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A mathematical framework simulating nerve fiber physiology



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ABSTRACT

Nerves are an important part in human body which not only controls the movement and locomotion of the body but also but also contains sensory receptors from other parts of the body which provide continuous feedback to the brain and spinal cord. Various diagnostic methods are used to detect the specific damage response of the nerve but they do not identify the precise location of the nerve damage. A computational nerve model may help to identify the exact location of nerve damage. Therefore, in this study the organization of a one-dimensional (1D) synthetic single element structural and functional model which typify the anatomy and physiology of the nerve is proposed. The geometrical model was developed using 1D linear Lagrange basis function while the functional model was developed by applying external stimulus and solving the bidomain model. The unmyelinated and myelinated nerve electrophysiological models were used to generate and propagate the action potential in this 1D synthetic single element model (SSEM). The nerve conduction velocity (NCV) was also computed in this proposed model and found that the myelinated nerve model has a higher NCV in contrast to unmyelinated model. This model will provide a platform for the development of the complete anatomical and functional model of the nerves in the various location of the body and may be helpful for clinician and physiologists in the evaluation and diagnosis of the structural as well as functional consequences of diabetic neuropathy in its initial stages.

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1. Introduction

Nerves serve a variety of complex and diverse tasks grouped into sensory, integrative and motor functions. Any impairments and progressive loss of and function, also known structure as neurodegeneration, can disrupt the sensations of touch, smell, taste, hear, sight, pain, heat and can hamper activities of speech, memory and body movements (Tortora and Derrickson, 2008). According to researchers, a large proportion of neurodegenerative diseases are associated with diabetes mellitus (McKusick, 2007; Ristow, 2004). Also, diabetic neuropathies affect up to 50% of diabetic patients all over the world (Vinik et al., 2006). Therefore, the diagnostic importance of neurodegenerative disorders has always prompted the research interests of clinicians and researchers worldwide.

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The electrophysiology of nerves is of real importance here. A nerve which is a collected term for several hundred axons, plus associated blood vessels and connective tissue can either be classified as myelinated nerve or an unmyelinated nerve, depending upon the presence or absence of a multilayer lipid and protein layer around the nerve axon called the myelin sheath. This sheath provides electrical insulation and speeds up the nerve conduction velocity through the axon (Tortora and Derrickson, 2008). Myelinated nerve axons have increased diameters of up to 22 µm resulting in enhanced conduction velocities of up to 130 mm ms⁻ ¹ as compared to unmyelinated nerves having a small diameter of up to 1.3 μm and slow conduction velocities of approximately 1.5 mm ms⁻¹ (Tortora and Derrickson, 2008). This translates that the axon diameter and degree of myelination along with temperature governs the speed of nerve impulse propagation through a nerve fibre (Giuliodori and DiCarlo, 2004).

Diagnostic techniques for diabetic neuropathies include both invasive and non-invasive methods but lack the ability to locate the exact position of damage (Oh, 2007; Kennedy and Inglis, 2002; Oh, 1990; Lauria et al., 2005). Our potential solution to this

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problem is application of mathematics to neuronal structure and dynamics in a computer based medium; neuronal computational modelling (Dayan and Abbott, 2003). Such mathematical and computational models offer valuable tools for investigating nerve activity across multiple structural locations of the human body independent of the need for an actual patient.

Most neuronal mathematical models are electrophysiological in nature and there are separate models to analyze the electrical activity of unmyelinated and myelinated nerve fibres (Rattay and Aberham, 1993). Two of the most popular electrophysiological are the unmyelinated Hodgkin-Huxley (HH) and myelinated Chiu, Ritchie, Rogert, Stagg and Sweeney (CRRSS) nerve models respectively. The HH nerve model was the first comprehensive mathematical model of electrical excitability in the nerve fibre (Hodgkin and Huxley, 1952). HH defined the membrane potential from non-linear ordinary differential equations (ODEs) of a single neuron. In contrast, the CRRSS is suitable for simulating the myelinated peripheral sensory nerve fibre in rabbits (Chiu et al., 1979; Sweeney et al., 1987). In this model, experimentation was performed on the myelinated nerves for the measurement of quantitative membrane currents using the voltage clamp analysis (Chiu et al., 1979; Sweeney et al., 1987). They used a set of ODEs to determine the membrane potential. Formerly, the electrical stimulation in the retinal ganglion cell and human cochlear neuron was done by employing HH model respectively (Greenberg et al., 1999; Rattay et al., 2001). Whereas, the electrical stimulation in the sciatic nerve of the lower limb was achieved by implementing the CRRSS model (Kim et al., 2007).

Therefore, this research work describes the framework for the development of structural and functional model representing the anatomical and physiological behaviour of the nerves in the human body. This modelling framework presents a practical standard in which biophysical nerve activity can be included with the structural and functional aspects of the nerves in various location of the body. This may ultimately prove to be a major diagnostic tool for examining the pathological consequences of diabetic neuropathy at its initial stages.

2. Methodology

In this work, a 1D synthetic single element model (SSEM) is proposed and developed. This SSEM represents the anatomy and physiology of the nerve fibres and their details are given below:

2.1. Geometrical nerve model

In this study, a 1D geometrical SSEM with a length of 20 mm, characterizing the structure of the nerve fibres was developed using a 1D linear Lagrange basis function in modelling software known as Continuum Mechanics, Image Analysis, Signal Processing and System Identification (CMISS).

The 1D linear Lagrange basis was selected due to the fact that nerve fibre act as a cylinder and model using 1D cable theory (Segev, 1998).

2.2. Nerve electrophysiological model

In this study, two nerve electrophysiological models namely unmyelinated HH and myelinated CRRSS models were used to simulate the action potential generation and propagation along the developed 1D SSEM respectively. These models were encoded using the CellML modelling benchmark (Hedley et al., 2001; Lloyd et al., 2004). Through these electrophysiological models, we were able to calculate the values of membrane potential, ionic currents, conductances, and their respective ordinary ODEs in CellML which resulted in the calculation of the action potential at a single grid point along the 1D SSEM. The details of these electrophysiological models are given as follows:

2.2.1. HH model

HH were the first physiologists who successfully experimented the behaviour of nerve excitation with squid giant axon using voltage clamp studies (Hodgkin and Huxley, 1952). The voltage gated channels of the cell membrane control the ion flow. The activation as well as inactivation gates of the channels permit the amount of influx and outflux of the ions responsible for the production of action potential. The equivalent electrical circuit of a HH model comprises of time dependent sodium and potassium channels, a time independent leakage channel, and a membrane capacitance, as shown in Fig. 1a.

• a: HH model equivalent electrical circuit representing the capacitor as the capacitance of the cell membrane, the voltage dependent sodium and potassium channels with arrows in their respective conductance symbol and voltage independent leakage channel with no arrow, the reverse potential of each channel i.e., v_{Na1} , v_{k1} and v_{L1} represented by the round shaped battery symbol. The arrows for the ionic currents are indicated from inside to outside (Hodgkin and Huxley, 1952).

2.2.2. CRRSS model

The popular CRRSS model describes the mammalian myelinated nerve response. The nodes of Ranvier are modelled by voltage gated sodium channels and the voltage gated potassium channel are absent in CRRSS model (Chiu et al., 1979; Sweeney et al., 1987). The single cell myelinated nerve fibre model is represented in Fig. 1b.

• b: The equivalent cable electric cell model represents nodes of Ranvier in a CRRSS model. This model shows voltage dependent sodium (arrow) channel and a voltage independent leakage channel with no arrows. The reversal potential of each channel is designated by v_{Na2} and v_{L2} , illustrated with a round shaped battery symbol. The arrows indicates the ionic currents

flowing from inside to outside of the membrane (Chiu et al., 1979).

The parameters and constants used in the HH and CRRSS models for determining the action potential along the SSEM are given in Table 1.



Fig. 1: Electrophysiological nerve electrical equivalent model

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Parameters	HH Model	CRRSS Model			
	Values and units	Values and units			
Membrane capacitance (v _m)	0.01 μF mm ⁻²	0.01 μF mm ⁻²			
Reverse potential (v _r)	-80 mV	-80 mV			
Sodium reverse potential (v _{Na})	115 mV	115.64 mV			
Potassium reverse potential (v _k)	-12 mV	N.A.			
Leakage reverse potential (v _L)	10.613 mV	-0.01 mV			
Sodium maximum conductance (v _{Namax})	1.2 mS mm ⁻²	14.45 mS mm ⁻²			
Potassium maximum conductance (v _{kmax})	0.36 mS mm ⁻²	N.A.			
Leakage maximum conductance (v _{Lmax})	0.03 mS mm ⁻²	1.28 mS mm ⁻²			
Nodal gap length (l)	N.A.	1.3 mm			
Diameter of the fibre (d)	N.A.	7 µm			
*N.A. denotes not applicable					

2.3. Functional nerve model

In order to translate the structural model into a functional model, a flow chart representing the various stages involved for action potential propagation (APP) along a SSEM is shown in Fig. 2. These stages are used for APP as well as NCV measurements along the 1D SSEM, characterizing the physiology of the nerve fibre. The geometrical 1D SSEM opted as an input for APP along the diverse parts of the elements. Grid points were allotted in the same vicinity as in the local coordinate space for this formed geometrical SSEM. The initial calculated values of the membrane potential, ionic currents and their respective ODEs obtained from the CellML based electrophysiological nerve models were passed to the equally space single grid points of this 1D SSEM. In finite element modeling (FEM), bidomain model (Tung, 1978) was selected to characterize the APP along the single grid point of the designed 1D SSEM by applying extracellular stimulus as it was formerly used to model neural tissue electrical activity (Kim et al., 2007). External stimulation was applied at time t = 0 and the membrane voltage value returned to the CellML based electrophysiological nerve model which computed updated values of membrane potential,

ionic currents and their particular ODEs and this parameters value were once more reassigned to the bidomain model for computing newer values of the action potential. The LSODA numerical integration technique was implemented in order to solve the differential equation implicitly of the electrophysiological nerve models for ionic current (Hindmarsh, 1983) and compute the action potential of each grid point in the element iteratively to a final desired time interval. The final APP output obtained from the bidomain model was then exported to the front end of CMISS namely continuum mechanics graphical user interface (CMGUI) and to UNEMAP. CMGUI is used for the 3D visualization of APP along the 1D nerve geometrical model and APP traces at different location (grid points) were displayed in UNEMAP.

3. Results and discussions

The 1D SSEM illustrating the structure of the nerve fibre is shown in Fig. 3. This 1D element has a length of 20 mm and divided into regular, evenly space grid points.

For functional model development, the aforementioned methodology was implemented in Fig. 3. There were 70 grid points in this 1D SSEM

that gave a spatial resolution of 0.29 mm. Here, an assumption was done that the diameter remained steady along the entire 1D SSEM. An extracellular stimulus current with a magnitude of 40000 μ A mm⁻¹ and duration of 0.5 ms was injected into the left end of the nerve fibre model using bidomain model (1D element) as shown in Fig. 4.

This magnitude of extracellular stimulus was strong enough to depolarize the nerve fibre and cause the APP along the representative 1D element nerve model that used both the unmyelinated HH and myelinated CRRSS models. Fig. 4 illustrates that the membrane voltage is at resting potential at 0 ms (*blue sphere*) with no extracellular stimulus. The membrane potential then starts propagating from 0 ms onwards. This subsequent APP is shown in Fig. 4. As a whole, Fig. 4 represents 1D SSEM setup using linear Lagrange basis function representing the physiology of the nerve fibre with regular, evenly spaced 0.29 mm grid point space.

This is represented at 0 ms, which is the resting state of the fibre. Action potential was propagated along a 1D SSEM from the leftmost grid point to the right end of the nerve fibres using HH model at various time steps i.e., 0.1 ms, 1 ms, and 3 ms.



Fig. 3: 1D single element model with regular space grid points

The NCV was used to characterize the action potential movement along the entire length of the SSEM of the nerve fibre and to determine the convergence of the bidomain 1D SSEM. It is calculated by (Eq. 1),

$$NCV = \frac{\sum D}{t_f - t_s} \tag{1}$$

where, D indicates the sum of the distances of individual grid points along the 1D SSEM of the nerve fibre, t_f represents the final time at which the maximum amplitude of the action potential reached

the other boundary end of the SSEM of the nerve fibre and t_s is the start time at which the extracellular stimulus current was applied to activate the bidomain nerve fibre.

The NCV at the various locations in the developed 1D SSEM was then determined using Eq. 1 using HH and CRRSS electrophysiological nerve models and is presented in Table 2. From Table 2, it can be easily determined that the NCV in the developed 1D SSEM was higher using myelinated CRRSS nerve model in contrast to unmyelinated HH nerve model over the same location i.e., distance of this 1D SSEM.



Fig. 4: 1D SSEM setup using linear Lagrange basis

Table 2: NCV measurements at different locations in the 1D SSEM using HH and CRRSS model

Distance (mm)	HH Model		CRRSS Model	
	Time (ms)	NCV (mm ms ⁻¹)	Time (ms)	NCV (mm ms ⁻¹)
4.3	1	4.3	0.2	21.5
8.6	1.9	4.5	0.36	23.9
12.9	2.8	4.6	0.5	25.8

Furthermore, a simulation research was executed to obtain numerical convergence in this 20 mm length 1D SSEM by varying the spatial resolution of the grid points. So, the grid points varied between 20 points ($\Delta x = 1mm$) to 100 points ($\Delta x = 0.2mm$). The NCV was then determined using Eq. 1 based on this range of spatial resolution by applying the simulations iteratively with a nerve fibre diameter of 5 µm and the intracellular and extracellular conductivities of 50 mS mm⁻¹ respectively. The effect of spatial resolutions on NCV from both the unmyelinated HH model and the myelinated CRRSS model is shown in Fig. 5. It can be concluded from Fig. 5 that the NCV declined as the number of spatial resolutions (grid point per mm) increased. It stopped declining as soon as the spatial resolution convergence of 0.29 mm was reached for both the HH and the CRRSS models respectively. In addition, the convergence of NCV achieved from the CRRSS model was higher than the NCV obtained from the HH model. This might be due to the certainty that it has a myelin sheath and also contains nodes of Ranvier (Giuliodori and DiCarlo, 2004). The converged NCV was 25.6 mm ms⁻¹ and 7.1 mm ms⁻¹ for the CRRSS and the HH models respectively.



Fig. 5: Convergence test for the spatial resolution using the bidomain model to examine the NCV in a 1D SSEM using the CRRSS and the HH model. With the increase in grid points per mm, the NCV was decreased. The minimum converged spatial resolution was ~3.5 grid point per mm or grid point spacing of ~0.29 mm

4. Conclusion

A framework for the development of structural and functional models, which embodies the anatomy

and physiology of the single nerve fibre, has been described. This framework presents a podium for the development of computational nerve models of the nerve networks in different regions of the human body. These models can be used to examine normal and abnormal neuronal function. This can eventually prove to be a useful diagnostic tool for investigating the pathological outcomes of diabetic neuropathy such as diabetic ulcers and other related conditions. In the prospective works, an electromechanical coupling between loading and sensory nerve feedback can be made in healthy as well as diabetic neuropathy patients.

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