# Brain Modeling:

A very simple Network model of the Human Brain

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#### Presentation Overview

- 1. Introduction: Brain, Neuron and Connections
- 2. Objectives:
- 3. Model and Implementation
- 4. First Results
- 5. A couple of comparisons
- 6. Conclusion
- 7. Limitations
- 8. Futur work

## 1- Introduction: Brain, Neuron and Connections

- The nervous system is a set of a billion of neurons that communicate with each-other through a billion of billions of synaptic connections. It is apparent to a set of electrical cables connecting nodes, forming a huge network. Most published studies on diseases of the nervous system face the problem of complexity of the nervous system as its whole.
- It is difficult to tell accurately the reason of a dysfunction; this is the case of Autism: Although symptoms of the disease are known, there is no consensus on the reasons of such a phenomenon. Some authors talk about spend energy related to long-distance connections, others talk of poor synchronization between the left hemisphere and the right hemisphere, others talk about poor quality of the gray matter, namely, the myelination of axons, key for inter-regional communication, ... Those are known through Infra-red and a bunch of scanning tools.
- It is very hard for Biologists to figure out what is really going on. We then need model with which we can manipulate different possibilities and observe the results.

# 2- Objectives

- Multi data processing with network implementation in order to mimick the Humain Brain, and try to identify potential reason for dysfunctions like Autism, Alzheimer, Parkingson, ...
- Build 2 X 3D networks with as much as neuron possible
   (Abnormal Network → Dysfunction Vs Normal Network → Safe).
- Analysing :
  - Simple Correlations between parameters such as ATP and length all along the signal processing.
  - How does a specific parameters change between those 2 Networks can make a difference
  - ... (if time allows)

#### A) Model

- I choose not to use in a first step the huge and enormous equations that rules neurons input and signal transfer we may see in the literature, in Electrical and physics. Instead, I used the simplified model of neuro-psychology, which is easier to implement. However, those equations in Electrical can be included in the model.
- Action Potential that decide whether a Neurone fires or not is not based on Boolean value but based on both time and neurotransmitter abundance.

```
∑ ( neurotrans X action ) = received inpulse
if received impulse >= Threshold then → Fire
```

• I consider the time in the entire Network, and this time is based on the time recorded at signal input (button press).

#### A) Model

- I started with 64 neurons in the Network (just to avoid long time running more than 1000 Neurons which is the final goal for now).
- I have 1 Main\_File that store the information on all Neurons (ID, myelinated or not, strength of myelin, if it is an Activator, if it is an Inhibitor, coordinates X, Y, Z in 3D space, Action\_Potential\_Threshold).
- I designed a graphical interface that allows to directely Update that File (In a random way, and also in specific way : Ex: one can choose how one want the Neuron\_X to behave with its neighboor).
- I have 64 Files That store each Neuron interaction (1 File per Neuron).
- With the graphical interface, one can directly update the behavior of any Neuron with its Neighboors. Then, there is 2 ways to update:
  - ✓ Either by area (you update interactions of neurons located in the selected area: the best choice for general analysis).
  - ✓ Either by ID (you update interactions of neurons taping is ID: this would be the best choice for very-very deep analyse, but extremely long to do with 1 million neurons).
- Regarding Neuron Interactions: it is information such as: Neighboors (consequentelly I have 64 lines), behave =
   Activation or Inhibition or Nothing (this depend on the previous Main\_file), Strenght of that interaction, duration of
   signal transmission, length with the Neighboor, ATP spent while the signal is travelling, Current time of signal
   perception.
- In the next slide, all stuff put in color are not random; They depend on the interaction with the neighboors on the Network.

#### A) Model

- Activation or Inhibition or Nothing (this depend on the previous Main\_file): When a Neuron is an activator, it is a (+1). When it is a inhibitor, it is a (-1). When He receives a signal, He transfers that signal to its neighboors. Then there is 2 cases: He may transfer a negative signal (inhibitory signal) or a positive signal (activator signal).
- If He transfers a positive signal, and that neighboor fires, that firing strengthens their relationship; in case of non-firing, it diminish their relationship
- If He transfers a negative signal, and that neighboor fires, that firing diminishes their relationship; in case of non-firing, it strengthen their relationship.
- Strength of that interaction: Dependent on quantity of neurotransmittor.
- Duration of signal transmission: depends on both distance between 2 neurons, and the force of the Myelin layer; The longer is the distance, the longer it takes the signal to travel. The thicker is the myelin layer, the faster the signal can travel.
- Length with the Neighboor: It s based on the Cartesian distance between 2 neurons. I added 0 10% margin because neural connections are almost never straight.
- ATP spent while the signal is travelling: Inversely proportional to the thickness of the myelin and the length of the axon.
- Current time of signal perception: This allows Matlab to follow the signals. Actually, there are no real signal: those are just timing to follow an imaginary signal through the network.

Edit text allow set-up Action potential Theshold, incrementation Vs decrementation jump

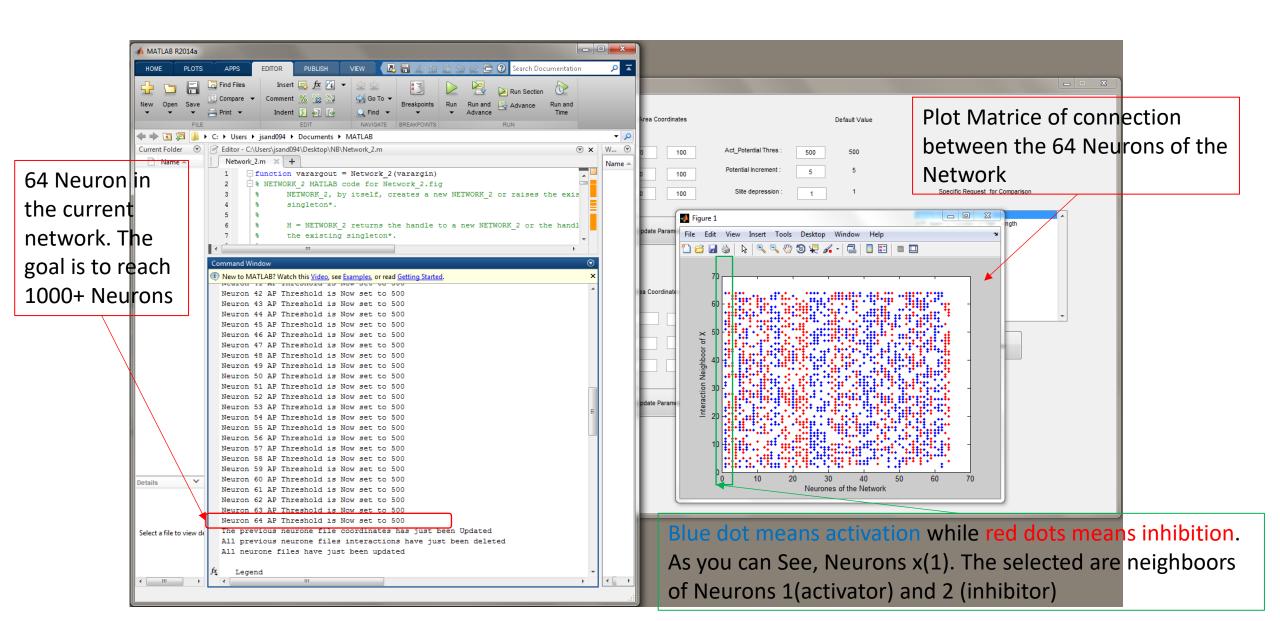
B) Implementation : done with

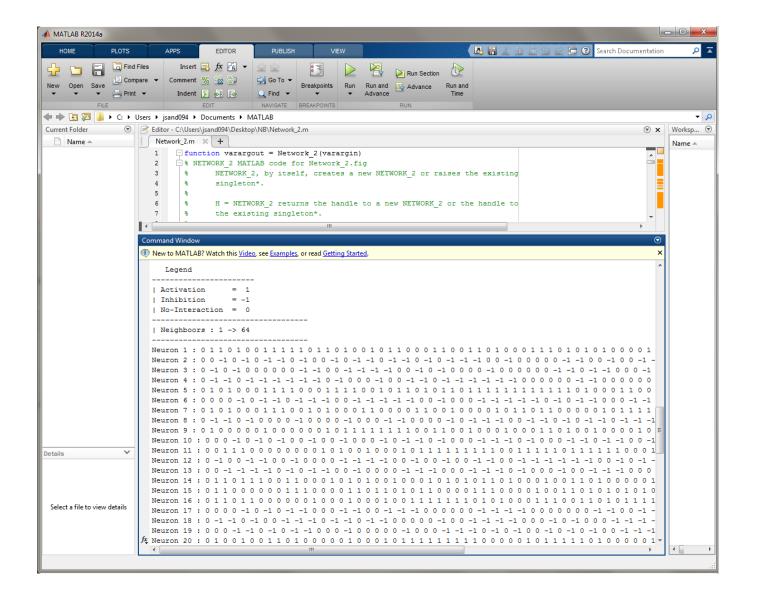
Edit text allows to modify specific areas

Matlab Network\_2 Area Coordinates Default Value Set Selected Neurons to be all Excitators Set Selected Neurons to be all Inhibitors X1...Xn Act\_Potential Thres Set AP to be reached at 100 Abnormal Network Set AP to be reached at 1000 Potential Increment Y1...Yn Set AP Threshold to another value Set Potentiation Increment to another value Starting point for Set slith depression to another value Z1...Zn Slite depression Specific Request for Comparison Set Neuro Abund the same value for all its Neighbo Set Myelin strength to the Input Value (1 - 100) each network after ATP Spent in Function of Path Lengtl Update Parameter Modify Specific Interaction Between 2 points --> loading the application Area Coordinates Default Value Set Selected Neurons to be all Excitators Act\_Potential Thres X1...Xn Set Selected Neurons to be all Inhibitors Set AP to be reached at 100 Set AP to be reached at 1000 Potential Increment Send Impulse Set Potentiation Increment to another value Set slith depression to another value Z1...Zn Set Neuro\_Abund the same value for all its Neighbo Slite depression Set Myelin strength to the Input Value (1 - 100) Once you select an item on the lisbox, Update Parameter Modify Specific Interaction Between 2 points --> you fill the edit text, then press There are 2 ways to send an inpulse. So far I am using update the one at left, but once the work will be completely done, we will have to use only the one at right

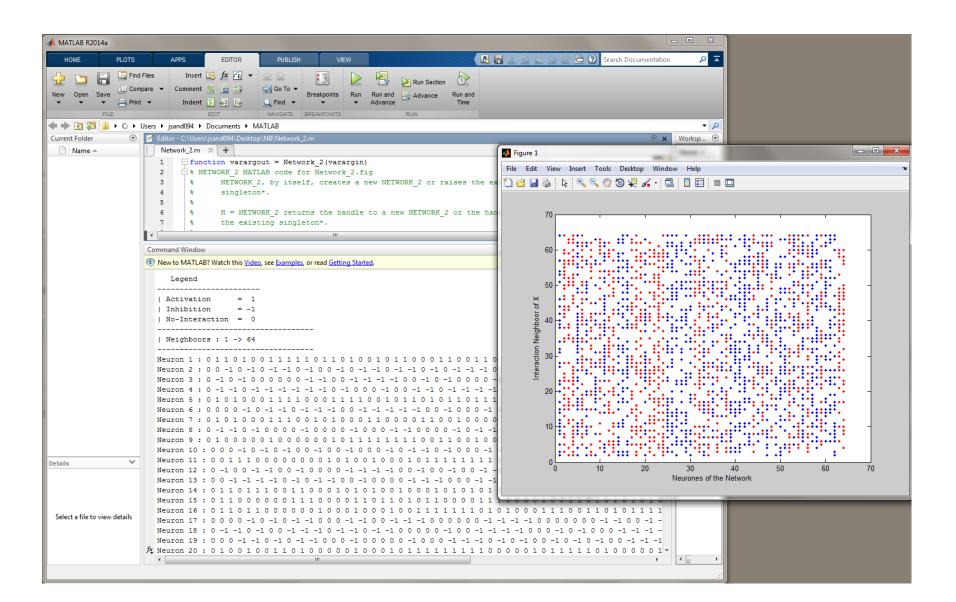
This button is to modify a specific interaction between 2 points. It opens a other frame. I will go there in a couple of slides.

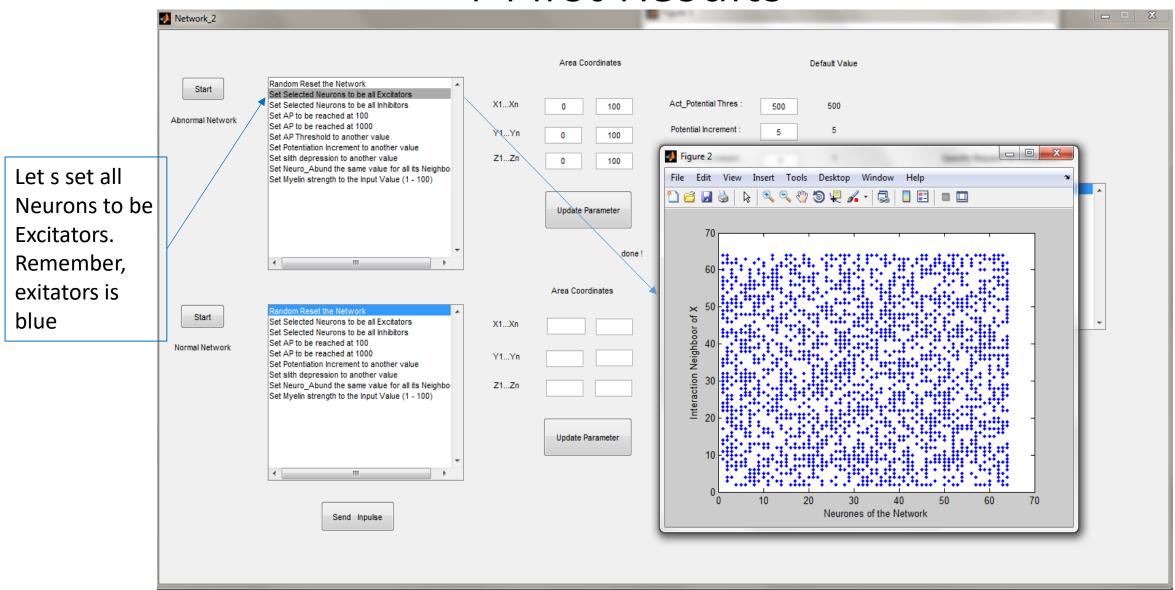
Then selected, it allow me to specify which test I want to perform



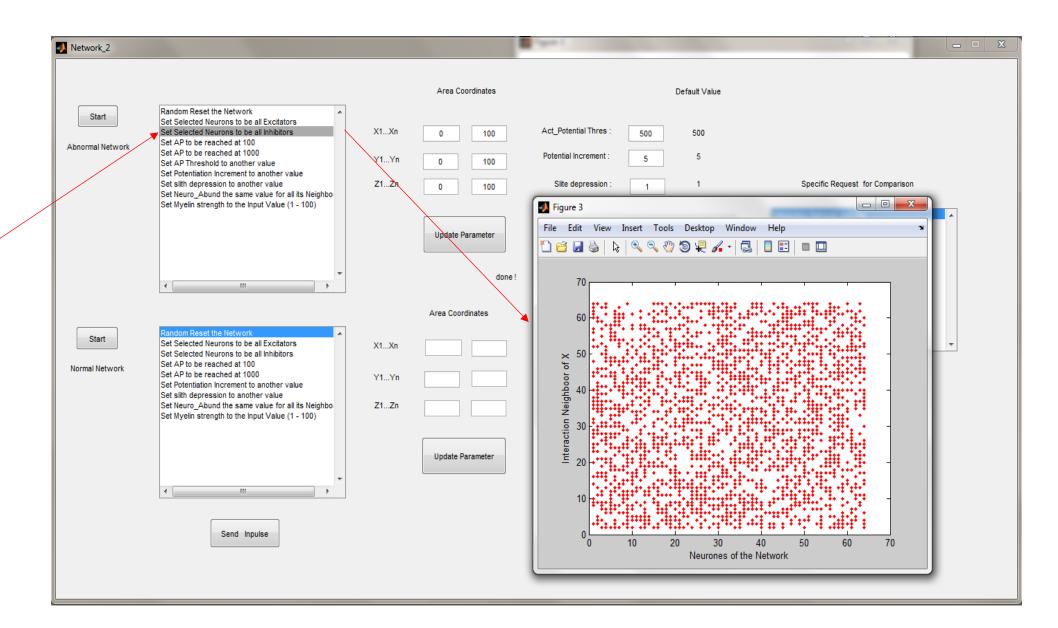


Another view of the Connection Matrice

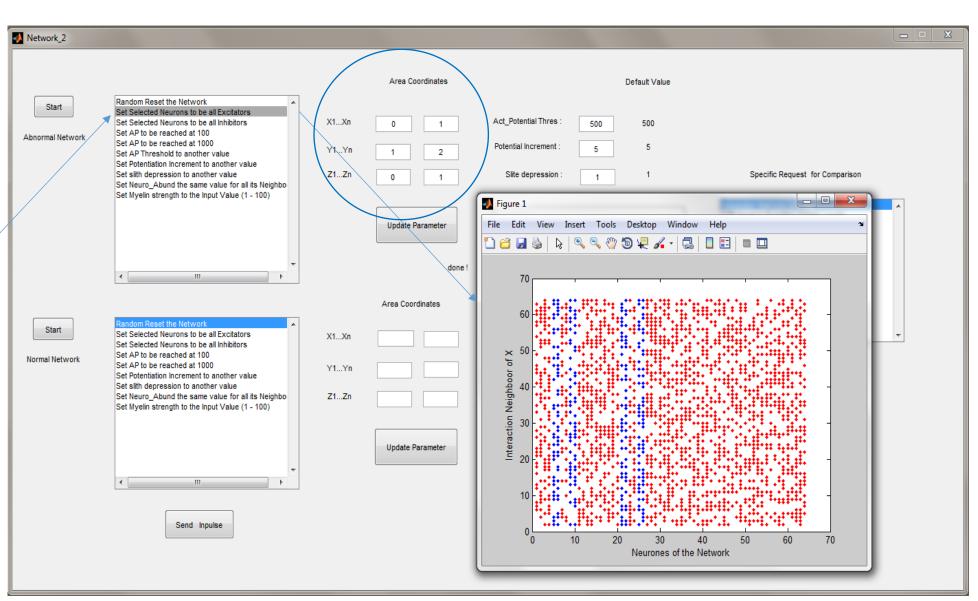


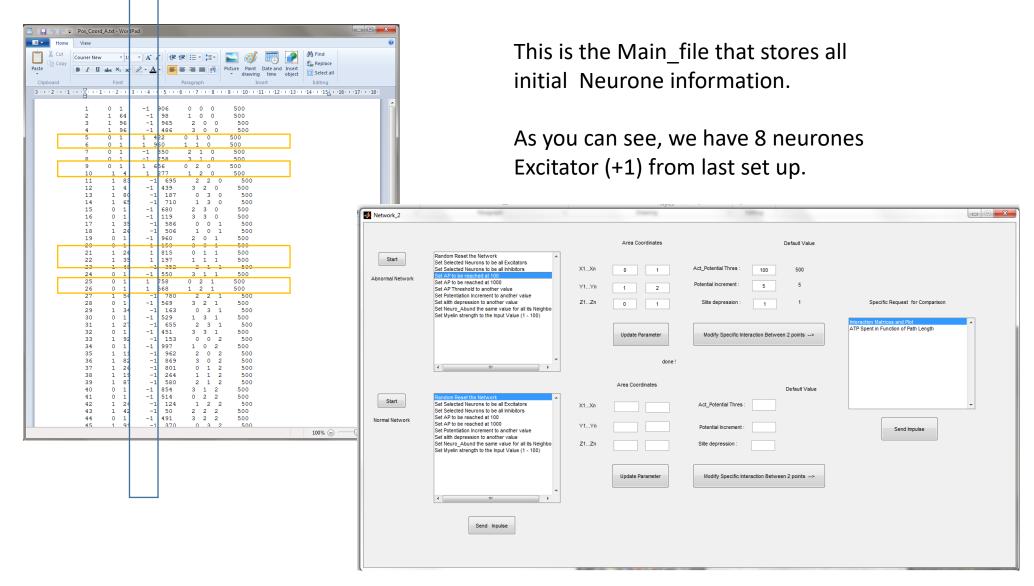


Let s set all neurons to be inhibitors. Remember, inhibitors is red.

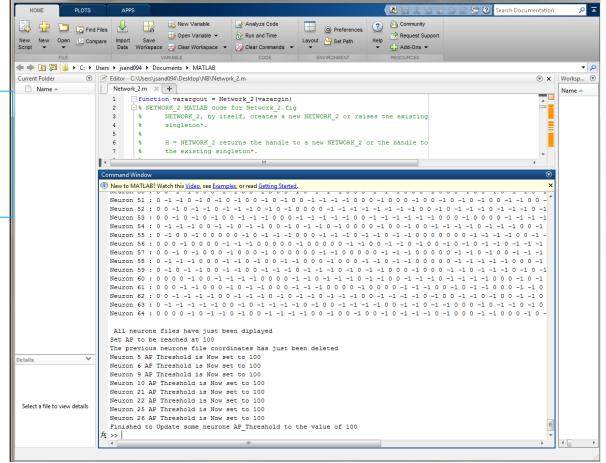


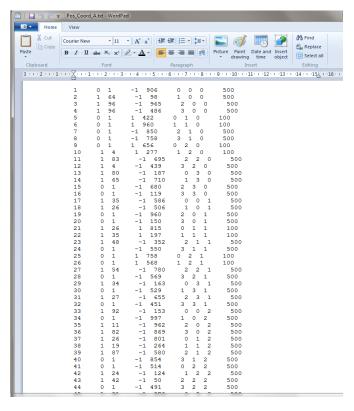
Let select only neurons which coordinates are inside selected Area, to be Excitators.





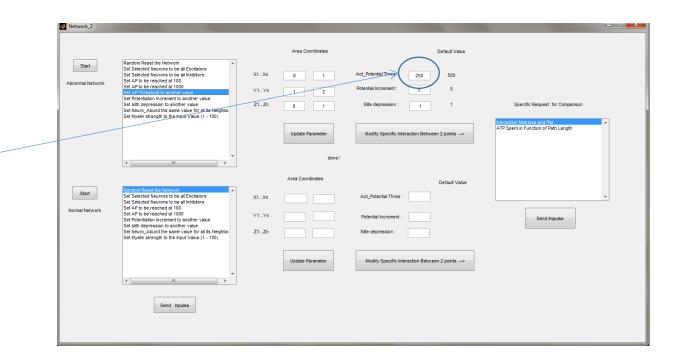
Set selected Neurons Action Potential to be reached at 100 neuro-transmitors. ▲ MATLAB R2014a

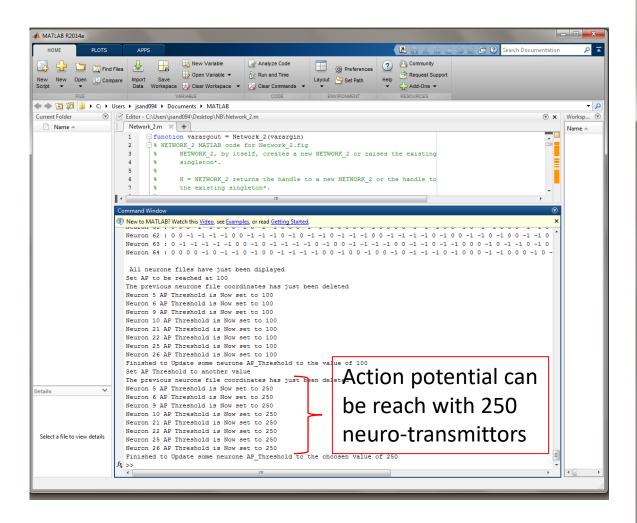


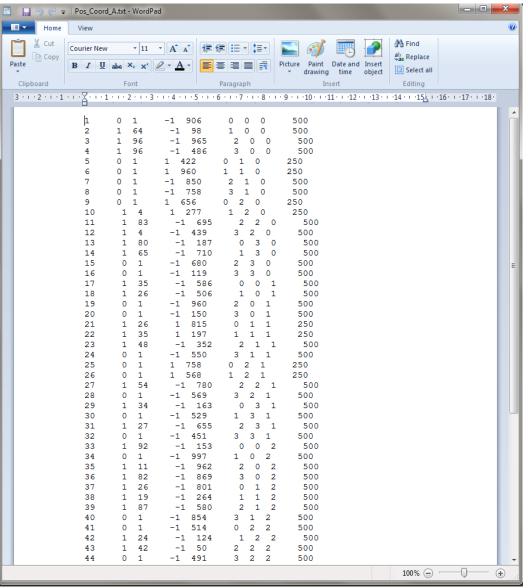


Set selected
Neurons Action
Potential to be
reached at
another number
of neurotransmitors.

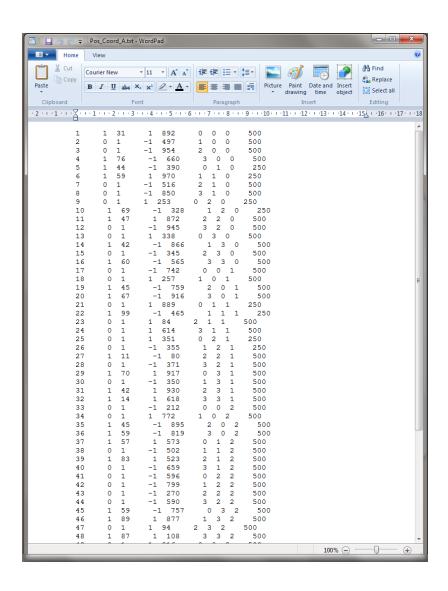
Random value here is 250

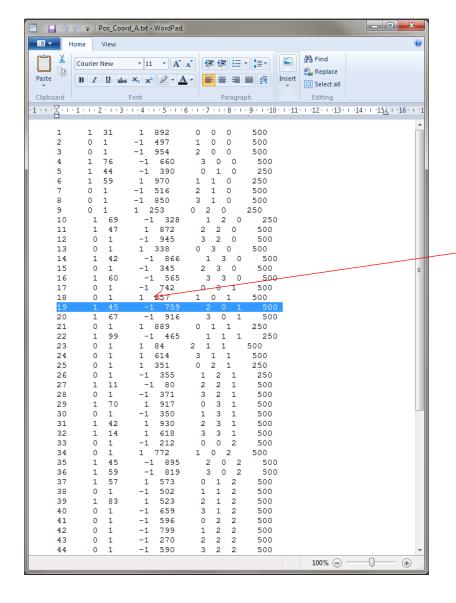






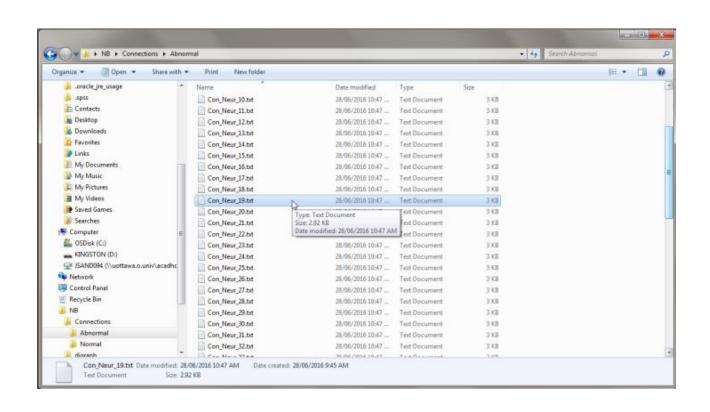
# Let s randomly take Neuron 19



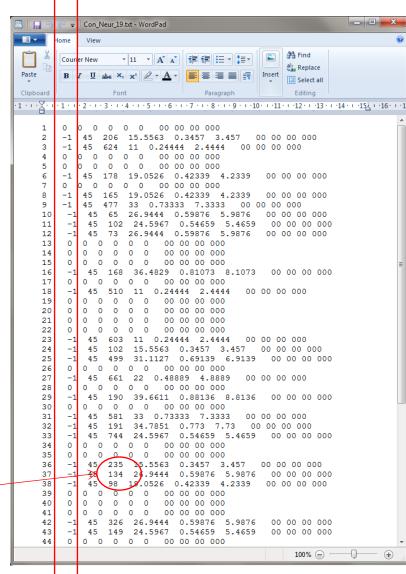


It is an inhibitor (-1)

He has either 0 or 1 inhibition connection with its neighboors



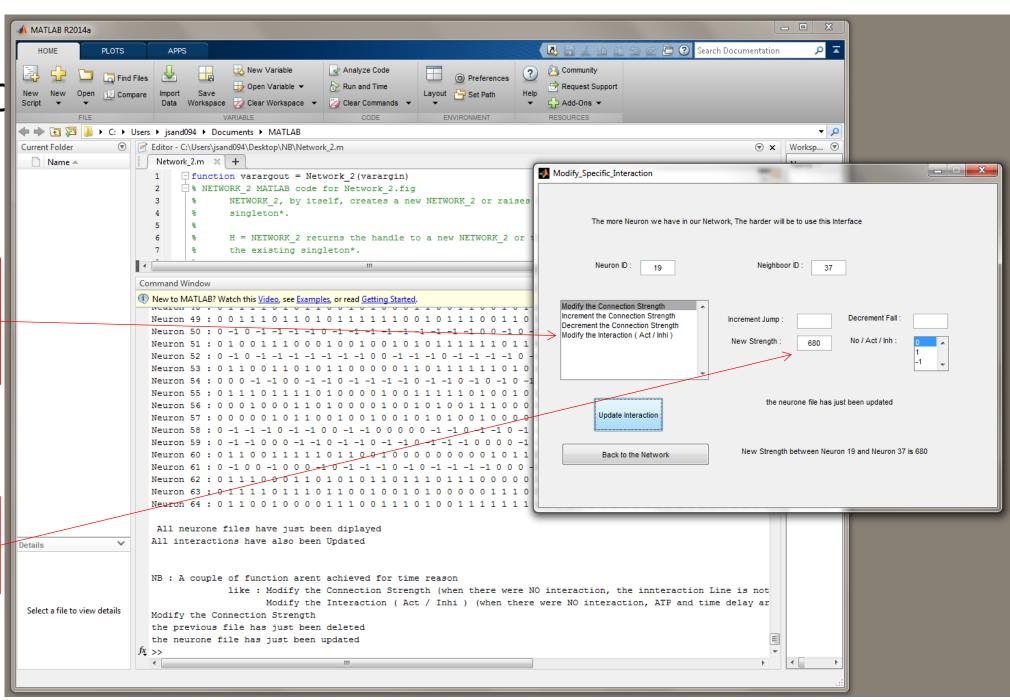
Its current strength with Neuron 37 is 134



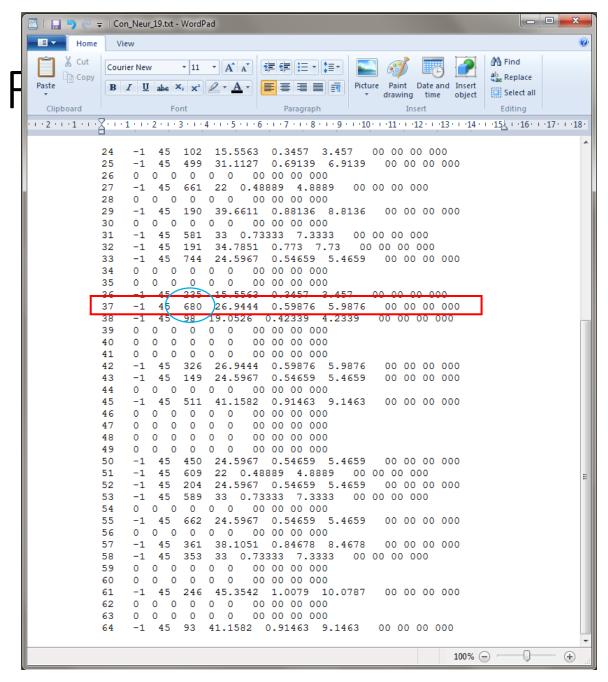
#### Mod

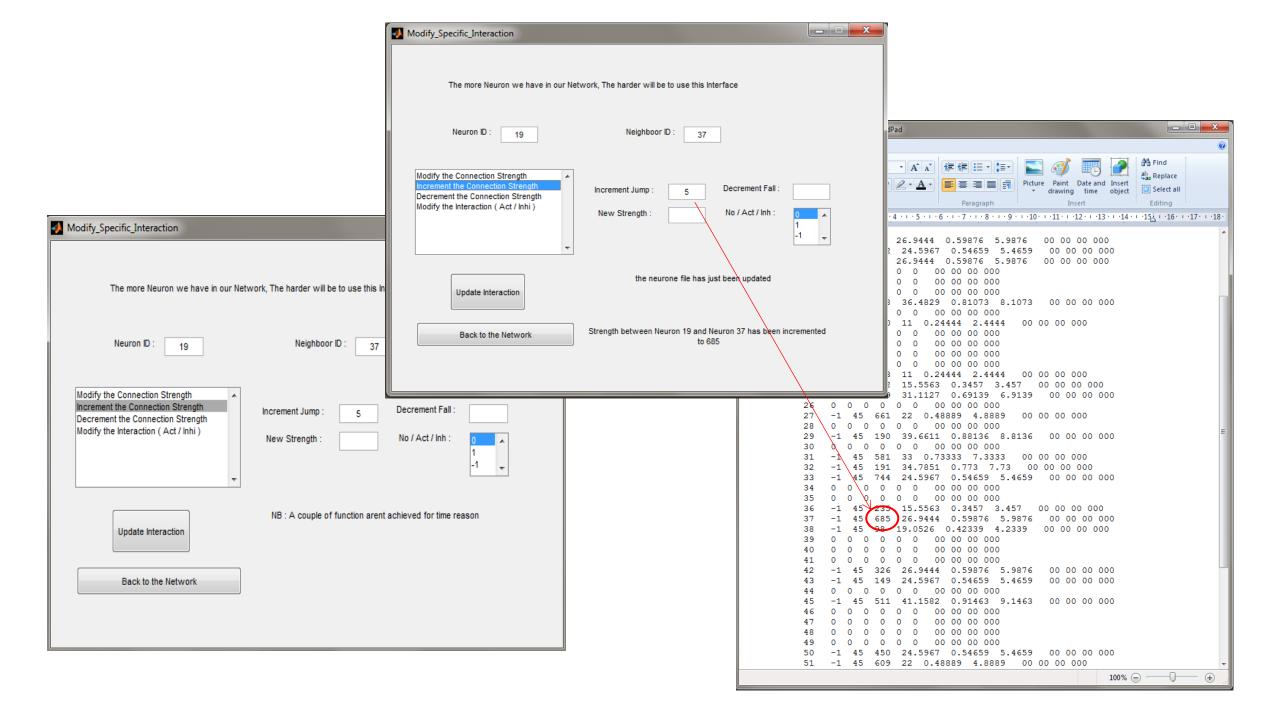
The last request in the listbox is not achieved for time reasons.

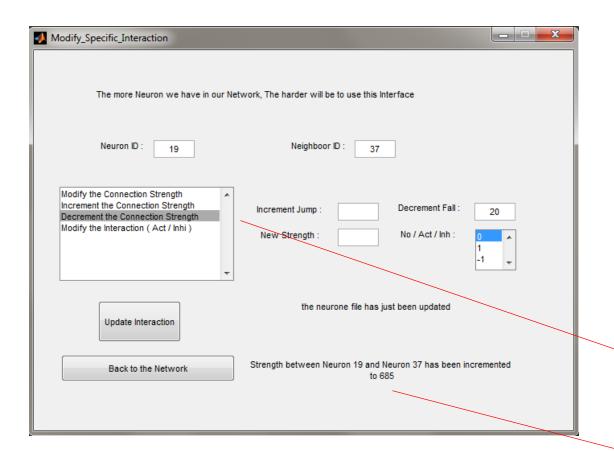
We update that strength to 680



# Back to the

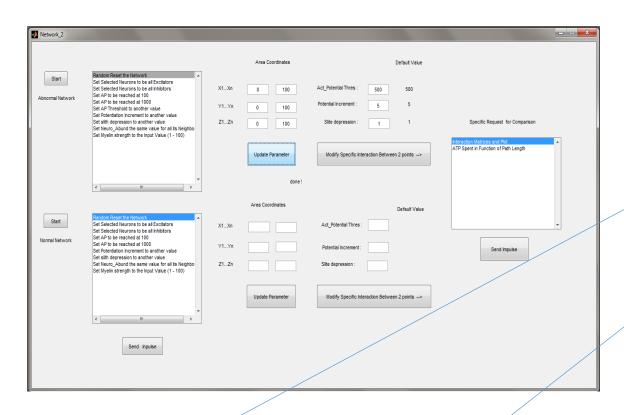




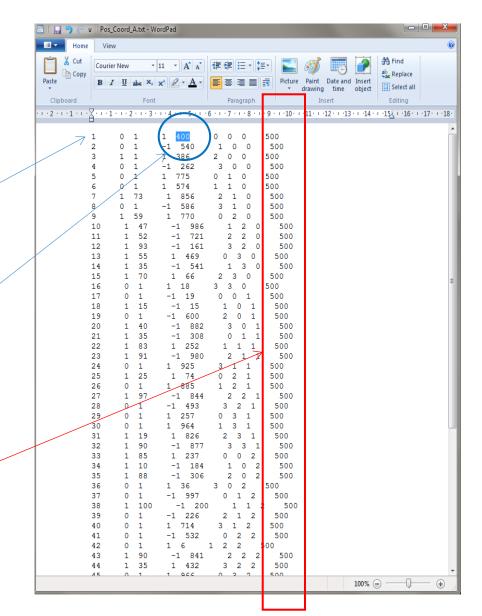


↑ Modify_Specific_Interaction	X
The more Neuron we have in our Network, The harder will be to use this Interface	
Neuron ID : 19	Neighboor ID: 37
Modify the Connection Strength Increment the Connection Strength Decrement the Connection Strength Modify the Interaction ( Act / Inhi )	Increment Jump:  Decrement Fall:  20  No / Act / Inh:  1 -1
Update Interaction  Back to the Network	Strength between Neuron 19 and Neuron 37 has been decreased to  665

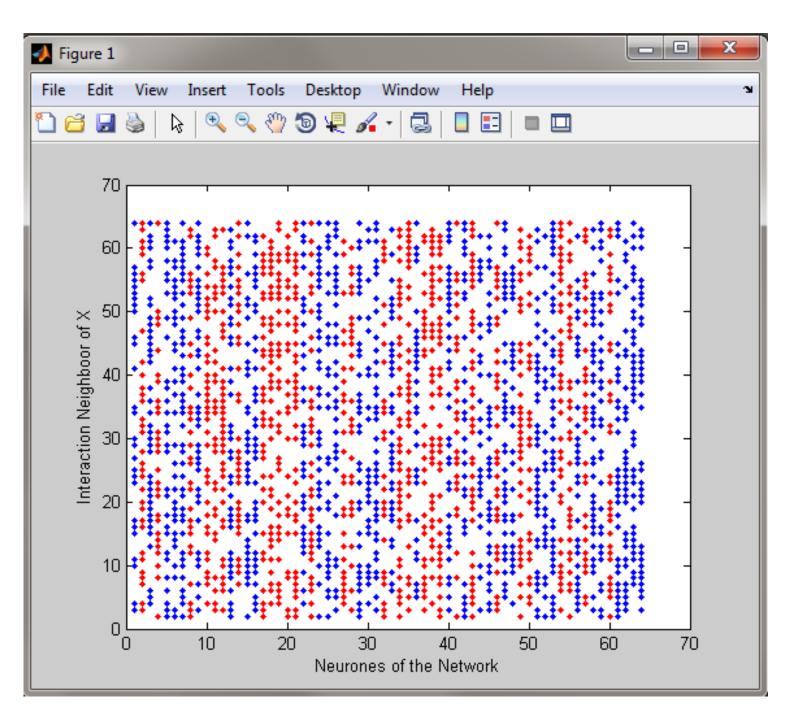
# Let s go back to the Network

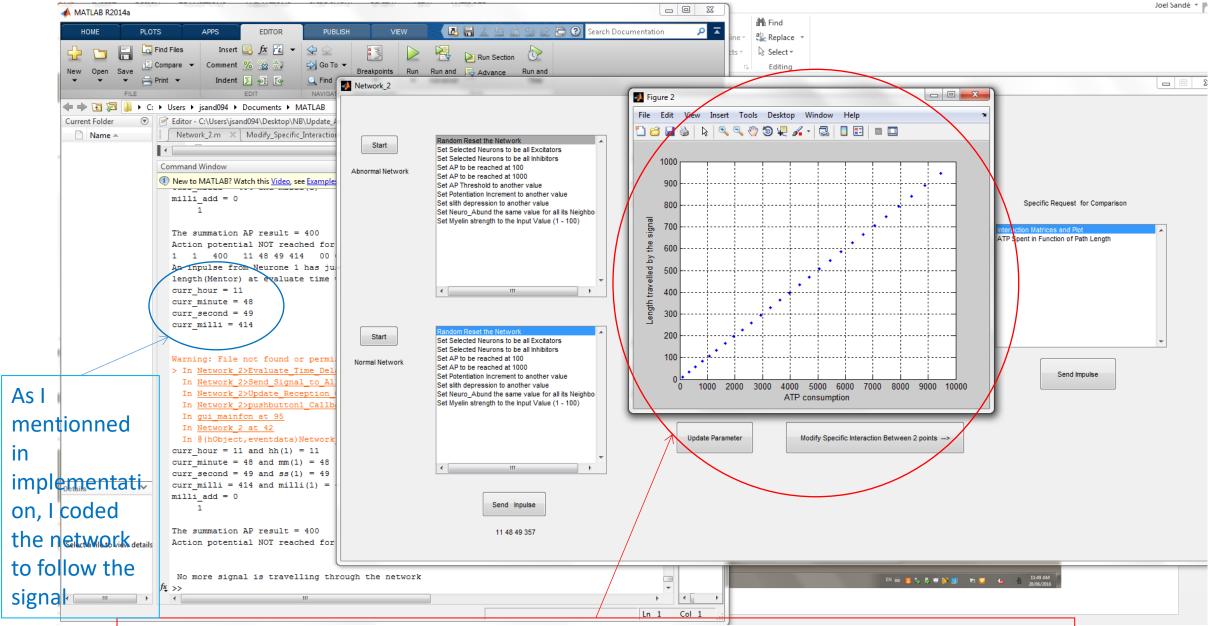


When one submit an impulse, the Neuron1 is the first neuron that it: Receptor Neuron. The last column is the action potential threshold. When neuron 1 will send its signal to its neighboors, for this simulation, I manually put its neurotransmitors (the power with which he excites its neighboors) to 100. In that case, he cannot excite none of its neighboors which threshold is 500 > 100.



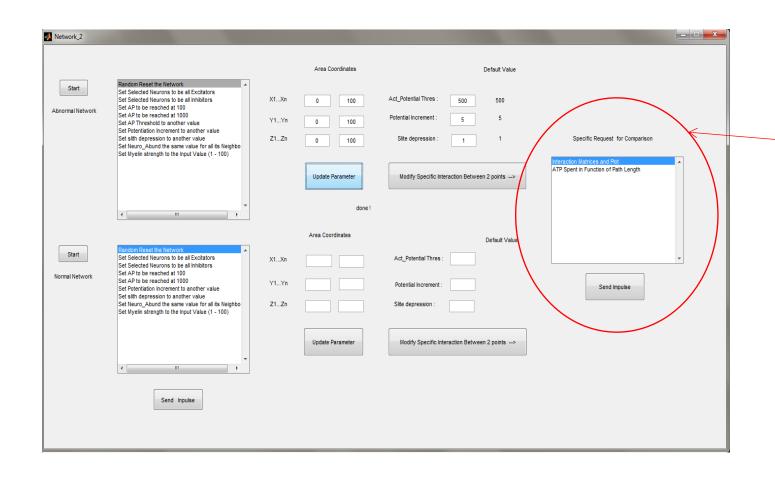
After updating the network, this is the interactions profile for this simulation





When the myelin is not altered, and we consider one neuron at the time, we can see a direct relationship between ATP consumption and Length travelled by the signal.

# 5-A couple of comparisons



This part is to compare the Normal Network and the Abnormal network. Right now, white I am writing this, I am reviewing a function that unfortunately saturates Matlab while running the whole simulation, because of iterations error. Matlab stores too much data, and stop running after.. 5 - 10 min. I will be able to do more simulations once I fixe that problem.

Current work

#### 6-Conclusion

• The conclusion we can make from slide 27 is that: When the myelin is not altered, and we consider one neuron at the time, we can see a direct relationship between ATP consumption and Length travelled by the signal.

 Not enough results on the network processing itself because of time constraints.

• However, session 4 provides good results regarding the network implementation.

#### 7-limitation

- As I don t have huge computer to do a multithread,
- 1) I cannot run simultanely both Abnormal and normal Network. I have to run one network at the time.
- 2) As I don t have huge computer to do a multithread, I can only send one signal and wait for that signal to disappear completely before I send another one

Solution: To use Supercomputers to do the simulations

#### 8-Futur work

- I found a function named Digraph available in Matlab 2016. A powerful function that :
  - ✓ gives the shorter path in a given network, from a point A to a point B,
  - ✓ topology that allow to connect neighbour neuron in specific pattern,
  - ✓ ... a lot of new stuff we can do, such as representing areas for the frontal lobe, visual lobe, ...
- Run more simulations varying different parameters :
  - ✓I can base the neuroplasticity on function available via matlab 2016 in addition to what I already did.
  - ✓ As the brain is lasy, I can easily choose in a simulation to keep only paths that spend the less energy possible and see how it impacts the network topology, using functionalities available on Matlab 2016.

### 8-Futur work

• I have a solution to implement data memory, and even, artificial intelligence within the system (algorithms based on this work). We can perform reminding based on synaptic-connections. What is interesting with those algorithm, they can give you (in output) only what you input, starting at a completely stupid stage : at the beginning there is no output possible until you start putting repetitively the same inputs.

### 8-Futur work

• I believe that to understand how disease like Alzheimer and Autism (memory disease) work, we have to get that kind of algorithm coded inside a network (and make difference between different compartments of the brain), and if that works well in the simulations, we can understand far beyond ATP and length issue, but also at the connection level.

→ this is not for today ☺