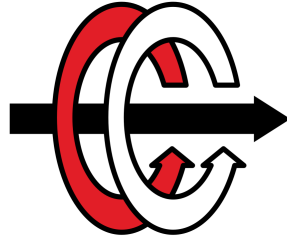




ORTA DOĞU TEKNİK ÜNİVERSİTESİ
MIDDLE EAST TECHNICAL UNIVERSITY



EE400 Summer Practice Report

Intern: Ahmet Akman 2442366

Supervisor: Dr. Johannes Zierenberg

Supervisor Contact: johannes.zierenberg@ds.mpg.de - +495515176475

Assigned Faculty Member: Prof.Dr. Engin Tuncer

Institution Name: Max Planck Institute for Dynamics and Self-Organization

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Electrical and Electronics Engineering Department

Middle East Technical University

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1 Introduction

2 About Institution

2.1 Institution Name

Max Planck Institute for Dynamics and Self-Organization.

2.2 Institution Location

Max Planck Institute for Dynamics and Self-Organization Am Faßberg 17 37077 Göttingen Germany

2.3 General Description

The Max Planck Institute for Dynamics and Self-Organization, located in Göttingen, Germany, is a prominent research institution primarily focused on the investigation of complex non-equilibrium systems, particularly within the fields of physics and biology. Its historical roots trace back to 1911 when Ludwig Prandtl initiated the establishment of a Kaiser Wilhelm Institute dedicated to the study of aerodynamics and hydrodynamics. This initial effort led to the formation of the Aeronautische Versuchsanstalt in 1915, which later evolved into the Kaiser Wilhelm Institute for Flow Research in 1924. In 1948, it became a part of the Max Planck Society. In 2003, it underwent a name change and became the Max Planck Institute for Dynamics and Self-Organization. Presently, it stands as one of the 80 institutes under the auspices of the Max Planck Society, contributing significantly to the understanding of intricate dynamic systems.

2.4 Organization Structure

The organization structure of the institute is given in Figure 1. I was part of the group led by Prof.Dr. Viola Priesemann which is indicated as *italic* on 1.

3 Project Background and Motivation

The research project for the internship is constructed upon the following set of previous research. In the field of neuroscience [4], initiated the idea of inhibition is a necessary part of a neuronal network for stability of the network. Then, the impactful paper [1] explained the dynamics of a sparsely connected random network with fixed parameters. Also, the role of inhibition was similar to the [4]. The phases of the network through these dynamics are also stated. These states can be summarized as follows also given in Figure 2. Synchronous regular (SR) is the state where the whole network is blinking in a synchronized way. So, the neurons are synchronized with each other, exciting each other together, and regular activity is observed. Synchronous irregular (SI) is the state where the individual neurons show irregular activity, but the global activity is still regular. Therefore, the rate of the random

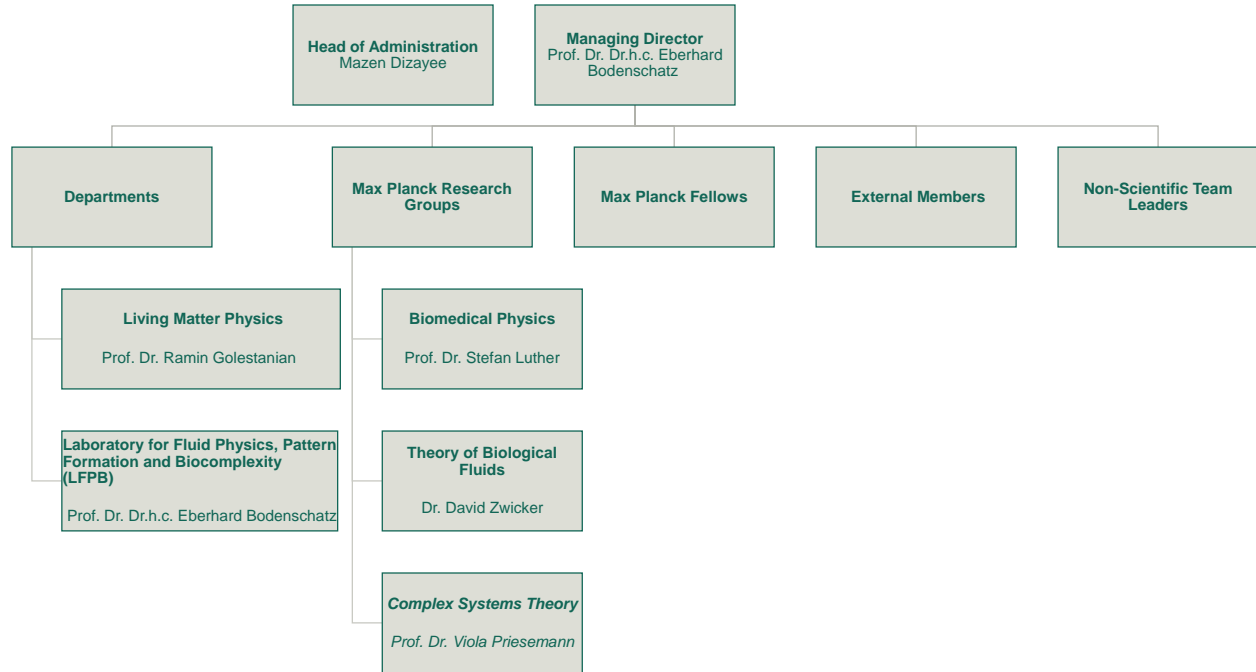


Figure 1: Basic organizational structure of MPI-DS.

network is still blinking in some sense. SI-fast and SI-slow characterize this as two different dynamical states. The asynchronous irregular state is the desired stable state where both the individual (per-neuron) and global activity show irregularity, that is, to small average overall activity not-blinking global network. [1] has been considered as a guiding baseline in the field, and many other network descriptions built upon this idea, including inhibition as a factor for stability. It is important to note that the [1] employs a fixed set of parameters on a completely randomly connected network.

3.1 Hypothesis

In neurobiology, one can classify the neurons as excitatory and inhibitory. A typical excitatory neuron is called a pyramidal neuron, whereas there are more than 20 mainstream types of inhibitory neurons. The idea is that the role of inhibition might be something other than just stabilizing the network. So, we hypothesize that only excitatory networks can be stable. To achieve the stability, the structural connectivity of the network should be reconsidered. [2] describes the 2D layered structure of the cortical networks in the brain.

4 Project Description

The project is built upon the hypothesized idea frame. First, the tools for the project are determined. For simulation implementation, Python programming language and a brain simulator package called Brian2 [3] are used. The simulations run on the HPC (High-performance computing cluster.), which is set to be used with the industry standard SGE

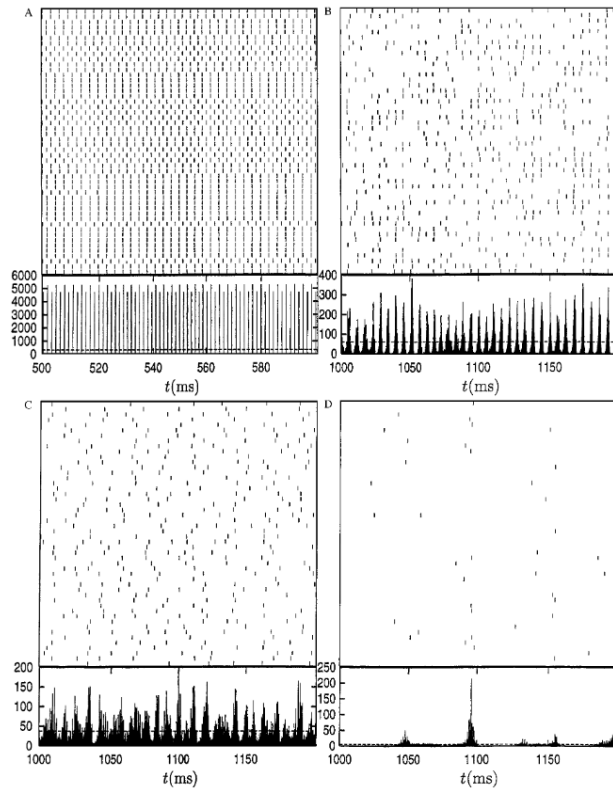


Figure 2: The states of the network. (A) Synchronous Irregular. (B) Synchronous Irregular-fast. (C) Asynchronous Irregular. (D) Synchronous Irregular-slow.

(Sun Grid Engine). The project timeline aligns with the report format presented in this document.

4.1 2D Connectivity

The 2D locality of the network is somewhat intuitive as the neurons in the cortical networks have a higher probability of connection to nearby neurons. In order to model this phenomenon, first, the N number of neurons is randomly located in the $1by1$ area. Then, according to the "smallest distance" between each pair, the probability map is constructed by zero-mean Gaussian. Figure 3 gives an example probability map for $N =$. In this formulation, the outdegree K of the neurons is fixed. That is to say, one neuron has a fixed number of outgoing connections. In other words, one axon has a fixed number of synapses. According to the probability map, the connection assignment process is conveyed as picking K number of neuron j 's to connect neuron i without replacement. A sample is given in Figure 4.

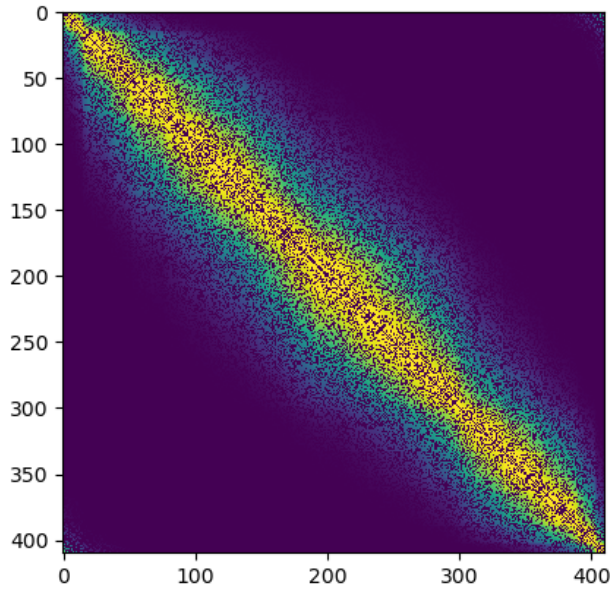


Figure 3: Sample probability map for $N = 512$

Also, the connections for one neuron are illustrated in Figure 5 as a part of the whole network for $N = 10000$ and $K = 100$, which is the standard network size through the rest of the study.

There are two important points that should be stated. First, the "smallest distance" is being calculated as if the 2D plane has periodic boundary conditions, creating 3D torus structure. Second, as the outdegree is fixed the σ value, which is the standard deviation of the Gaussian function has two limitations. One dependent on the system size, the other one is dependent on the outdegree. Since the periodic boundary conditions apply, larger σ loses

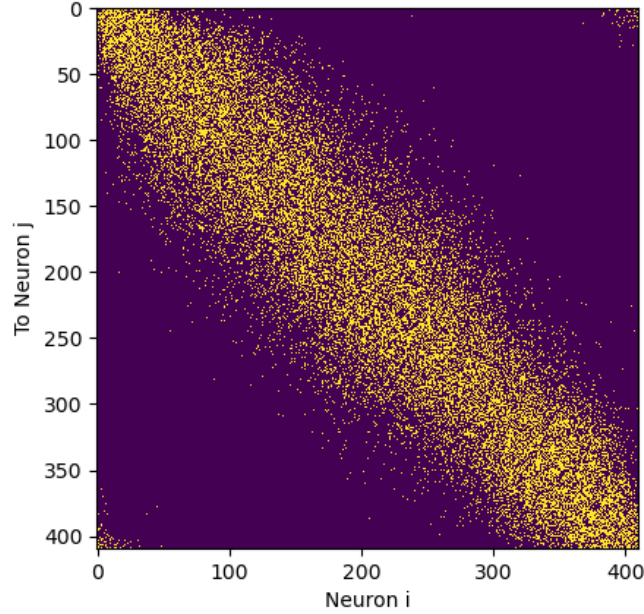


Figure 4: Sample connection map for $N = 512$

its effectiveness. On the other hand, as the σ goes smaller and smaller it becomes impossible distinguish between two different values because of the fixed number of selections, so one can not go more local in that sense. To be able to illustrate this situation the Figure 6. On x axis the σ value for the probability map is ranged. On y axis the *effective* σ which is the average distance of the connected neurons, is given. This plot allowed us to choose a small and useful enough σ value.

4.2 Baseline

First, the simulations that form a basis are conveyed before introducing regulative mechanisms. Let us build our model one by one. The differential equations that determine the behavior of a neuron are given in equation 1

$$\begin{aligned} \frac{dv}{dt} &= -\frac{(v - v_{leak_i})}{\tau_{mem_i}} + noiseparam \times sqrt(\frac{2}{\tau_{mem_i}}) \times xi \\ \frac{dI_{exc}}{dt} &= \frac{I_{exc}}{\tau_{exc_i}} \end{aligned} \quad (1)$$

The main parameters are taken from [1] for consistency. The parameters and the maximum noise value are given in Table 1. This is the model for spiking neurons as the v_{thres} exceeded the neuron spikes and the membrane voltage returns to the reset value. The d_{syn} is the axonal delay parameter where the spikes take their time to reach the destination neuron. The τ_{ref} corresponds to the refractory period in which the neuron can not spike for that

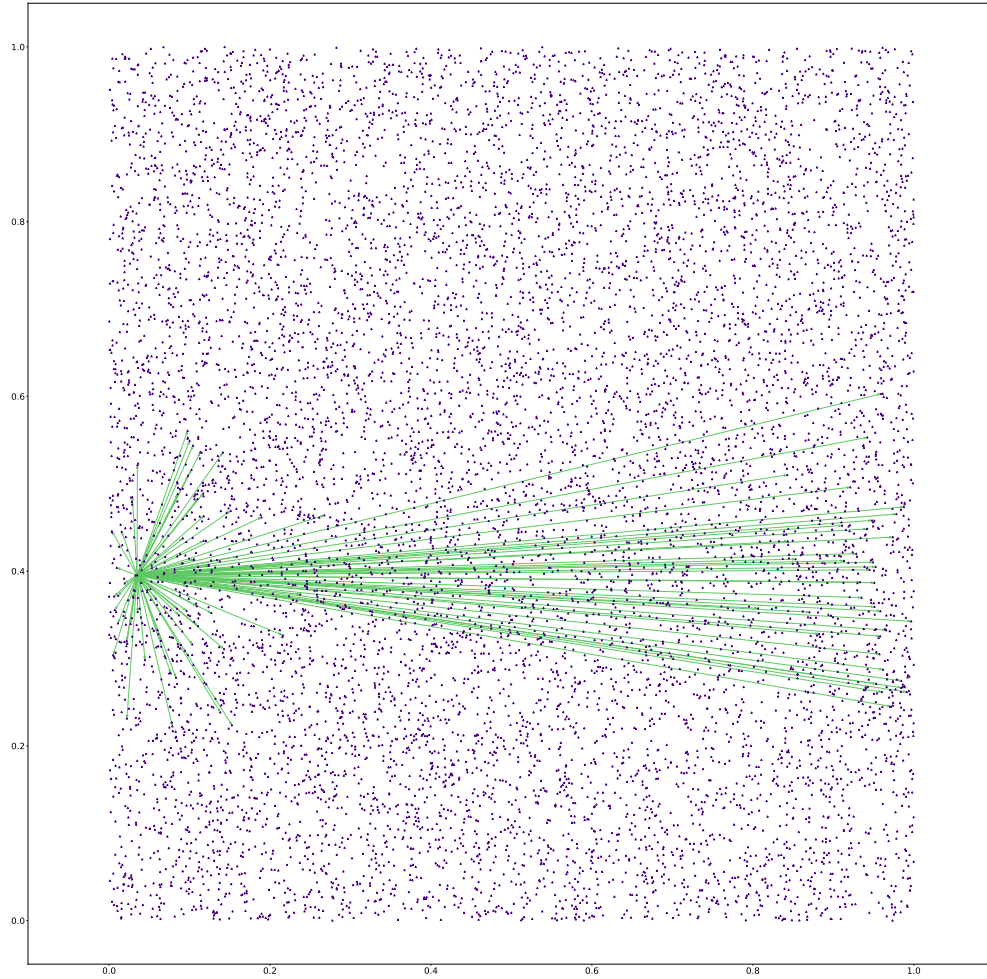


Figure 5: Connections of a neuron on the bigger network size.

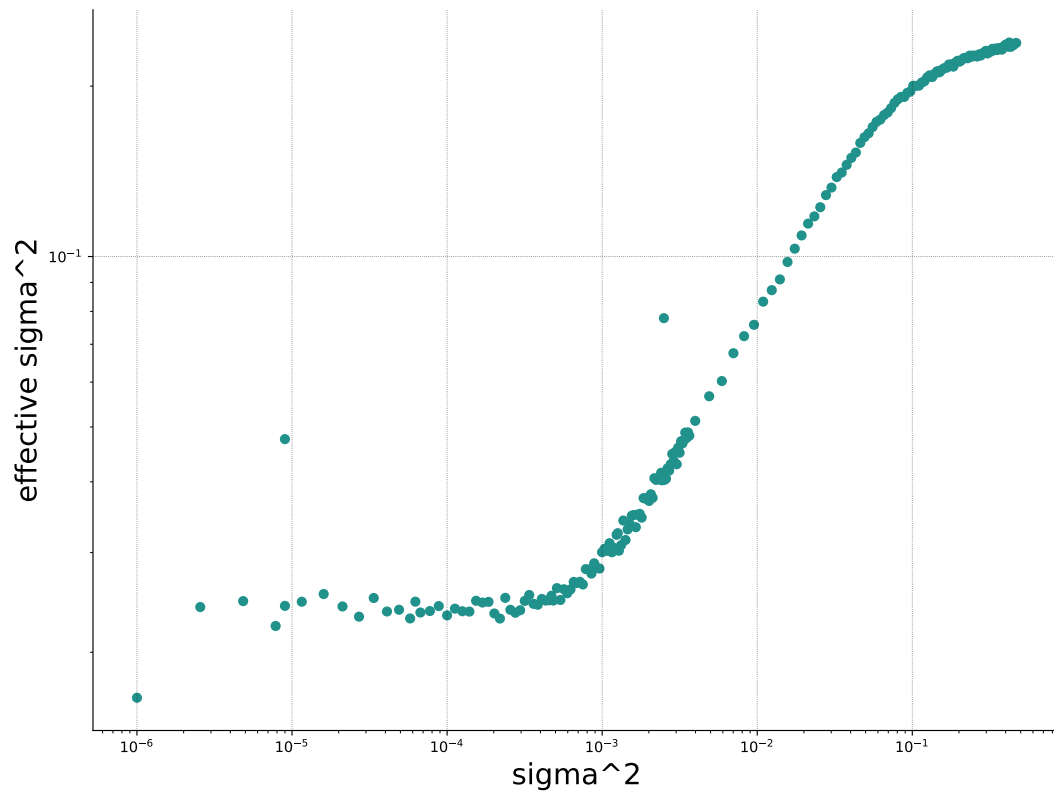


Figure 6: effective σ^2 vs σ^2 .

amount of time after spiking. The reversal parameter is implemented in order to scale the code easily in case of introduction inhibition. So, if a neuron gets too much inhibition in a time interval, the reversal parameter can not exceeded. That is biologically realistic since, in such a situation, the synaptic interface gets saturated.

Parameter	Value	Maximum Noise
τ_m	20.0 ms	1.0
$\tau_{\text{syn_exc}}$	5.0 ms	0.2
v_{leak}	0 mV	0
v_{thres}	20 mV	0.5
v_{reset}	10 mV	0.5
τ_{ref}	2 ms	—
d_{syn}	1.5 ms	—
v_{rev}	-10.0 mV	reversal

Table 1: Parameter Values

It should be noted that different from [1], temporal and parameter noise is involved in our model in order to make it stochastic, thus realistic. Also, to make sure that introducing %5 noise does not make the system deviate from where it should be, the simulations are also run in no-noise settings, and it is verified that the variability does not make the system go crazy.

As we have compiled our basic neuronal model, the coupling parameter J is swept through different values in order to find the threshold where the neuronal activity jumps. The activity is measured as a rate per second per neuron from a few milliseconds of the simulation. Therefore, the resultant rate gives necessary information about sustaining activity. In this baseline, the external drive can be summarised as follows. For 5 seconds, each neuron gets a very high external Poisson rate of excitation. So, the network gets charged up. Then, the external drive was suspended, and the simulation continued for 15 seconds. As a result, the $J(\text{mV})$ vs activity plot presented in Figure 7 is obtained. Also, when we zoom in on the time development of the four closest points to the jump point, as presented in 8, one can see that even though each network settles for a small amount of time to an intermediate non-full-bursting state, two of them dies out afterward.

The project is steered towards employing regulatory mechanisms for coupling parameter J by forming a baseline.

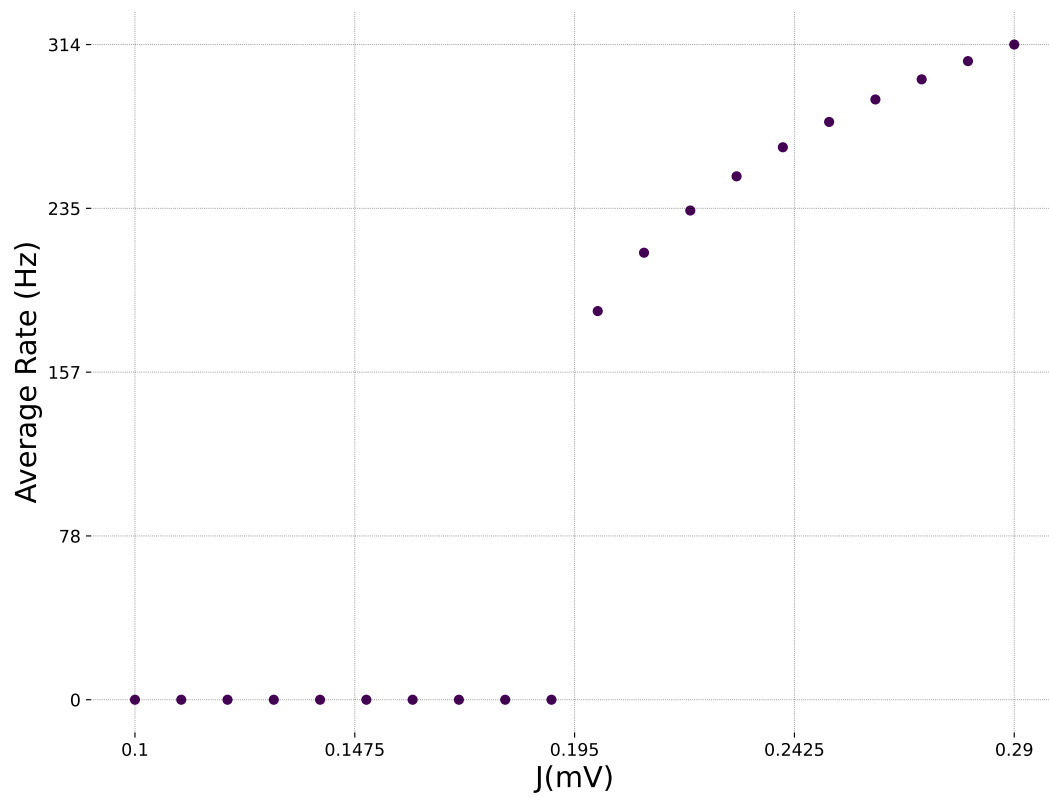


Figure 7: Rate (Hz) vs J(mV) for baseline.

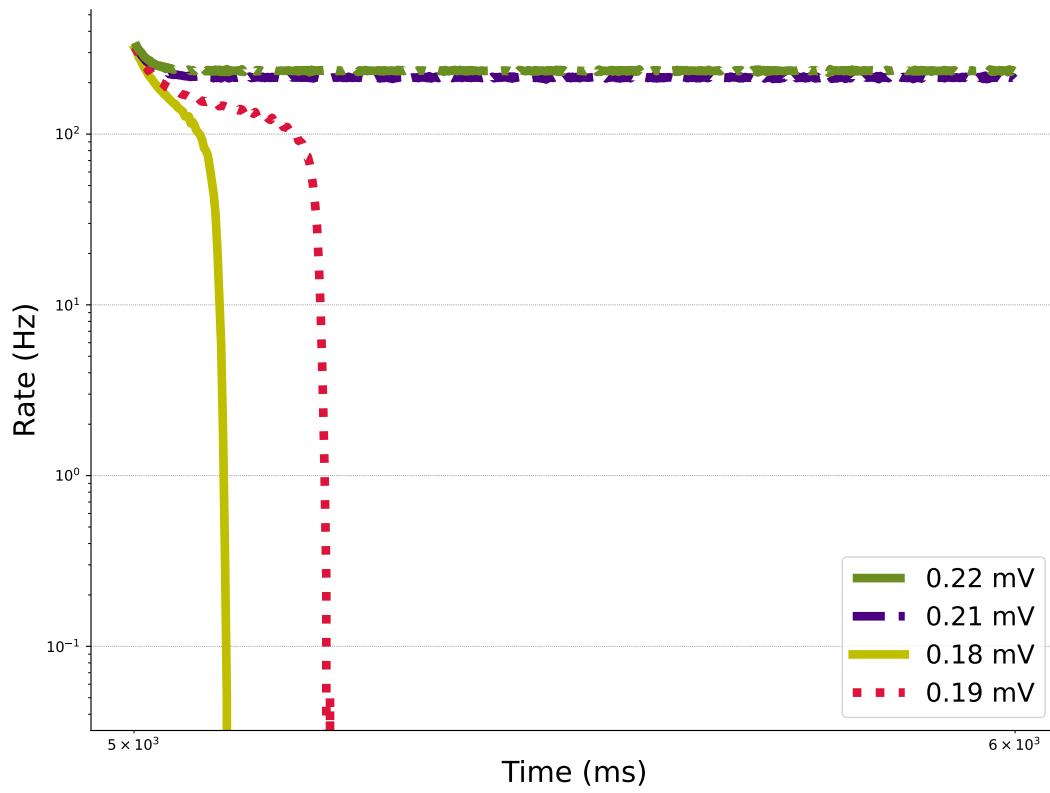


Figure 8: Rate (Hz) vs time (ms) for baseline samples in logscale.

4.3 Homeostatic Regulation

4.4 Future Projection on the Project

5 References

References

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6 Appendix