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Microscopic Modelling of non-linear GJ channels

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 $\mathbf{1} gr$

On Gap Junctions

- What is a Gap Junction?
- Cluster of gap junction channels
- Linking structure between neighbouring cells • Provides direct passage of molecules and ions



Figure : Schematic diagrams of a standard connexin molecule and gap junction channel. (Del Corso et. al., 2006)

Non linear GJ model - 0D

• GJ current:

 $I_j = G_j(t, V_j)V_j,$

where:

- V_i transjunctional voltage.
- Conductance: $G_i(t, V_i) = g_{i,0}g_i(t, V_i)$.
- The amplitude of the junctional conductance:
- $g_{i,0} = 68nS$ for Cx43 cells, i.e. $g_{i,0} = 2nS$ for Cx43KO.
- Gating variable: $g_i = g_i(t, V_i)$

$$\frac{dg_j}{dt} = \frac{g_{\infty}(V_j) - g_j}{\tau_{\infty}(V_j)}$$

Assumptions

- Use only homotypic Cx43 and Cx45 GJ channels.
- GJ channels are at the perimeters of the cells, on the membrane.
- The rest of the membrane has only ionic channels. Use Beeler Reuter ionic model.

2D microscopic model

- What are they made of?
- Proteins connexins.
- 6 connexins = 1 connexon (hemichannel) • 2 connexons = 1 gap junction channel
- Cardiac cell proteins: Cx43, Cx45 and Cx40.



Figure : Predicted configurations of connexons and gap junction channels for two different connexins. (Desplantez, 2004)

- Where are they located in cardiac cells?
- Mostly on the longitudinal ends where they compose the intercalated disks
- The behaviour of transversal GJ channels is not well understood



• Fit experimental data to find normalised $g_{\infty}(V_i)$.



• Fit experimental data to find $\tau_{\infty}(V_i)$.





Main problem:		
$\sigma^i \Delta u_i = 0,$	in Ω_i ,	
$\sigma^e \Delta u_e = 0,$	in Ω_e .	
Ionic model:		
$\partial_t V_m + I_{ion}(V_m, \mathbf{h}) = -\sigma$	$\sigma_i \nabla u_i \cdot \mathbf{n},$	
$\partial_t V_m + I_{ion}(V_m, \mathbf{h}) = -\sigma$	$[\nabla_e \nabla u_e \cdot \mathbf{n}, \} \text{on } \Gamma_{io}$	n
$\partial_t \mathbf{h} = f_{ic}$	$_{on}(V_j,\mathbf{h}), \]$	
GJ model:		
$G_j(V_j,g_j)\cdot V_j=-\sigma_i abla$	$^{7}u_{i}\cdot\mathbf{n},$ on Γ .	
$\partial_t g_j = f_j (V_r)$	$(m, g_j), \int 0111 j$	
Boundary conditions and stin	mulus:	
$\sigma_e \nabla u_e \cdot \mathbf{n} = 0,$	on $\partial \Omega_e \setminus \Gamma_{ion}$	1,

Figure : Immunohistochemical analysis of Cx43. Left: Bar = $10\mu m$. (Beauchamp, 2004) Right: top view(A), lateral view(B). Bar = $5\mu m$. (Beauchamp, 2012)

- What about electrical behaviour?
- The dual voltage patch clamp
- Non-linear behaviour
- Time dependence
- Dependent on connexin arrangement.





Figure : GJ currents in time, fixed V_i . Left: Homotypic Cx43/Cx43 cell pairs. Right: Cx43KO/Cx43KO cell pairs. for the fixed V_i .

on Γ_{qr} , $u_e = 0,$ for $t \in [t_0, t_0 + t_{stim}]$, $u_e = U_{app},$ on Γ_{app} .

- σ_i, σ_e intra and extracellular conductivities • **h** gating variables for ionic model • $V_m = u_i - u_e$ transmembrane potential
- $V_i = [u_i]$ transjunctional potential

Numerical analysis - 2D test case

• Domain: 2×1 cells, cell size $100 \times 20 \mu m$, distance $1 \mu m$. Mesh and time step: $dx = 1 \mu m$, dt = 0.02 m s. - BR ionic model. Semi-implicit time scheme. FEM. Iterative method. Test: no gap junctions.



Macroscopic effects

- Primary cultures of Cx43 and Cx43KO ventricular myocytes.
- Macroscopic velocity calculated from the difference in activation times





Figure : Left: A mixed patterned cell culture at low magnification. Right: Decrease in velocity of propagation w.r.t. presence of Cx43 cells.



On going work

- Current observations: without GJ there is no AP propagation from cell to cell.
- To be tested: finer mesh, smaller cell distances.
- Include different types of GJCs on larger domains.
- Compare velocities with the experimental observations.

Future work:

• Move to homogenised model.

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