

Synaptic activity and excitability modulates information transfer in Purkinje cells: a modeling study Allan D. Coop¹, Hugo Cornelis¹, Fidel Santamaria²

Introduction

Synaptic information is transfered from synapses, to dendrites, to the soma. This is particularly true in Purkinje cells, since there are no backpropagating action potentials.





Mutual information between synaptic input: Spikes

Dendritic segments Dendritic currents Whole dendrite

Traditional approaches collapse the information capacity of a neuron into a point source process at the soma. Here we want to understand how the membrane excitability and synaptic activity affect how synaptic information is coded in a large dendritic tree.

Methods

Purkinje cell model

The simulations consisted in randomly activating all the excitatory and inhibitory synapses at constant Poisson firing rates. We used four different combinations of excitatory and inhibitory synaptic activity that resulted in the same firing rate at the soma of the Purkinje cell. We ran simulations for up to 400 s saving the value of all dendritic and synaptic currents every 100 µs. In order to avoid initial condition effects the first 5 seconds of all traces were not used for the analysis. Simulations were run with a pre-release version of the new GENESIS 3 software (http://www.genesis-sim.org/) in a cluster at UTSA (http://www.cbi.utsa.edu).

Statistical analysis

In order to simplify the analysis we monitored the total value of the synaptic or dendritic currents. We also chose to use the total excitatory current because then the results of our study could be mapped to dynamic current clamp experiments.

For the purposes of comparing the changes due to background activity we normalized the value of all currents from 1-100 and binned the data in 1000 equally spaced bins. All the analyses described here were performed with the normalized current values.

Initial characterization of currents was done by calculating the histograms under all the different combinations of synaptic activity. Further analysis consisted in calculating the cross-correlation between the excitatory synaptic input (IGIu) and dendritic currents.

Mutual information

The entropy was calculated as

$H(x) = \sum_{i} p(x_{i}) \log_2 P(x_{i})$

where p is the probability of seeing value x_i . The conditional entropy was calculated as

H(y|x)=∑_i∑_ip(x_i,y_j) log₂ p(y_j |x_i)

In our case, x is the input signal (IGIu) and y any of the dendritic currents. Conditional probability distributions matrices were calculated based on the binned traces. Finally, the mutual information was calculated:

$I(y \mid x) = H(y) - H(y \mid x)$

It is well known that the value of I can be biased due to the binning process and finite size of the data being analyzed (Panzeri et al., 2007). We used a recently developed toolbox in Matlab (Natick, MA) that allows the accurate calculation of the different information measurements and compensation for potential biases (Magri et al., 2009). The value of I can be biased if the joint probability distribution of the two traces being analyzed is scattered and does not fill out the joint probability space (1000 x 1000 entries). The ratio N/m has been shown to determine the strength of such a bias, where N is the number of non-zero entries in the joint probability distribution and m number of non-zero entries of probability distribution of the stimulus. If N/m is less than 1 then the value obtained from calculating the mutual information is biased. All our simulations had an N/m > 1.

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