + a \quad (2)$$

x : activator, reproducing behavior of voltage during the course of a spike
 y : inhibitor, inhibiting production of x
 a : bifurcation parameter - threshold parameter: determines whether the system excitable ($a > 1$) or exhibits periodic firing ($a < 1$)

Bifurcation Analysis and Nullclines / Fixed Points:

- $\dot{x} = \dot{y} = 0$, x - *nullcline* : $x - \frac{x^3}{3} - y = 0$, y - *nullcline* : $x + a = 0$
- *equilibrium (fixed) point* : $(x_A, y_A) = (-a, -a + \frac{a^3}{3})$
- *Jacobian Matrix*, analyses the stability of fix points with eigenvalues
- $\lambda_{1,2} = \frac{1-a^2 \pm \sqrt{(1-a^2)^2 - 4\varepsilon}}{2\varepsilon}$, $\varepsilon = 0.005$, for $|a| > 1$ the fixed point is stable, for $|a| < 1$ unstable

Fixed point (x_A, y_A) : Separating excitable and inhibitory region.

Stability : resting towards (x_A, y_A)

Change of stability to instability: periodic oscillations

1.2 Related Figures

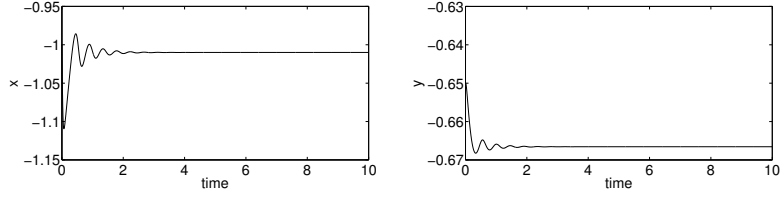


Figure 1, Simple FHN model, x and y plots as time series. $a = 1.01$, and $\varepsilon = 0.005$

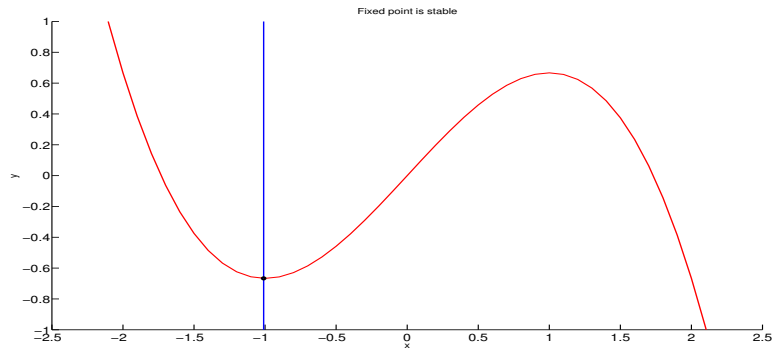


Figure 2, Simple FHN model, x versus y in state space $a = 1.01$, and $\varepsilon = 0.005$

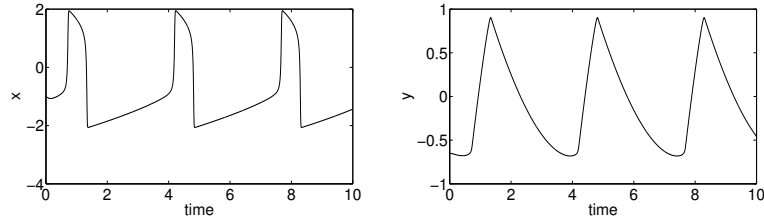


Figure 3, Simple FHN model, x and y plots as time series. $a = 0.97$, and $\varepsilon = 0.02$

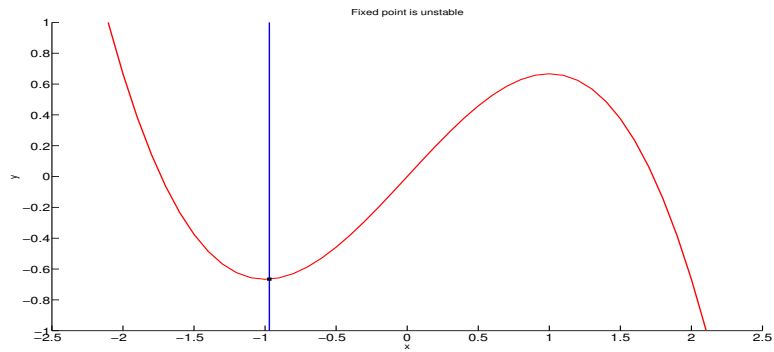


Figure 4, Simple FHN model, x and y plots as time series. $a = 0.97$, and $\varepsilon = 0.02$

1.3 Extended FitzHugh-Nagumo Model

$$\varepsilon \dot{x} = x - \frac{x^3}{3} - y \quad (3)$$

$$\dot{y} = x + a - \gamma y \quad (4)$$

Extended FHN model : now \dot{y} depends an additional linear inhibitory term γy
Bifurcation Analysis and Nullclines / Fixed Points:

- $\dot{x} = \dot{y} = 0$, x -nullcline : $x - \frac{x^3}{3} - y = 0$, y -nullcline : $x + a - \gamma y = 0$
- equilibrium (fixed) point : $@fsolve(x(x - \frac{1}{\gamma}) - \frac{x^3}{3} - \frac{a}{\gamma}) \rightarrow x(a, \gamma) = x_f$
- Jacobian matrix eigenvalues : $\lambda_{1,2} = \frac{(1-x_f^2-\gamma\varepsilon) \pm \sqrt{(1-x_f^2-\gamma\varepsilon)^2 - 4(\gamma\varepsilon x_f^2 + \varepsilon - \gamma\varepsilon)}}{2}$

Bifurcation point depends now not only a but also γ .

Fixed point can be now *stable*, *unstable*, *saddle*.

Let us make determinant of Jacobian matrix positive to eliminate saddle points:
 $\det(J) = \varepsilon(\gamma(x_f^2 - 1) + 1) > 0$, then we should choose $0 < \gamma < 1$

1.4 Related Figures

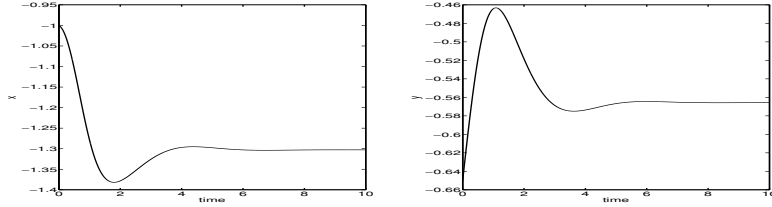


Figure 5, Extended FHN model, x and y plots as time series. $a = 1.30$, $\gamma = 0.005$ and $\varepsilon = 0.4$

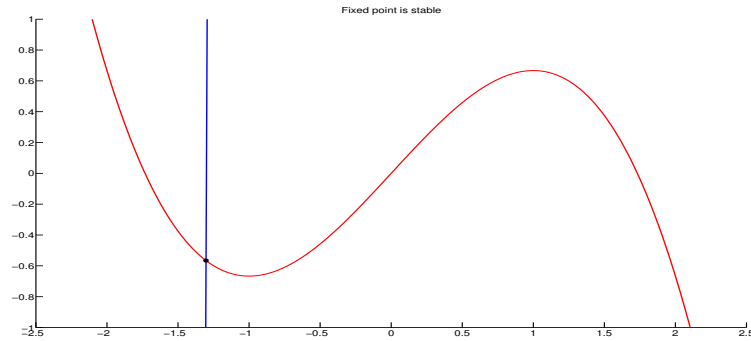


Figure 6, Extended FHN model, x and y in state space. $a = 1.30$, $\gamma = 0.005$ and $\varepsilon = 0.4$

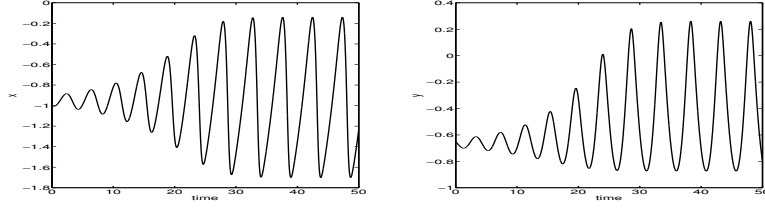


Figure 7, Extended FHN model, x and y plots as time series. $a = 0.95$, $\gamma = 0.005$ and $\varepsilon = 0.4$

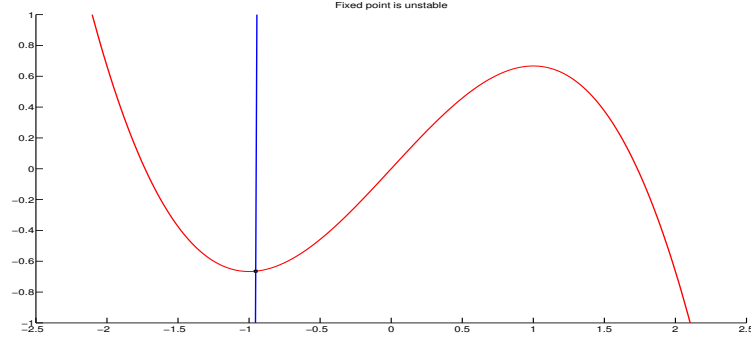


Figure 8, Extended FHN model, x and y in state space. $a = 0.95$, $\gamma = 0.005$ and $\varepsilon = 0.4$

2 Execution of Python and MATLAB Scripts

- Create a matrix with threshold:

```
python threshold_matrix.py <arg1> <arg2>
python threshold_matrix.py A.txt 0.5
```

that means the matrix A is converted with threshold value 0.5 into another matrix. The elements of new matrix are $[0, 1]$, the values below the given threshold are 0, otherwise 1. The new matrix is saved as $A_{r0.5}.dat$

- Simulation of neural activity: time evolution of activator and inhibitor

```
python fhn_time_delays.py <arg1> <arg2> <arg3> <arg4>
```

< arg1 > : f_{ij} , functional connectivity matrix
 < arg2 > : d_{ij} , matrix; distances between nodes in brain
 < arg3 > : c , coupling constant
 < arg4 > : D , noise strength

- [VUK13] - some theoretical approach to the command-line above:

$$\dot{u}_i = g(u_i, v_i) - c \sum_{j=1}^N f_{ij} u_j(t - \Delta t_{ij}) + n_u \quad (5)$$

$$\dot{v}_i = h(u_i, v_i) + n_v \quad (6)$$

where c is coupling strength, f_{ij} is the connectivity matrix, Δt_{ij} is time delay due to finite signal propagation velocity between nodes, n_u is the noise factor. Δt_{ij} is calculated as $\Delta t_{ij} = \frac{d_{ij}}{v}$, distance matrix divided by velocity and noise factor is includes the noise strength D .

The functions g and v are modeled very similar to FitzHugh-Nagumo model introduced before:

$$\dot{u} = g(u, v) = \tau(v + \gamma u - \frac{u^3}{3}) \quad (7)$$

$$\dot{v} = h(u, v) = -\frac{1}{\tau}(u - \alpha + bv - I) \quad (8)$$

- The outcome of the *fhn_time_delays.py*:

$$simfile = \begin{bmatrix} 0 & u_{11} & v_{11} & u_{21} & v_{21} & . & . & . & u_{N1} & v_{N1} \\ dt & u_{12} & v_{12} & u_{22} & v_{22} & . & . & . & u_{N2} & v_{N2} \\ 2dt & u_{13} & v_{13} & u_{23} & v_{23} & . & . & . & u_{N3} & v_{N3} \\ 3dt & . & . & . & . & . & . & . & u_{N4} & v_{N4} \\ 4dt & . & . & . & . & . & . & . & u_{N4} & v_{N4} \\ . & . & . & . & . & . & . & . & . & . \\ t_{max} & . & . & . & . & . & . & . & u_{NN} & v_{NN} \end{bmatrix}$$

- Observe attractors's activity as time series

MATLAB >> calcBOLD.m >> calcBOLD('simfile')

The program *calcBOLD.m* firstly eliminates all the u_i time series from the input i.e. *simfile* and plots the total time versus all the u_i series.

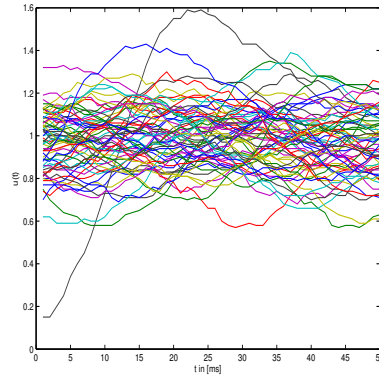


Figure 9, Extended FHN model, x and y in state space. Threshold applied to f_{ij} matrix is $r = 0.5$, coupling constant $c = 0$, noise strength $D = 0.05$ and velocity of signal propagation $v = 70m/s$.

- Simulated Bold activity with Baloon-Windkessel model

The resulting time series of the neural activity u is used to infer the BOLD signal observed in fMRI data via Baloon-Windkessel model. The puspose of the project is to be able observe how well the simulated BOLD signal correlates with the emprical fMRI signal. Here is an example of how the simulated BOLD signal might look like.

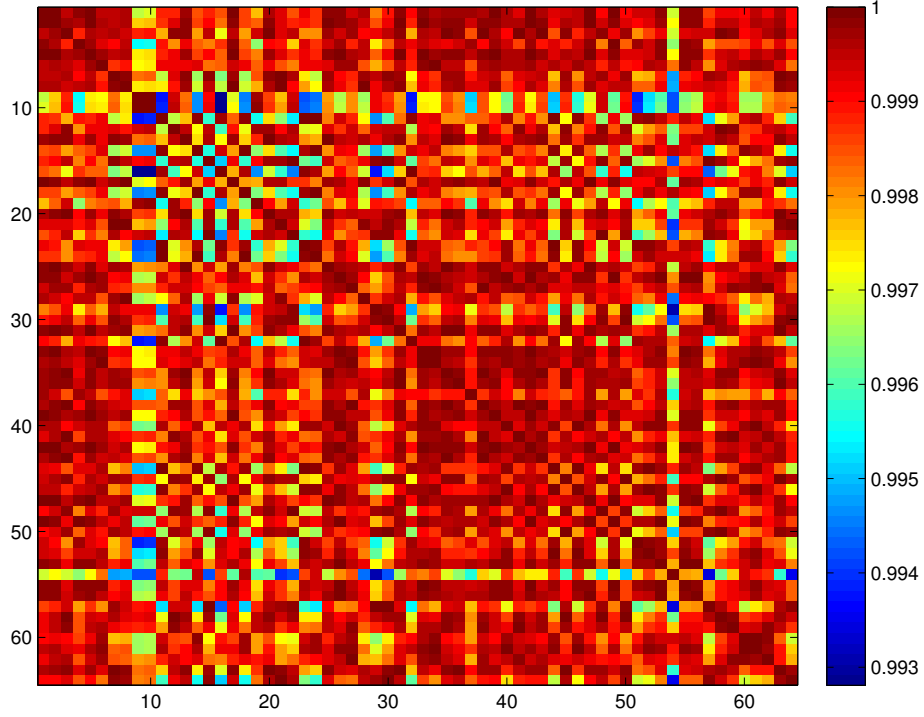


Figure 10, correlation matrix of sBOLD, coupling strength $c = 0$ and noise strength $D = 0.01$, $v = 70m/s$.

3 How the FHN model is modified in those papers: [VUK13], [GHO08], [GHO08a]

The FHN model used in [VUK13] has been introduced in equation 7 and 8, let us compare them with equations 1 and 2.

$$\begin{aligned} \dot{x} &= \frac{1}{\varepsilon} \left(x - \frac{x^3}{3} - y \right) & \dot{y} &= x + a - \gamma y \\ \dot{u} &= \tau \left(\gamma u - \frac{u^3}{3} + v \right) & \dot{v} &= -\frac{1}{\tau} (u - \alpha + bv - I) \end{aligned}$$

- Instead of $\frac{1}{\varepsilon}$ in equation 3, a time constant τ is introduced in equation 7. Beside of changed parameters, the most important difference is that, the inhibitor \dot{v} depend in equation 8 on a negative attractor $-u$ and there occurs also one additional term called I ; the external stimulus, but that is assumed to be 0 in papers.

3.1 Related Figures

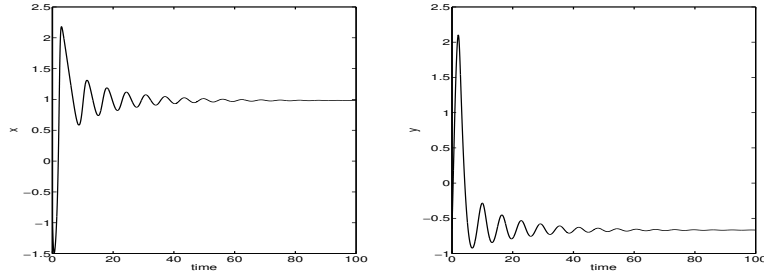


Figure 11, $\alpha = 0.85$, $\gamma = 1.0$, $b = 0.2$, $\tau = 1.25$

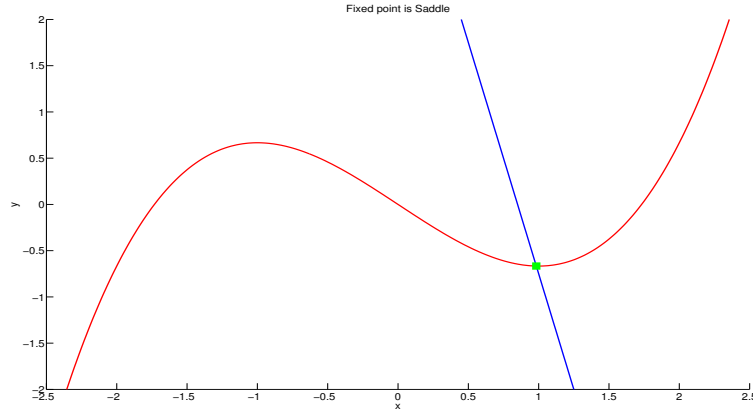


Figure 12, $\alpha = 0.85$, $\gamma = 1.0$, $b = 0.2$, $\tau = 1.25$