

Assessing Retino-Geniculate-Cortical Connectivities in Alzheimer's Disease with a Neural Mass Model

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Abstract—Longitudinal studies have shown that increase of mean frequency within the theta band may be considered as an early symptom of progression into Alzheimer's Disease (AD). Also, slowing of mean frequency within the alpha band has long since been known to be a definitive marker in AD. This work is aimed at developing a better understanding of alterations in neuronal connectivity underlying Electroencephalogram (EEG) changes in AD. Specifically, connectivity changes in the dorso-lateral geniculate-cortical pathway are studied using a neural mass computational model. Connectivity parameters in the model are informed by the most recent experimental data on mammalian Lateral Geniculate Nucleus (dorsal). A slowing of the mean power spectra of the model output is observed with increase in both excitatory and inhibitory parameters in the intra-thalamic and thalamocortical pathways and a decrease of sensory pathway synaptic connectivity. The biological plausibility of the results suggest potential of further model extension in AD research.

I. INTRODUCTION

Currently specified diagnostic symptoms for Alzheimer's Disease (AD) correspond to a stage where irreversible damage has already occurred in specific neural pathways [1]–[3]. The biggest challenge in treatment of Alzheimer's Disease (AD) today is in finding a biomarker that can detect the disease well in advance and prior to such irreparable pathological alterations [4]–[7]. Recent advances in this direction have suggested retinal alterations as an early indicator of AD [2], [8]. This is in line with impaired visuo-cognitive skills reported in Alzheimer's Disease [9], [10]. An increase in theta rhythms and decrease in alpha rhythms (commonly referred to as 'slowing of mean EEG') are well known symptoms in EEG of patients with dementia of the Alzheimer type [11], [12]. It is now believed that similar thalamocortical mechanisms underpin the generation and behaviour of both alpha and theta rhythms [13], [14]. Furthermore, slowing of the mean EEG has long since been considered as an indication of deteriorating cognitive abilities [15]–[17]. In previous research [18], [19] we have used a classic neural mass computational model (Alpha Rhythms model (ARm)) [20], [21] to study the underlying neuronal synaptic aberrations as a cause of changed EEG in AD. Our results showed a slowing of the alpha rhythm with an increase of inhibitory connectivity parameter in the model, while the excitatory connectivity parameter was held at its initial values.

More recently, we presented a modified version of the ARm firstly by introducing a synaptic connectivity in the retino-geniculate pathway, and secondly by introducing synaptic

connectivity parameters based on experimental data from the dorsal Lateral Geniculate Nucleus (LGNd) [22]. This model was used for a similar study (as in [18], [19]) on both alpha and theta frequency bands. The model output displayed a definite slowing of the dominant frequency within the alpha and theta bands with increasing values of the connectivity parameters in the intra-thalamic loop and decreasing values of connectivity parameter in the retino-geniculate pathway. These results, and research reports of a probable link between the retinal pathway aberrations and AD [2], [8]–[10], inspired us to carry out the work presented in this paper. We propose to study the the modified version of the model (in [22]) with an emphasis on the retino-geniculate synaptic connectivity and its impact on the model behaviour in context to EEG behaviour in AD. Furthermore, the model output is studied by a simultaneous variation of connectivity parameters, unlike a single parameter variation approach in all our earlier works. The output of the model mimics cortical EEG [20] and corresponds to EEG changes associated with synaptic alterations in the retino-geniculate-cortical pathway. We speculate that such an approach might help to identify an EEG biomarker corresponding to visual modality related deficiencies observed in AD.

In Section II, we provide an overview of the model used in this work. The results are presented in Section III. Discussion of the results and future work is mentioned in Section IV.

II. A SIMPLE NEURAL MASS MODEL OF THE RETINO-GENICULO-CORTICAL PATHWAY

The model used in this work is shown in Figure 1 and is based on the classic Alpha Rhythm model (ARm) originally proposed by Lopes da Silva [20] and later modified [21]. It has two representative neuronal populations: Thalamocortical Relay (TCR) cells of the Lateral Geniculate Nucleus (LGN) and cells of the Thalamic Reticular Nucleus (TRN). The TCR cells receive excitatory afferents from the retinal cells and inhibitory feedback from the TRN cells. The TRN cells receive excitatory feed-forward afferents from the TCR cells. The resting state retinal afferents correspond to closed eyes of a subject and is simulated by a stochastic signal $\Psi_{\mu,\varphi}(t)$ with mean μ and standard deviation φ and expressed in spikes/sec [20], [21], [25], [26]. Excitatory and inhibitory synapses are represented

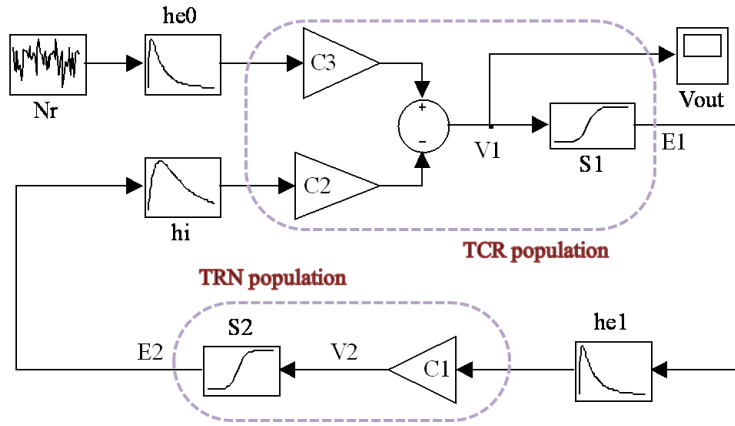


Fig. 1. The neural mass thalamocortical circuitry as a modification of the ARm and as simulated in our work with Simulink[®] in Matlab. The connectivity parameter values in the model are based on most recent experimental data (from the cat Lateral Geniculate Nucleus (LGN)) available from literature [23], [24] and represent the proportion of excitatory or inhibitory synapses from respective afferents on a single dendritic terminal of a cell. The block N_r represents retinal input to the model in an eyes-closed resting condition, simulated by a stochastic signal $\Psi_{\mu,\varphi}(t)$ with mean μ and standard deviation φ . All other blocks in the model are defined in (1)–(3). The parameter values for all blocks in the figure are provided in Table I.

by a response function defined as [25]–[28]:

$$h_{e,i}(t) = \frac{A_{e,i}}{\tau_{e,i}} t e^{-\frac{t}{\tau_{e,i}}}, \quad (1)$$

where $\{e, i\}$ represent excitatory or inhibitory synapses respectively, A is the synaptic amplitude and τ is the rise-time (time to reach 90% of the maximum value) of the kernel function $h_{e,i}(t)$. The ‘collective’ (commonly referred to as ‘lumped’) average membrane potential $V_{mp}(t)$ of a neuronal population is expressed as:

$$V_{mp}(t) = h_{e,i}(t) \otimes \xi(t), \quad (2)$$

where $h_{e,i}(t)$ is as defined in (1) and $\xi(t)$ is the pre-synaptic spike rate and expressed in spikes/sec. The membrane potential at the collective soma of a neuronal population is converted by a sigmoid operator $S(\cdot)$ to an average spike rate $E(t)$, representing the collective spiking frequency of the population and defined as [26], [28], [29]:

$$E(t) = S[V_{mp}(t)] = \frac{2e_0}{1 + e^{\nu(s_0 - V_{mp}(t))}}, \quad (3)$$

where e_0 is the maximum firing rate of the population, s_0 is the resting membrane potential and ν is the sigmoid steepness parameter.

We consider synaptic connectivity, unlike fibre connectivity considered in the original model [20], [21]. Furthermore, our choice of parameters is based on most recent available data obtained from cat LGN (dorsal) as reported in [23], [24]. The synaptic connectivity parameters in the model represent the proportion of a certain type of synapse on a neuronal population with respect to the total number of synapses converging on a single dendrite of the population from different sources. In Figure 1, C_1 represents the proportion of (excitatory) synapses made by the collaterals of the thalamocortical efferents from the TCR on the TRN population and is expressed as a percentage of the total number of synaptic vesicles at the synaptic junctions of a single dendrite of the population.

TABLE I
PARAMETERS USED IN THE MODEL AND DEFINED IN (1) AND (3). THE VALUES OF THE CONNECTIVITY PARAMETERS C_1 , C_2 AND C_3 ARE AS OBTAINED FROM MOST RECENT EXPERIMENTAL DATA FROM THE CAT LGN AND REPORTED IN [23], [24]. THE VALUES OF PARAMETERS IN (1) ARE AS USED IN [26], [28] WHILE THOSE USED IN (3) ARE AS IN [25], [27].

Parameters	Units	Values
μ	spikes per sec (sps)	550
φ	sps ²	120
A_e	mV	3.25
A_i	mV	22
τ_e	msec	10
τ_i	msec	20
ν	mV ⁻¹	0.56
e_0	V	25
s_0	mV	6
C_1	% of total (TRN)	35
C_2	% of total (TCR)	28
C_3	% of total (TCR)	10

Similarly, C_2 represents the inhibitory synapses by the TRN cells on the TCR population while C_3 represents the excitatory synapses by the retinal afferents on the TCR population. The experimental data corresponding to these synaptic connectivity parameters in the model are — $C_1 : 30 - 40$; $C_2 : 24 - 31$; $C_3 : 7 - 12$. We refer to these values as the ‘normal range’ of connectivity values indicating that these are obtained from a pathologically intact LGN(d).

The output of the model is the simulated membrane potential of the TCR population and mimics occipital EEG. All parameter values are defined in Table I. The connectivity parameter values in Table I are referred to in this work as their ‘basal’ values (synaptic connectivity values in a simulation of pathologically unaltered brain condition) and are evaluated as mean of the normal range of values obtained from experimental data and as mentioned above. Results are discussed in the Section III.

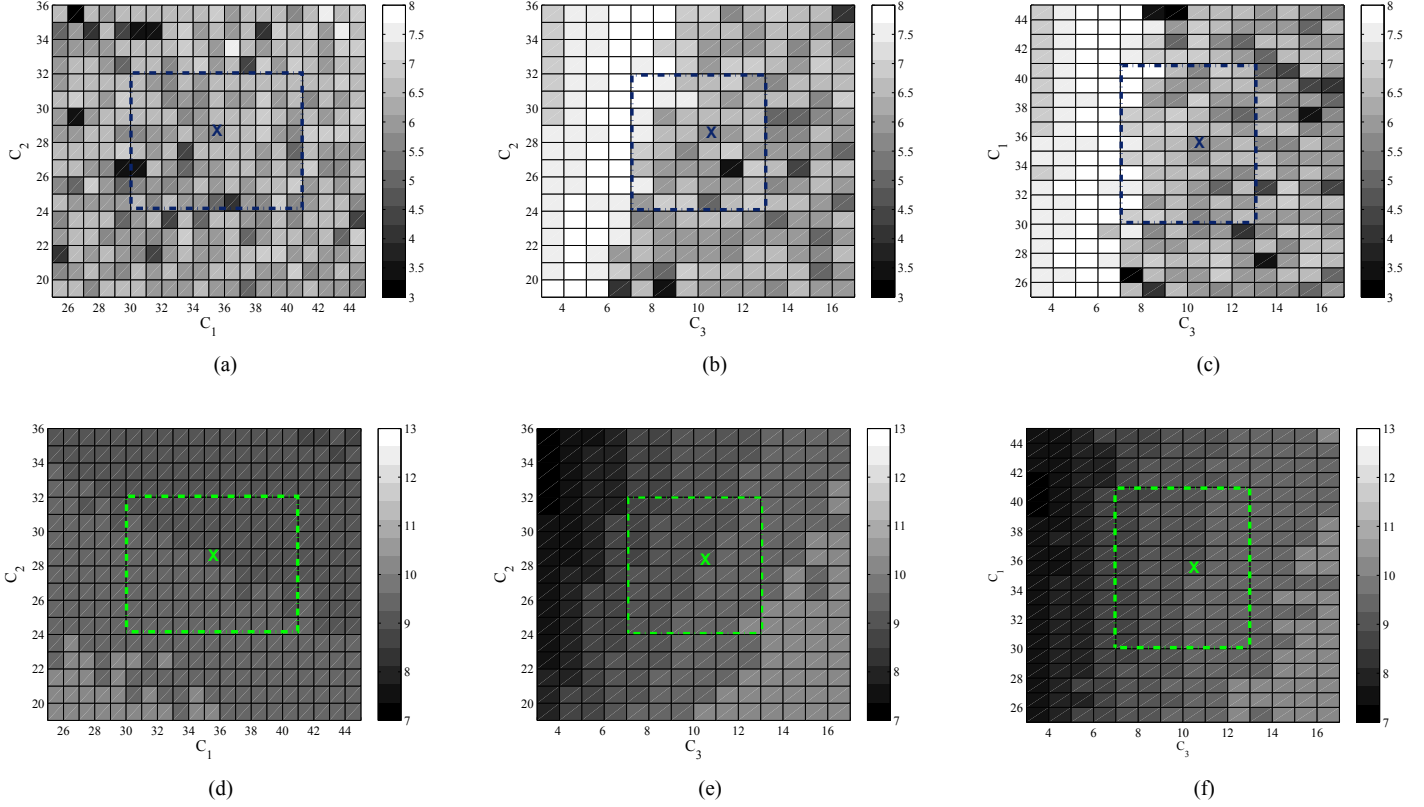


Fig. 2. Plots showing average dominant frequency behaviour of the model output within the (a)–(c) theta ($4 - 7 \text{ Hz}$) and (d)–(f) alpha ($7.5 - 13 \text{ Hz}$) bands. The parameter variation ranges are — $C_1 : 25 - 45$; $C_2 : 19 - 36$; $C_3 : 1 - 17$, while the normal range (range of values in a pathologically intact brain) for each of the parameters are — $C_1 : 30 - 40$; $C_2 : 24 - 31$; $C_3 : 7 - 12$. The basal value (a base value used to benchmark the model output oscillatory behaviour representing that of a pathologically unaltered brain) of each parameter is taken as the mean of the normal range of values for the respective parameter and is shown in Table I. The 'X' in each plot identifies the output response when all three parameters are at their basal values. The region within the square in each plot show the boundaries of the normal range of values of each parameter. (a)(c) Variation of the excitatory C_1 and inhibitory C_2 parameters in the thalamocortical pathway when the excitatory parameter C_3 in the retinal afferent pathway is at its basal value 10. (b)(e) Variation of C_3 and C_2 when C_1 is at its basal value 35. (c)(f) Variation of C_3 and C_1 when C_2 is at its basal value 24.

III. EXPERIMENTAL METHODS AND RESULTS

To simulate abnormal pathological conditions, the connectivity parameters in the model are varied above and below their basal values. The output power spectra of the simulated 'EEG' from the model is observed within theta ($4 - 7 \text{ Hz}$) and alpha ($7.5 - 13 \text{ Hz}$) frequency bands. Similar observations were made in our previous work by varying the connectivity parameters at arbitrary intervals about their basal values [18], [19]. In this work, we perform a global parameter search in order to study the dynamics of the output spectra at a higher resolution of parametric variations. Thus, we vary the connectivity parameters to every consecutive integer value within their respective normal ranges. Further, the values are varied up to ± 5 of their maximum and minimum values respectively. The parameters of the synaptic function as well as the sigmoid function are also different than those in our previous work and are based on physiologically plausible values as reported and used in [26], [28].

Recently, we have demonstrated that the choice of input mean μ and the standard deviation φ in this model as well as in ARm are crucial to the output oscillation frequency [19],

[22]. In this work, we adjusted these values by trial simulations so as to obtain an output dominant frequency of $10.5 - 11 \text{ Hz}$ within the alpha band and $4 - 6 \text{ Hz}$ within the theta band while the connectivity parameters are set at their basal values.

Model simulation is implemented using the $4^{th}/5^{th}$ order Runge-Kutta ODE solver within the Simulink[®] environment in Matlab. Total simulation time is 10 seconds with a sampling frequency of 250 Hz. The output from each simulation is clipped on the time axis so as to abstract the values from the start of the 4^{th} sec. to the end of the 8^{th} sec. as is done in experimental studies of EEG [21], [30]. The output vector thus obtained is bandpass filtered between $3 - 8 \text{ Hz}$ (to extract the theta rhythm components) and $6.5 - 14 \text{ Hz}$ (to extract the alpha rhythm components) using a Butterworth filter of order 10 in Matlab. The power spectra of this filtered output is computed in Matlab using a Welch periodogram with hamming window of segment length $1/4^{th}$ the size of the sampling frequency and overlap of 50% [18], [19], [31].

The results of variations of the connectivity parameters on the dominant frequency of the output spectra are shown in Figure 2. Each plot shows variation of model output dominant

frequency corresponding to variation of any two parameters; we show the cases when the third parameter is at its respective basal value. Compared to the other plots, there is a clear lack of systematic change in Figure 2(a) indicating that the connectivity parameters C_1 and C_2 have very little influence on the output behaviour within the theta band. In contrast, the parameter C_3 seem to play a major role on the theta band output behaviour. In both Figures 2(b) and 2(c), we see that there is a sudden increase in dominant frequency within the theta band for parameter values below the normal range minimum ($C_3 < 7$). On the other hand, the parameters C_1 and C_2 show a marked influence on the alpha band behaviour with a slowing within the alpha band for increasing values of C_2 observed in Figure 2(d). Furthermore, the average dominant frequency of oscillation within the alpha band is also reduced with increasing values of C_1 . These results agree with our earlier work with the ARM [18], [19], in spite of change in model parameters and structure. Figures 2(e) and 2(f) show a consistent slowing of the alpha rhythm with decreasing values of C_3 . The diagonal patterns in Figure 2(e) indicates an overall diminishing of the dominant frequency within the alpha band with increasing values of C_2 . However, the diagonal patterns are much steeper in Figure 2(f) showing a lesser impact of C_1 on the slowing of dominant frequency, thus agreeing with Figure 2(d). Overall, Figures 2(d)– 2(f) show that all the three parameters interact with one another within the alpha band.

A series of multi-way ANOVAs (done using the Statistical ToolboxTM in Matlab) show that the effects of variation of all three parameters on the average dominant frequency, shown in Figures 2(b)– 2(f), are statistically significant ($P \ll 0.001$). However, effects of variation of C_1 and C_2 on average dominant frequency within the theta band, shown in Figure 2(a), are not significant statistically ($P > 0.001$). The results are discussed in the following section.

IV. DISCUSSION AND FUTURE WORK

In this work, we present a modified version of the Alpha Rhythm model originally proposed by Lopes da Silva, subsequently extended [21] and validated in our previous work [18], [22]. We endeavoured to use the model in studying the underlying neuronal mechanism corresponding to changes in mean EEG observed in Alzheimer’s Disease patients. Here, we present a preliminary exhaustive global connectivity parameter search in order to study the model output behaviour within the theta and the alpha bands at a higher resolution than reported in our previous works.

Results presented in Section III show that there is an overall slowing of the mean dominant frequency in the output spectra when values of synaptic connectivity in the retinal pathway are below its normal range. This is indicated by the step increase in the dominant frequency within the theta band and a decrease within the alpha band. The output behaviour is thus in agreement with various research reports stating the vital role of several aspects of the visual modality as a biomarker in detection of Alzheimer’s Disease [8], [10], especially in its early stages [2], [9]. Furthermore, increase in theta band

power is reportedly a symptom of early stage Alzheimer’s Disease [11], [32]. Since the retina is the most accessible part of the nervous system [2], [33], we speculate that such simulated results indicating underlying loss in neuronal synaptic connectivity might suggest new directions of experimental investigation in Alzheimer’s Disease and possibly lead to new diagnostic and therapeutic tools.

The overall slowing of mean output frequency within the alpha band with increasing values of both the excitatory (C_1) and inhibitory (C_2) connectivity parameters in the thalamo-cortical loop is contrasting to the increase in alpha band average dominant frequency seen with increasing values of the excitatory parameter (C_3) in the retinal pathway. This agrees with results presented in our earlier work [22] where we have hypothesised that such behaviour indicate a vital role of the inhibitory cell population in the model. As the inhibitory cells represent characteristics of the TRN cell population, the results conform to experimental reports stating the important role of the TRN in modulating thalamic oscillatory behaviour [34]–[36].

Although parameters used in the model presented here are based on experimental data, there is ample scope for incorporating more biological plausibility — for example, the cortico-thalamic connectivities are now known to play a vital role in the thalamic as well as cortical oscillations [37]–[40]. Furthermore, experimental data suggests that the majority of synaptic inputs to the thalamic cells are from the cortex [23], [24], [41]. Alzheimer’s Disease is characterised by not only cortical, but also thalamic atrophy [42], [43]. In a recent study (under review for publication), we have shown that retinal connectivity with the thalamic inhibitory interneuron (IN) population in the model plays a more important role in slowing of alpha band power as compared to connectivity with the Thalamic Relay Cells (TCR). This is a significant result as firstly, an essential role of inhibitory brain circuitry in modulating brain rhythms is re-iterated; and secondly, this gives a direction to future research where the role played by IN cell population may be investigated more thoroughly than until now. We anticipate that such a research direction may provide vital clues on the retinal and thalamic role in modulation and alteration of cortical EEG in neuro-degenerative diseases such as AD.

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