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# The Roles of K+ Currents in Human Articular Chondrocyte Electrophysiology:

# A Computational Perspective

**Harish Narayanan**[1](#foot78)   
*Center for Biomedical Computing,*   
Simula Research Laboratory, Lysaker, Norway

**Mary M. Maleckar,**   
*Center for Biomedical Computing,*   
Simula Research Laboratory, Lysaker, Norway

**Robert B. Clark,**   
*Faculty of Kinesiology,*   
University of Calgary, Calgary, Canada

**James R. Wilson,**

Innovate Calgary

Calgary, Canada

**Wayne R. Giles,**   
*Faculty of Kinesiology,*   
University of Calgary, Calgary, Canada

### Abstract:

We have developed a computational model for studying the electrophysiology of the human chondrocyte. This model is based on an initial experimental data set, which has identified the main K+ currents that are expressed in these isolated single cells form knee joints of healthy adult human donors. The applicability and utility of this model are illustrated by focusing on the role of a novel 2-pore K+ current in regulating the chondrocyte resting potential and modulating intracellular Ca2+ homeostasis. This model is demonstrated to be useful for integrating available data from electrophysiological, PCR and gene array experiments. It is also an important adjunct for rationalization of working hypotheses, design of new experiments and understanding the principles and limitations of patch clamp methods applied to the isolated human chondrocyte.

*Key words:* chondrocyte; electrophysiology; potassium currents; computational model

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**INTRODUCTION**

Articular cartilage is considered to be an aneural, avascular, alymphatic component of the flexible connective tissue that covers the articulating ends of diarthroidal joints (1-3). It is essential for the required stability and low friction movement of the associated long bones. This connective tissue consists of an extracellular matrix (ECM, composed primarily of collagen, elastin and proteoglycans, (as detailed below) and only one type of cell - the chondrocyte. Chondrocytes are responsible for synthesis and homeostasis of the extracellular matrix. Articular cartilage is exposed to cyclical mechanical forces. In fact, this dynamic environment is essential for the health of the tissue (2) Chondrocytes occupy only approximately 1-10% of the total volume of articular cartilage in mammals ([4](chondrocyte-model.html#CarneyMuir1988),[5](chondrocyte-model.html#Halletal1996)) and play no direct mechanical role. Instead, mechanical support is provided by the ECM, which is composed of (a) collagen fibers that give the tissue the ability to resist tension, (b) negatively-charged gel-like proteoglycans (PGs) that are trapped within the collagen mesh and allow the tissue to bear compression (1,8) and (c) synovial fluid within the articular capsule which acts as a lubricant, thus ensuring low friction movement of the bones (7). The primary role of the human chondrocyte is to maintain viable cartilage by balancing macromolecular synthesis and breakdown (see e.g. 2,7,9).

In a variety of conditions (including healthy aging), or as a consequence of injury, progressive chondrocyte damage and dysfunction may occur (10-12). As a result, the balance between matrix synthesis and degradation is altered and the low friction environment of the joint may be reduced or lost (9). Under these conditions there is often an inflammatory response within the articular joint. This is one factor that increases the early development of osteoarthritis: a thinning of the cartilage layer which causes painful, bone-against-bone friction (1,11). The progression of osteoarthritis (12,13) and the ability of chondrocytes to respond to perturbations in the extracellular environment (14) are associated with deficiencies in chondrocyte volume regulation (15). It is known that physical damage to cartilage is more frequent in the setting of altered osmolarity (16). In turn, there is evidence that these volume changes are linked to an abnormal resting potential in chondrocytes (17), which may be due to changes in K+ and/or Cl- channel activity. Direct experimental investigations that address possible functional relationships between chondrocyte electrophysiology and pathophysiology are challenging due to the small cell size of the mature chondrocyte and the associated limitations of *in vitro* electrophysiological studies. In fact it is not certain that conventional patch pipette methods (18,19) can accurately determine the resting potential of isolated single chondrocytes (20). Accordingly, for the purpose of integrating available data and attempting to understand its functional significance, we have developed a detailed biophysical model of human chondrocyte electrophysiology. This model can facilitate investigation of electrophysiological mechanisms, which regulate chondrocyte physiology and pathophysiology and are the targets of intra- and intercellular signaling pathways responsible for biomechanical roles including excitation-secretion coupling.

Early signs of osteoarthritis may develop even in young individuals following articular joint injury. This is thought to be due to a phenomenon denoted 'chondrolysis', a condition in which accelerated loss of articular cartilage occurs over a short time period (21-23). Several clinical studies have also documented that this significant pathophysiological response can occur as a result of postoperative administration of bupivacaine, a local anesthetic. Experimental work has confirmed that bupivacaine can cause profound effects including chondrocyte death in both isolated cell experiments (24) and associated animal studies (25-28). However, the cellular and subcellular mechanisms responsible for this toxic response (characterized by increased apoptosis) are not completely understood. In this regard it is interesting to note that a family of two-pore K+ channels that have been recently identified in human articular chondrocytes can be inhibited by bupivacaine (29). Since bupivacaine-induced block of these and other (29) 2-pore K+ channels can result in a pronounced depolarization of the chondrocyte it is important to gain a detailed understanding of the functional relationships between membrane potential, intracellular Ca2+ levels (that can regulate apoptosis) and cell volume which can modulate excitation-secretion coupling.

The main goals of this work are (i) to develop the first detailed mathematical model of chondrocyte electrophysiology and, (ii) to use this model to investigate the relationships between membrane potential and essential cellular functions including excitation-secretion coupling and stretch sensitivity.

**MODEL DEVELOPMENT AND METHODS**

Our model has been constructed on the basis that the 'target system' consists of a single chondrocyte as an isolated cell, which resides in deep regions of articular joint cartilage. This specialized extracellular environment was modeled by specifying external concentrations [Na+]o , [K+]o , [Ca2+]o, [H+]o and [Cl-]o within physiologically-relevant, slightly hyperosmostic ranges (see Table [1](#tab:external-concentrations-2)). The chondrocyte cell membrane is known to express a number of voltage- and ligand-gated ion channels as well as pumps and exchangers (30).

Patch clamp experiments done on enzymatically isolated individual human chondrocytes obtained with permission from a knee replacement surgery program (The Southern Alberta Transplant Service) provide the basis for this model development. These chondrocytes have resting membrane potentials that range from -30 to -60 mV when superfused with normal Tyrodes solution and studied using antibiotic-permeablized (amphotericin) patch clamp methods (28). This significant range of resting membrane potential values may reflect the heterogeneous physiological state of these cells. However, as we have illustrated previously, some of this variability is likely to result from the fact that in these very small, approximately spherical cells (diameter, 7 microns; capacitance 10-15 pF) the patch pipette recording method is being applied very near its technical limitations (20). That is, the input resistance of the target cell (chondrocyte) is very large (1-10 gigohms), and the seal resistance between the surface membrane of the chondrocytes and the polished surface of the glass pipette has a typical value of 1-5 gigohms. Thus, the actual chondrocyte membrane potential is underestimated due to the current flow through the seal resistance that in most circumstances results in a depolarization. The Appendix of this manuscript describes the known biophysical principles which, when combined, regulate the ways in which the combination of patch pipette seal resistance and intrinsic input resistance result in apparent values for the resting potential in small cells such as the human chondrocyte.

The main objective of this first order model is to obtain insights into the ionic mechanisms that underlie the resting potential in the human chondrocyte *in situ*. The raw data that forms the basis of this model is derived mainly from our previously published recordings of the K+ currents in human (28), mouse (31) and canine chondrocytes (32). Other elements summarized diagrammatically in Figure 1 below are obtained from published literature on isolated chondrocytes from rabbit, canine and equine tissue.

[Figure 1 near here]

## The Atypical Environment of the Chondrocyte

As noted in the Introduction, the chondrocyte is situated in a physiological environment that differs significantly from that of most other mammalian cells. This information is summarized in Table 1. The large electrochemical gradients for Na+, K+ and Cl- are very similar to those in other mammalian systems. These gradients lead to the requirement for an ATP-dependent Na+/K+ pump mechanism that is assumed to be electrogenic. Note however, that the extracellular fluid within the articular joint is hypertonic (approximately 320 mOsm vs. blood plasma which is approximately 280 mOsm). In addition the extracellular pH of the synovial fluid that bathes the chondrocyte is somewhat acidic, pH 7.2, (8,24) although apparently the normal intra- to extracellular pH gradient can be identified.

[Table 1 near here]

The substantial number of fixed negative charges on immediately adjacent proteoglycans attract free cations (e.g. Na+) and exclude free anions from the matrix. As a result of this cation accumulation there is an osmotically driven water influx. (10,17).

The articular joint receives little or no vascular supply. As a result the synovial fluid must supply adult articular cartilage with the required small amounts of nutrients as well as oxygen to e.g., maintain Na+/K+ pump activity (33). Metabolic byproducts are removed mainly by diffusion (34,35) A second consequence of the avascular nature of this tissue, is that the chondrocytes generate ATP by substrate-level phosphorylation during anaerobic respiration. This generates H+ ions as a byproduct, which further lowers the pH in this micro environment. The dynamic changes in mechanical loading within the knee joint during activity also exposes chondrocytes to very significant fluctuations in vector forces and shear. The resulting mechanical changes would be expected to activate mechano- or shear-sensitive ion channels (36-40).

Experimentally reported values for the external cation concentrations differ markedly from those of normal extracellular fluid. We have incorporated these values for extracellular ion concentrations.

**Table 1:** Consensus Values of Ion Concentrations for Chondrocyte Electrophysiology ([8](chondrocyte-model.html#Wilkinsetal2000)).

|  |  |  |  |  |  |  |  |  |  |  |  |  |  |  |  |  |  |  |  |  |  |  |  |  |  |  |  |  |  |  |  |  |  |  |  |  |  |  |  |  |  |  |  |  |  |
| --- | --- | --- | --- | --- | --- | --- | --- | --- | --- | --- | --- | --- | --- | --- | --- | --- | --- | --- | --- | --- | --- | --- | --- | --- | --- | --- | --- | --- | --- | --- | --- | --- | --- | --- | --- | --- | --- | --- | --- | --- | --- | --- | --- | --- | --- |
| |  |  |  |  |  | | --- | --- | --- | --- | --- | |  | Cytoplasm | Matrix | Serum/Synovium |  | | (mM) | 40 | 240-350 | 140 |  | | (mM) | 120-140 | 7-12 | 5 |  | | (mM) | 8.e-5 | 6-15 | 1.5 |  | |  | 60-90 | 60-100 | 140 |  | |  | 20 | 15 | 23 |  | |  | 0.17 | 0.30 | 0.81 |  | | pH (mM) | 7.1 | 6.6-6.9 | 7.4 |  | | Osmolarity (mOsm) | -- | 350-450 | 300 |  | |

**MATHEMATICAL FORMULATIONS FOR IONIC PUMP AND EXCHANGER CURRENTS**

### Background Leakage Currents

The model accounts for background leakage of Na+ and K+ through the use of time-independent channels whose mathematical expressions are based on standard Hodgkin-Huxley formalisms:

|  |  |
| --- | --- |
|  | (1)  (2) |

The Nernst potentials for Na+ and K+ are computed based on their respective intra- and extracellular concentrations:

|  |  |
| --- | --- |
|  | (3)  (4) |

As indicated in the section below there is evidence that adult mammalian chondrocytes from a number of species express time independent and/or ligand gated Cl- channels. We have incorporated this type of current into our model by treating it as a background current, specified by the equations below.

|  |  |
| --- | --- |
|  | (5) |

where

|  |  |
| --- | --- |
|  | (6) |

is the Nernst potential due to the difference in Cl- concentration inside and outside the cell.

### Ion Pumps and Exchangers

**Na+/K+ Pump**

The active extrusion of Na+ ions from the chondrocyte is assumed to be achieved by the turn over of an electrogenic Na+/K+-ATPase. Mobasheri et al (33) have characterized some of the functional properties of an electrogenic Na+/K+ pump in bovine articular chondrocytes. Cell volume is maintained by altered balance of leaks and pumps to hold cell water constant. In this model, we employ the following sodium-potassium pump formulation (42).

|  |  |
| --- | --- |
|  | (7) |

Figure X shows a representative current-voltage curve for this electrogenic pump.

**Na+ /Ca2+ Exchanger**

As in many other cell types, the Na+/Ca2+ exchanger plays a key role in Ca2+ homeostasis in articular chondrocytes (41). We model this electrogenic exchange process using the following mathematical expression (42).

|  |  |
| --- | --- |
|  | (8) |

**Na+ /H+ Exchanger**

Chondrocytes express a Na+/H+ antiporter (43,44) which allows the cell to regulate pH and establish an intra- to extracellular pH gradient. This pH gradient is essential for maintaining intracellular Na+ levels and the intracellular pH indirectly regulates a number of essential enzymatic processes. In order to model this channel, we use the following functional form described (45).

|  |  |
| --- | --- |
|  | (9) |

where,

|  |  |
| --- | --- |
|  |  |

### Potassium Currents

This model is based on mathematical expressions for a total of 4 distinct K+ currents that have been identified from electrophysiological studies done on mammalian chondrocytes. Two of these, a delayed rectifier K+ current which we denote IK-DR; and a K+ current due to a 2-pore K+ channel which we denote IK-2P have been studied in detail in our laboratory (28,32), and by others (45-57). The remaining two K+ currents, a Ca2+-activated K+ current (IK-Ca) and an ATP-dependent K+ current (IK-ATP) have been identified in other patch clamp studies. These are also included.

a) Delayed Rectifier K+ Current: IK-DR

A time- and voltage-sensitive K+ current has been identified in the human articular chondrocyte (28-32). The biophysical properties of this current and the details of its pharmacological blockade suggest that it is generated by the family of alpha subunit K+ conductances denoted Kv1.4 or Kv1.6. Panel A of Figure 3 shows an I-V relationship generated under physiological conditions, that is, a normal electrochemical gradient for K+. Panel B of this Figure shows the steady-state activation curve and Panel C illustrates the kinetics of activation and deactivation within this range of membrane potentials. The fits to this data provide the basis for the simulation of this current in our model.

In this work, the mathematical expression for the delayed rectifier is based on descriptions for the ultra-rapidly rectifying potassium current in human atrium (48).

|  |  |
| --- | --- |
|  | (10) |

In the above equation aur and iur are time-dependent channel activation and inactivation, and are defined in the model via the following expression:

|  |  |
| --- | --- |
|  |  |

Figure [2](#fig:potassium-currents)c shows the current voltage curve for this current. The continuous I-V curve has been fitted to our experimental data (32).

[Figure 3 near here]

b) 2-Pore K+ Current:

Our recent work has identified recording conditions under which an additional K+ current generated by what is believed to be the TASK family of 2-pore K+ channels (28). These channels show no detectable time dependence. An additional characteristic of this subclass of 2-pore channels is their significant increase in conductance in response to an increase in pH or alkalinization of the extracellular medium. A peak I-V curve for this current is shown in Panel A of Figure 4. Figure 4B illustrates the enhancement of this current when extracellular pH was changed from XXXXX.

Note that these recordings were made under conditions of elevated K+ so that the size of the currents could be increased to the level that they could be detected and biophysical properties could be resolved. Before this data can be incorporated into this mathematical model it needs to be corrected appropriately. The correction that has been employed is based on the Eisenman principle (54) that the conductance of an ion selective channel scales according to the square root of the concentration of that permeant ion. The I-V curve in Figure 4C shows the original data recorded in isotonic K+ with the expected reversal potential of 0 mV together with the corrected data assuming external K+ to be 5.4 mM with a corresponding reversal potential of approximately -85 mV.

[Figure 4 near here]

Using the classic Goldman-Hodgkin-Katz equation (54) for voltage-gated, time-independent single-species ion channels, this current can be represented by:

|  |  |
| --- | --- |
|  | (11) |
|  |  |

and Figure 5a shows the current-voltage curve fitted to experimental data (28). Our experimental work also demonstrated that this particular K+ current was strongly inhibited by bupivacaine. An effective concentration of bupivacaine also resulted in a significant depolarization of the resting potential (see Discussion).

[Figure 5 near here]

c) Ca2+ Activated K+ Current

Virtually all recordings of global K+ currents in human chondrocytes include a significant and very noisy outward current at depolarized potentials (55,56). The spontaneous fluctuations of this current suggest that it is generated by the so-called large conductance variant of the Ca2+ activated K+ current family (57). We have not explored the biophysical properties of this current or its pharmacological profile in any detail. However, it is consistently present in our recordings and a Ca2+ activated K+ current has also been described in isolated chondrocytes from rabbits, dogs and horses (47). Figure 6 below shows represented data. The formulation for the Ca2+ activated K+ current in this model is given by the set of equations below.

[Figure 6 near here]

In the present formulation, we model the (large) Ca2+-activated K+ channel using a functional form defined in previous papers (57,58).

|  |  |
| --- | --- |
|  | (12) |

where,

|  |  |
| --- | --- |
|  |  |

Figure 7b shows the current-voltage relationship for this current fitted to our experimental data.

[Figure 7 near here]

ii) Intracellular Ca2+ Homeostasis

The intracellular Ca2+, [Ca2+]i is maintained at very low levels by incorporating an electroneutral, ATP requiring Ca2+ pump. In accordance with our previous models of the atrial myocyte (42,48) the intracellular Ca2+ buffering in this model is assumed to be due mainly to calmodulin within the cytosol. The expressions which govern this Ca2+ buffering are given below:

(12)

MOLLY: PLEASE ADD CALCIUM BUFFERING EQUASIONS

d) ATP-sensitive K+ Current

An ATP-sensitive K+ current has been reported in chondrocytes from a variety of mammalian species (59). None of our experimental work has addressed this possibility and there are no reports of this current being activated in human chondrocytes. However, the hypoxic environment in which the chondrocyte is placed makes it plausible that this current could be turned on during normal biomechanical activity.

[Figure 8 near here]

e) Cl- Current

In a variety of mammalian chondrocyte preparations one or more types of Cl- conductance has been identified (59,c.f. 30). This quasi linear Cl- conductance has also been identified in human chondrocyte cell lines (61), and in experiments in which selected transmitters or pharmacological agonists (e.g., histamine) have been applied (62).

f) Transient Receptor Potential or TRP Current

Ligand gated channels that exhibit properties that correspond to some of those of TRP ion channels are expressed in mammalian chondrocytes (63). The TRPV4 family is prominently expressed in mouse chondrocytes (64,65). For that reason this type of ion channel has been incorporated into this initial model of the human chondrocyte. The rationale for doing this is based on the likelihood that this type of conductance is needed to explain electrophysiological responses to naturally occurring paracrine substances (e.g., ATP) or to cytokines that are liberated in the setting of acute or chronic inflammation (e.g., XXX). A further reason for incorporating this type of conductance into this early model is that this type of ion channel exhibits significant permeability to both Na+ and Ca2+ (68). Either or both of these cation species are likely to be important in regulating excitation secretion coupling and could modulate cell volume. It is also known that this type of ion channel can couple to purinergic receptors and/or to specific connexin proteins which function in the context of intercellular electrotonic communication, or as hemi-channels (see below).

[Figure 9 near here]

|  |  |
| --- | --- |
|  | (13) |

g) Connexin Mediated Current Flow

The chondrocyte from adult humans functions as an isolated cell. As a result it may seem that consideration of connexin function is irrelevant. However data from humans during early adolescence suggest that the growth plate of articular joints is characterized by small groups of directly opposed chondrocytes with expression of selected members of the connexin family being detectable using standard immunohistochemical approaches (66). Moreover it is now known that even in the case of adult isolated cells prominent ATP release can be observed (67). One plausible mechanism for this chemical or mechanical release is transient opening of HEMI channels due to expression of either pannexin or connexin subunits. For these reasons our model incorporates a connexin mediated conductance which, however, under physiological conditions is shut off or has a value of 0 pS.

## Implementation and Utilization of this Model

All individual transmembrane ionic currents above are defined by Equations 2-12. The ODE system (68) is solved for the primary vector of unknowns: Vm, [Na+]i , [K+]i , [Ca2+]i , [H+]i , [Cl-]i , aur and iur. All initial conditions were chosen from previous calculations that were run to steady state. This equation system is solved using LSODE (68). The corresponding GNU Octave code is available free and open source.

In this first order model we do not account for spatial variations in quantities of interest. As a result the human chondrocyte can be modeled by the following set of ordinary differential equations (ODEs) in time.

|  |  |
| --- | --- |
|  | (14) |

where,

|  |  |
| --- | --- |
|  |  |

**RESULTS**

## Model Characteristics

Experimental conditions described in [Clark et al. (13)](chondrocyte-model.html#Clarketal2011) were first replicated in the simulations to fit significant potassium currents used in the model. For this, the exterior concentrations of the various species in the model were set to experimental values ( [K+]o = 5 mM, [Na+]o = 140 mM, [Ca2+]o = 2 mM, pH = 7.4 for IK, Ca2+-act and IKur ) and ( [K+]o = 5 mM, [Na+]o = 145 mM, [Ca2+]o = 2 mM, pH = 8.5 for IK2pore ), and the potential was linearly ramped over 1 s from -130 mV to +100 mV. The evolution of the concentrations and currents were tracked in this period, and multiple simulations of this nature were used to fit parameters for IK2 pore, IK, Ca2+-act and IKur). Figure [2](#fig:potassium-currents) shows these individual currents and their fit to experimental values from [Clark et al. (13)](chondrocyte-model.html#Clarketal2011). The other currents used in the model are significantly smaller in magnitude, and have not been specifically fit due to lack of experimental data. (Figure [3](#fig:other-currents))

After this primary parameteristion, the overall model behaviour was studied under a linear voltage ramp from -130 mV to +90 mV under conditions matching experiments ([K+]o = 5 mM, [Na+]o = 140 mM, [Ca2+]o = 2 mM, pH = 7.4) to reveal that the overall voltage-current behaviour of the chondrocyte model reproduces the experimental data quite closely ([13](chondrocyte-model.html#Clarketal2011)). This comparison is shown in Figure [4](#fig:overall-behaviour) along with corresponding time evolution of the total current in the model.

The initial conditions used in the simulations were steady state values of the solution under same conditions used for the different numerical experiments. Figure [5](#fig:concentrations) shows that, when starting from a steady state solution, the concentrations in the model do not evolve over a relatively long simulation period of 30 min. The initial conditions for the concentrations used in the computations were [Na+]i = 2.814 mM, [K+]i = 121.59 mM, [Ca2+]i = 2.371e-06 mM, [H+]i = 6.188e-10 mM, [Cl-]i = 13.209 mM. When the model is perturbed from these conditions, it returns to steady state values in a similar time frame.

Model development has proceeded in defined stages starting with implementation of the minimal requirements for generation of a resting potential.

i) Background Currents

Output from our initial simulations is shown in the current voltage (I-V) relationship in Figure 2. This I-V curve illustrates what are termed 'background currents'. In this model these include the resting Na+, K+, and Cl- fluxes. Also shown is the small outward current that is generated by the electrogenic Na+/K+ pump that has been scaled to achieve a steady-state intracellular Na+ concentration of 10-12 mM. The remaining background currents, shown in Figure 2B, are those that are generated by the Na+/Ca2+ exchanger under resting and steady-state conditions corresponding to an intracellular Na+ concentration of 12 mM, and assumed free intracellular Ca2+ concentration of 3 x 10-8 M. In the mammalian chondrocyte there is also evidence for a background flux of Na+ and H+. Equations which capture this electroneutral ion transfer are included since it is anticipated that pH regulation and regulation of intracellular Na+ levels are of importance in both physiological and pathophysiological settings.

The scaling of current densities is dictated by the known input resistance of the cell, which is approximately 2 Gigohms. This value is denoted on the I-V curve by the thickened trace in the region of the range of resting membrane potentials that has been reported.

DISCUSSION:

i) Practical Applications of this Model

In the present state of its development, our model represents an original but only semiquantitative tool that can be used as one part of our multidisciplinary studies of electrophysiological and biophysical principles of the human chondrocyte. Our initial emphasis was on investigating the basis of the resting potential (Em) since our data set consists mainly of K+ current measurements in this non-excitable cell. Moreover, it is known that even very small changes in the resting membrane potential can strongly modulate factors that govern intracellular Ca2+ signaling and homeostatic mechanisms. There is also evidence that relatively small alterations in membrane potential contribute to dynamic regulation of cell volume. Both the strength and duration of ligand gated conductance changes are dependent upon membrane potential (Em) since this sets the electrochemical driving force.

The computations shown in Figures 2 through 6 in fact do not fully explain the ionic basis for the resting membrane potential in the human chondrocyte. Rather, they serve to illustrate that a range of membrane potential values is to be expected. This is because the net outward current that sets the resting membrane potential is very small even though it is the algebraic sum of background conductances as well as currents due to the electrogenic Na+/K+ pump and Na+/Ca2+ exchanger. In many recording situations the leak current through the patch pipette seal resistance can influence (and could even dominate) the observed 'resting potential'. In addition spontaneous changes in this seal resistance can give the impression of an unstable resting potential in these cells.

Nevertheless, our computational work does provide a basis for addressing the question, 'what are the physiological roles of each of the four K+ currents that have been identified and characterized in human chondrocytes'? At the outset one can appreciate from the computed I-V curves that any, or all of these currents could: i) significantly hyperpolarize the resting potential or ii) repolarize the chondrocyte after it had been depolarized by e.g., the effects of mechanical activity or by a ligand gated conductance.

An example of this, illustrated in Figure 7, is based on simulation of the electrophysiological response observed following activation of the TRP channels that are known to be expressed in human chondrocytes. A characteristic TRPV4 I-V curve is shown in Figure 7A. The effect of this channel mediated current on the resting membrane potential can be deduced by adding this current to the steady-state I-V relation illustrated in Figure 7C. The resulting net current is shown in Figure 7B. From this it is clear that activation of TRP channels results in a transient depolarization of the chondrocyte. However and possibly more importantly, it is also apparent that TRP channel activation results in a significant influx of Ca2+ and Na+ under circumstances in which the chondrocyte membrane potential is negative to approximately -40 mV. The TRIP channel induced depolarization is limited in magnitude and eventually transformed into a repolarization by the voltage dependent activation of the delayed rectifier K+ current, IK-DR. It follows that under circumstances in which the TRP channel results in a significant increase in intracellular Ca2+, the Ca2+ dependent K+ current, IK-Ca will also be activated. This will hyperpolarize the membrane potential and this potential will be maintained until intracellular Ca2+ and/or Ca2+ dependent signaling mechanisms reset to resting values.

[Figure 10 near here]

Consideration of the physiologic milieu within the articular joint identifies conditions under which some of these K+ currents may be modulated significantly. For example, the extracellular fluid is hypertonic. The effects of osmolarity on voltage gated K+ currents have been studied extensively. In brief, as a consequence of changes in surface charge shielding alterations in superfusate tonicity can significantly shift the steady state gating of currents such as IK-DR. Our experimental work is done under fixed isotonic conditions. The literature would suggest that an increase in osmotic strength would cause a shift to the right, in the depolarizing direction by perhaps 5 mV. This biophysical correction would make it unlikely that IK-DR makes any significant contribution to the resting potential. However, this alteration would position this current to initiate prompt repolarization following any transient depolarization, while also ensuring that the input resistance of the cell at the resting potential remained very high. This is a critical factor since a minimum of ion transfer is desirable in the relatively hypoxic milieu of the articular joint.

ii) Limitations of this Model

As presented, this mathematical model of chondrocyte electrophysiology provides a reliable platform for explaining and evaluating the experimental data in this field. This first order model also has proven utility for rationalizing and integrating genomic data from expression array profiles and as part of ion channel/antiporter drug target initiatives. Iterations based on the semiquantitative approach that is made possible by this model can also be useful for designing new experiments that are aimed at revealing cellular mechanisms that govern/modulate excitation-secretion coupling. Finally, given that the chondrocyte is in a unique, but yet not completely defined microenvironment our model provides a basis for explaining: i) the known biophysical effects of alterations in ionic strength of synovial fluid on ion channel voltage-dependent gating (zeta potential effects), ii) one common approach in attempts to account for the effects of cyclic stretch on ion channels - strain dependent alterations in channel gating voltage dependence or kinetics (cf. YYYY), and iii) much needed approaches for detecting and determining the limitations of present patch clamp technology that can bias, if not dictate, resulting data sets and influence their interpretation.

We recognize that at this stage our model has significant limitations. These include but are not limited to:

a) The absence of any comprehensive account or set of mathematical descriptors for intercellular Ca2+ homeostasis. Extension of the model to include simulations of the Ca2+ is needed before it can be used to explore the roles of Ca2+-activated K+ and/or Cl- currents.

b) Essential aspects of excitation-contraction coupling including the sources for the Ca2+ which triggers and supports secretion and the reuptake or extrusion mechanisms for this Ca2+.

c) Mathematical expressions that would allow rationalization of what has been termed 'the AM and FM modes of Ca2+ signaling' which will require consideration of intracellular Ca2+-dependent phosphorylation and dephosphorylation reactions and consideration of the Ca2+-dependence involved in transcriptional regulation of ion channel, antiporter and pump target molecules.

d) An improved understanding of and development of mathematical approaches that may account for the response of the chondrocyte to the ambient hypoxia within the articular joint. Initially this is likely to require formulation of equations for hypoxia-activated K+ channels.

e) Adaptation of published mathematical formulations for alteration of ion channel activity arising from cyclic biomechanical stretch or from transient changes in shear forces that impinge upon the chondrocyte.

Voltage-gated Sodium Channels

These have been found in some rabbit chondrocytes and some osteo-arthritic cartilage. They are likely constantly inactivated due to the voltages the chondrocyte. We instead model  as a sort of constant background Na+ leak.

External stimulation (matching experiments, e.g. cyclic stimulation):

|  |  |
| --- | --- |
|  | (13) |

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CONFLICT OF INTEREST:

None of the authors have any conflict of interest items to declare.

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**Some mechanism for sensing external pH**: ASIC channels (related to ENaC) are opened by extra-cellular protons (in the acidic environment of the chondrocytes ) and mediate an increase in Ca2+. There is also some other possible mechanism involving something called connexin-43.

**Small calcium-activated Potassium ISK**: A few reports suggest this might exist ([14](chondrocyte-model.html#BarrettJolleyetal2010),[5](chondrocyte-model.html#Halletal1996)). Tiny magnitude, by definition and role unknown. So we ignore it.

**Voltage-gated calcium channels**: T, L-type VGCC found in some cartilage. Others refute. Supposedly, aggrecan and collagen synthesis induced by electrical stimulation relies on this channel. We will claim not important for RMP, just tissue growth, and thus we do not consider.

**Epithelial sodium channels (EnaC)**: Not clear what role this plays in chondrocytes, though it has been identified. People speculate it has something to do with mechanotransduction  contributes to RMP. It is perhaps defective during osteoarthritis. We will point out we do not look at this as it pertains to mechanics.

**Aquaporin channels:** AQP or some other mechanism for transport of water seems super important to the functioning of the cell. Studies show loss of volume regulation with inhibition of AQP. But we will point out that we do not model it because it pertains to mechanics.

**NMDA channels:** This is an excitatory neuro-transmitter receptor. It is possibly linked to mechanotransduction.

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**Figure Legends**

### Figure [1](#fig:chondrocyte-model).

An illustration of the ion-selective channels included in the mathematical model of the chondrocyte.

### Figure [2](#fig:potassium-currents).

Potassium current-voltage relationships which have been fit to experimental values (in red) from [Clark et al. (13)](chondrocyte-model.html#Clarketal2011). The external concentrations correspond to the experimental conditions: [K+]o = 5 mM, [Na+]o = 140 mM, [Ca2+]o = 2 mM, pH = 7.4, except for IK2pore , where [K+]o = 145 mM, pH = 8.5.

### Figure [3](#fig:other-currents).

I-V relations for the other currents. These are not fit to experimental data, but used to tune simulation results.

### Figure [4](#fig:overall-behaviour).

Overall behaviour of the model when voltage is ramped from -130 mV to +90 mV in 1 s. The model output closely matches experimental data (red) from [Clark et al. (13)](chondrocyte-model.html#Clarketal2011).

### Figure [5](#fig:concentrations).

Time-evolution of the concentrations over 1800 s to show that the initial conditions we have chosen for the model were at steady state. The initial conditions for the concentrations used in the computations were [Na+]i = 2.814 mM, [K+]i = 121.59 mM, [Ca2+]i = 2.371e-06 mM, [H+]i = 6.188e-10 mM, [Cl-]i = 13.209 mM.

### Figure 6

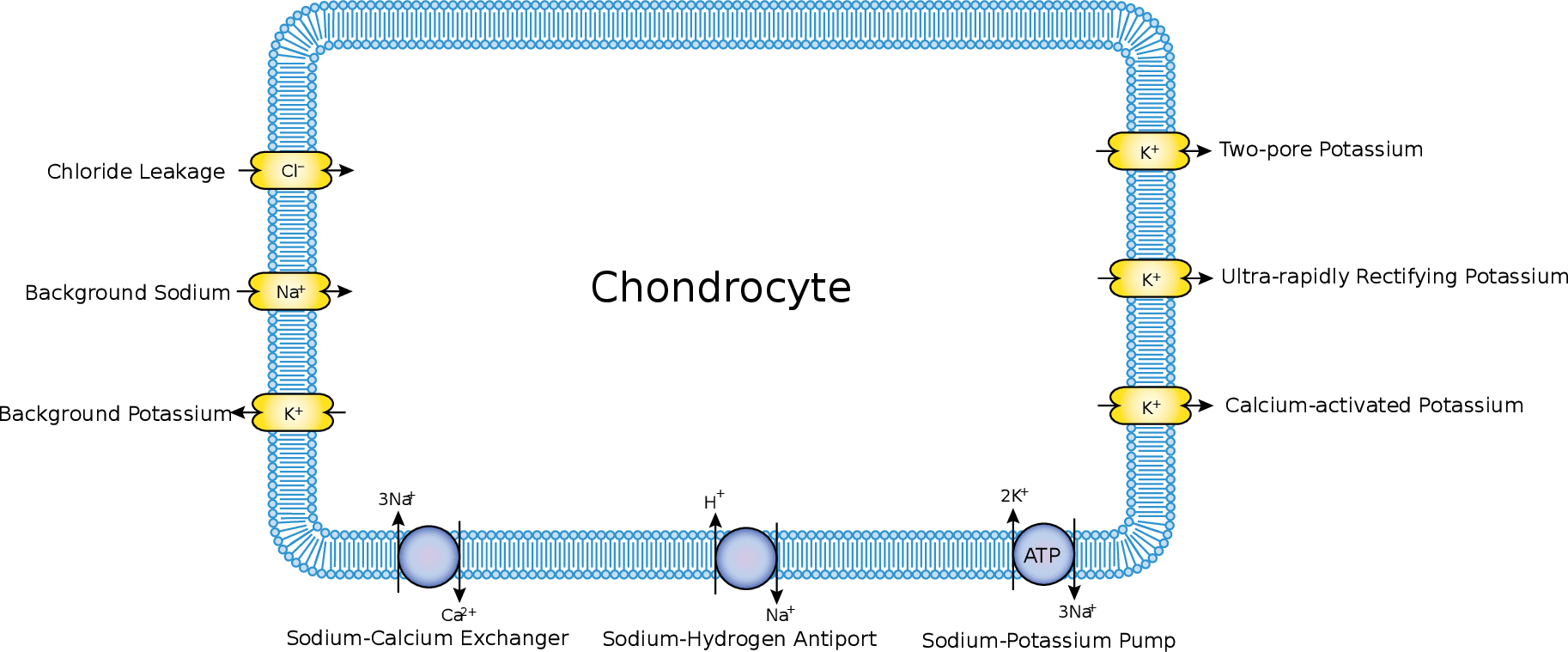
Evolution of the resting membrane potential with varying external potassium concentration. Note that while it is slightly more positive than experiments, it matches the qualitative behaviour quite closely ([13](chondrocyte-model.html#Clarketal2011)).

## Figure 7.

When the amount of IK2pore is varied from 100% to 0% (by blocking with increasing amounts of BUP), the RMP increases. These simulations were carried out at two different values of external concentrations [K+]o = 5 mM and [K+]o = 25 mM and results compare favorably with experimental data ([13](chondrocyte-model.html#Clarketal2011), Fig. 8B).

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**Figure 1:**



**Figure 2:**

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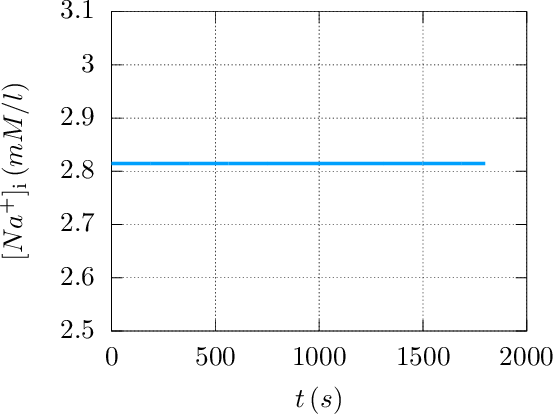
**Figure 3:**

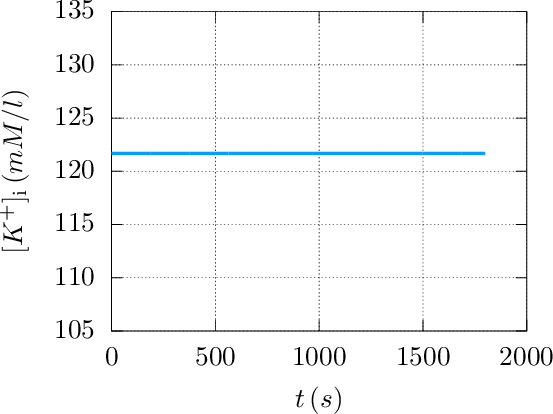
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**Figure 4:**

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**Figure 5:**

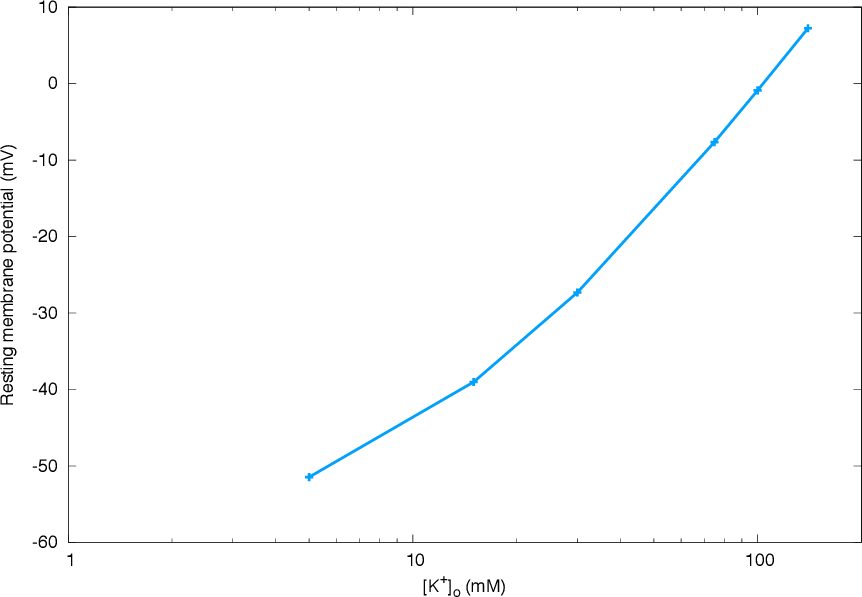


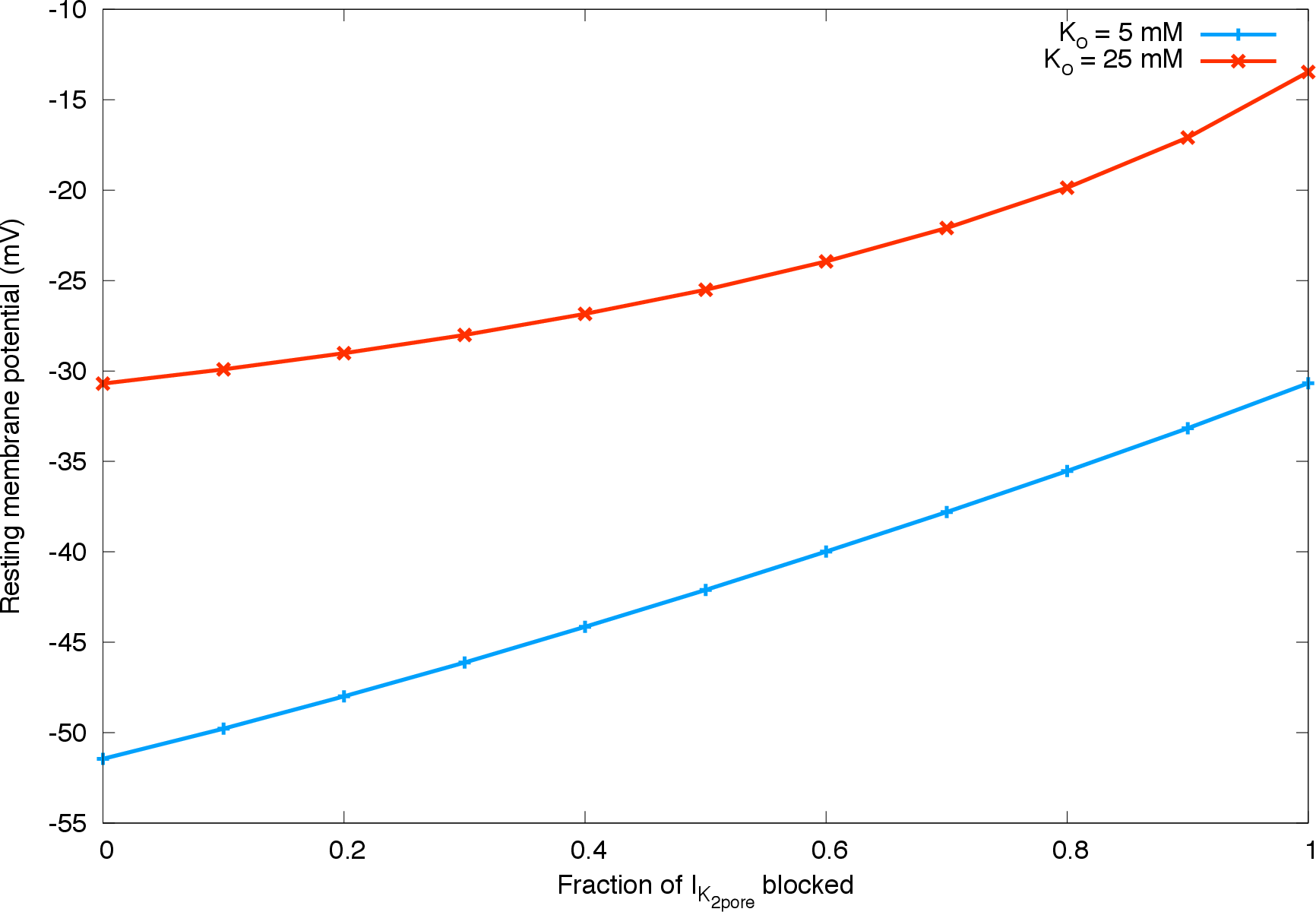


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**Figure 6:**

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**Figure 7:**



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#### Footnotes

...Harish Narayanan[1](chondrocyte-model.html#tex2html1)

Corresponding author. Address: Center for Biomedical Computing, Simula Research Laboratory, P.O. Box 134, 1325 Lysaker, Norway Tel.:+47 4003-4801, Fax: +47 6782-8201

Harish Narayanan 2012-10-05

APPENDIX:

Our model of seal resistance re: membrane potential