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Physiological Effects of the Electrogenic Current Generated by the Na⁺/K⁺ Pump in Mammalian Articular Chondrocytes

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Abstract

Background: Although the chondrocyte is a nonexcitable cell, there is strong interest in gaining detailed knowledge of its ion pumps, channels, exchangers, and transporters. In combination, these transport mechanisms set the resting potential, regulate cell volume, and strongly modulate responses of the chondrocyte to endocrine agents and physicochemical alterations in the surrounding extracellular microenvironment.

Materials and Methods: Mathematical modeling was used to assess the functional roles of energy-requiring active transport, the Na⁺/K⁺ pump, in chondrocytes.

Results: Our findings illustrate plausible physiological roles for the Na⁺/K⁺ pump in regulating the resting membrane potential and suggest ways in which specific molecular components of pump can respond to the unique electrochemical environment of the chondrocyte.

Conclusion: This analysis provides a basis for linking chondrocyte electrophysiology to metabolism and yields insights into novel ways of manipulating or regulating responsiveness to external stimuli both under baseline conditions and in chronic diseases such as osteoarthritis.

Keywords: articular cartilage, chondrocyte, Na⁺/K⁺ pump, Na⁺-K⁺-ATPase, electrogenic pumps, ion channels, mathematical model, osteoarthritis (OA), rheumatoid arthritis (RA)

Introduction

The biomechanical and physiological functions of mammalian articular joints are essential for locomotion, postural stability, proprioception, and motor learning. ¹⁻⁴ At the tissue and cellular levels, this results in requirements for a wide range of dynamic motion coupled with remarkable stability. An enabling component of this system is articular joint lubrication, made possible by coordinated activity and secretion of biological lubricants from the two principal cell types in synovial joints: chondrocytes and synovial fibroblasts. ^{5,6}

It is noteworthy that in the setting of chronic diseases of the articular joint, for example, rheumatoid arthritis (RA) or osteoarthritis (OA), significant changes in chondrocyte and synovial fibroblast function take place, and these contribute to reduced secretion of joint lubricants, ⁷ attenuated boundary lubrication, and altered joint loading. Reduced tribology results in increased boundary friction, load-induced wear of articular cartilage, impaired joint function, and eventual loss of proprioception. ⁸

The specific cell physiology-oriented focus of this study is the articular chondrocyte, which plays a key role in extracellular matrix (ECM) synthesis and degradation in all vertebrates. The presence of mature healthy chondrocytes and a full functional repertoire for these cells are essential for normal articular joint motion. Some of the important classes

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of transducer elements at the level of individual chondrocytes are ion channels, exchangers, or pumps that are expressed in the surface membrane. The significance of these integral membrane proteins, functioning individually or in concert, was first recognized with respect to the ability of chondrocytes to quickly and accurately volume regulate in response to even small changes in osmotic strength of the surrounding synovial fluid. The surrounding synovial fluid.

Within the last decade, a number of different ion selective channels and exchangers have been identified and shown to play essential roles in chondrocyte physiology. 16,19-24 Anv such ion channel being functional in chondrocytes may seem somewhat surprising to researchers working on excitable cells, since the chondrocyte is considered to be a nonexcitable cell. However, despite chondrocytes being nonexcitable cells, their dynamic membrane potential serves many cellular roles, ²⁵ is metabolically active, ¹⁶ especially during hypertrophy, ²⁶ and is capable of responses to commonly used drugs for treating hypertension.²⁷ Metabolic activity occurs at lower levels, especially in fully differentiated and senescent chondrocytes. 28,29 Indeed, chondrocyte precursors (i.e., the chondroblasts), which proliferate during joint development, and even some developmentally mature chondrocytes, are more metabolically active and are capable of robust ion channel mediated responses to biomechanical, pro-inflammatory, 30-36 and immunometabolic factors. 36,37 Essential to this fundamental physiological regulation is the ability of the chondrocyte to set and maintain an appropriate, stable resting membrane potential as mediated by ubiquitous ion channels, pumps, and exchangers in the plasma membranes of these cells.

Our main goal in this study was to obtain additional information concerning the role of the electrogenic Na⁺/K⁺ pump (also known as the Na⁺, K⁺-ATPase) in regulating the resting potential of the mammalian chondrocyte and thus ensure robust and stable physiological responses. A key starting point and motivation for our study was the previous demonstration of the presence of Na⁺/K⁺ pump proteins in the chondrocyte surface membrane using immunological and autoradiographic techniques and the molecular characterization of its multiple α , β , and γ isoform subunits. ^{39–41} This dataset, based on demonstration of relatively high affinity, saturable binding sites for ouabain (a classical sodium potassium pump antagonist), combined with more recent molecular studies that have identified transcripts and proteins of each of the known subunits $(\alpha, \beta, \text{ and } \gamma)$ of this integral membrane protein, ⁴² strongly suggests that in the mammalian chondrocyte the electrochemical gradients for sodium and potassium are established and maintained by this active or ATP-requiring pump mechanism. 43–45 Very recent proteomic studies have taken an agnostic and unbiased molecular discovery approach to exploring the "surfaceome" of chondrocytes, confirming the presence of multiple Na⁺/K⁺ pump isoforms in these cells.²⁴ These findings considered in the context of the extensive literature on Na⁺/K⁺ pump cell physiology^{46,47} make it very likely that in the chondrocyte the classical "coupled stoichiometry" that is characteristic of this enzyme is 3 Na⁺ pumped out of the cell, coupled with 2 K⁺ pumped into the chondrocyte cytoplasm (Fig. 1A). The resulting electrogenic current, oriented in the outward direction, would be expected to be one of the important factors in establishing the stable chondrocyte resting membrane potential. As shown in our previous article,³⁸ when this consideration is combined with the fact that the chondrocyte is a functional single cell with exceptionally high membrane resistance, it is likely that the electrogenic current due the Na⁺/K⁺ pump (although it is very small) can provide a 10–20 mV contribution to the chondrocyte resting potential.

The mathematical modeling that forms the basis of this article represents a continuation of our studies of chondrocyte electrophysiology, with an emphasis on understanding the basis for the resting potential and its physiological implications for cartilage function. Modeling and simulation here serve as knowledge integrators, unifying results from diverse tissues and experimental sources to offer new insight and to advance new hypotheses. Theoretical studies of this type are thus essential components of ongoing efforts to address important gaps in the background knowledge of the chondrocyte phenotype, mechanisms for cell volume regulation, transmitter and paracrine modulation, responses to anabolic and proinflammatory mediators in the context of ECM turnover, identification of early disease markers in chondrocytes, and perhaps most importantly, drug development for OA.

Methods

Simulations essential for the illustrations in this article were performed using our previously published model of the resting membrane potential of the chondrocyte and related intracellular calcium homeostasis.³⁸ In the present study, the physiological roles of the Na⁺/K⁺ pump are explored more fully by carrying out simulations in consideration of the unique environment of the chondrocyte in the synovial joint in terms of the *temperature* and the *ionic composition* of the extracellular milieu.

Temperature considerations

While temperature measurements of, for example, the human knee vary substantially dependent on the method, intrasurgical measurements suggest values between 31.5°C and 33.5°C for a healthy knee joint 48–50; as these values arise in the likely context of trauma and/or inflammation, a truly healthy knee joint at rest might have a temperature closer to 25°C±3°C, as suggested by external measurement methods. In contrast, joint temperatures are elevated in the context of RA (34–36°C) and OA (30–37°C), as probably attributable to chronic inflammation in RA and low-grade inflammation in OA. Thus, we have here considered the basal, "room" temperature of the previously published model, 23°C, to reflect a near-physiological temperature for the healthy synovial joint and chondrocyte environment; simulations also consider, in contrast, an elevated joint capsule temperature of 37°C to consider downstream effects on Na⁺/K⁺ pump function and the chondrocyte in the pathophysiological context.

Extracellular ionic milieu

In addition, the parent model (Fig. 1C) has been modified such that it can generate data sets that take into account the exceptional ionic milieu of the immediate microenvironment of the chondrocyte (see, e.g., Fig. 2A inset). We use this expanded model to explore a putative role for the electrogenic Na⁺/K⁺ pump current in regulating the resting membrane in articular chondrocytes with particular focus on conditions likely to be found in vivo, that is, given the unique ionic environment of the chondrocyte matrix.

Differential subunit expression and pump function

In addition, knowledge that chondrocytes express a number of different isoforms of the subunits that make up functional sodium potassium pumps is taken into account. The contributions of subtype-specific combinations of functional sodium potassium pumps are illustrated by adjusting the affinity constants for $[\mathrm{Na^+}]_i$ and $[\mathrm{K^+}]_o$ binding, as explored further in Figure 4. We illustrate and discuss the resulting profiles and patterns of electrogenic pump currents and related changes in membrane potential.

Overall, our findings can be used to inform ongoing and future patch clamp electrophysiological studies. They also have the potential to guide investigators in the design and implementation of novel planar recording methods.^{51,52} In the case of ion exchanger and pump activity generated by intracellular organelles, these approaches appear to be required for resolving and understanding these very small changes that can regulate fundamental properties of non-excitable cells such as the chondrocyte.

More specific information as to the current simulations performed using the model, including details regarding implementation in Matlab and parameter sets, may be found in the Supplementary Data. The model itself may be accessed at https://github.com/mmaleck/chondrocyte

Results

The sodium potassium (Na⁺/K⁺) pump makes an important contribution to setting up and maintaining the membrane potential.⁵³ The main goal of this study was to utilize mathematical modeling to explore the contribution of the electrogenic Na⁺/K⁺ pump to the resting membrane potential in chondrocytes isolated from healthy adult human articular joints. Na⁺/K⁺ pump expression has been demonstrated in primary bovine chondrocytes isolated from healthy joints,³ human chondrocytes in healthy and diseased joints in situ, 41 and human chondrocyte-like cell lines. 42 The functional presence of the Na⁺/K⁺ pump in chondrocytes has been further documented by related analyses of the molecular properties (isoform composition and expression levels),³ upregulation of the Na⁺/K⁺ pump in response to changes in extracellular Na⁺ concentration 40,54 and activity as demonstrated by ouabain binding, 40,43,44,54 and p-nitrophenylphosphatase activity in situ in both healthy and pathological samples. 41 Importantly also, the basis for the chondrocyte resting potential has previously been analyzed using mathematical modeling. ^{38,55–57}

However, none of these studies focused on defining the functional roles of the electrogenic current generated by the Na $^+$ /K $^+$ pump under pathophysiological conditions, for example, mimicking temperatures measured in OA ($\sim 37^{\circ}$ C), as opposed to the likely temperature found in healthy articular joints ($\sim 23^{\circ}$ C), also conditions that are typical of electrophysiological studies done using patch clamp methods under "room" temperature. Our first set of simulations involved making only one change to the parameters that govern the electrophysiological behavior of our human chondrocyte model. In this formulation, the mathematical expression for the electrogenic Na $^+$ /K $^+$ pump includes the ability to account for temperature differences by changing the Q₁₀ parameter that regulates the turnover rate of this enzyme. Based on the standard Q₁₀ (3.0) for a membrane-bound integral membrane

enzyme⁵⁸ the change from the baseline conditions (23°C) in this model to pathophysiological conditions (37°C) resulted in a substantial increase in the steady-state electrogenic current (Fig. 1D). These two superimposed steady-state current-voltage (I-V) plots clearly demonstrate that over a broad range of membrane potentials, and in particular, in the range of membrane potentials (-70 to 0 mV) that is most relevant to chondrocyte physiology, the outward electrogenic current increases by a factor of approximately 3-4. The curvilinear characteristic of both steady-state I-V curves reflects the intrinsic voltage dependence of the enzymatic reactions that result in coupled net electrogenic (3 Na⁺ in for 2 K⁺ out) fluxes that produce the so-called "pump current." ⁵⁹ These first findings strongly suggest that the Na⁺/K⁺ pump could modulate the resting membrane potential of the chondrocyte in health and disease.

The next set of mathematical simulations was performed in an attempt to illustrate the importance of the electrogenic current generated by the Na⁺/K⁺ pump for regulating or stabilizing the chondrocyte membrane potential. Figure 2A, based on our previously published mathematical model, 36 illustrates our working hypothesis for the ionic basis for this resting membrane potential. In brief, a background Na⁺ current (see also Supplementary Data and Supplementary Fig. S1) interacts with outward currents generated by 2-pore K⁺ channels and the outward electrogenic Na⁺/K⁺ pump current. Current generated by the Na⁺/Ca²⁺ exchanger and a Cl- conductance are also present but these are not the focus of this study. Note that under room temperature conditions (23°C) this complement of currents balances to generate a resting potential of approximately -40 mV, considering synovial ionic concentrations and generic affinities of the sodium-potassium pump for sodium and potassium ions, as published previously.³⁸ In the modified model used here (Supplementary Data), synovial electrolyte concentrations generate a chondrocyte resting membrane potential of approximately -60 mV. Note that the unique ionic milieu of the chondrocyte matrix may further hyperpolarize this resting membrane potential significantly with respect to the predictions using synovial ionic concentrations, even at room temperature/under healthy conditions (approximately -67 mV).

The analogous superimposed I-V plots that comprise Figure 2B were computed under conditions designed to mimic pathophysiological conditions (37°C) the chondrocyte may experience, for example, in OA. Specifically, the temperature dependence of the Na⁺/K⁺ pump was adjusted based on a Q_{10} value of 3.0, whereas the temperature dependence of the ion channel-mediated fluxes was adjusted using a Q₁₀ value of 1.2 for each.^{60,61} The combination of the very small sizes of the baseline (resting conditions) channel-mediated current in this small and very high resistance cell and the strong temperature dependence of the electrogenic current of the Na⁺/K⁺ pump results in an increased outward current due mainly to the Na⁺/K⁺ pump. Although still relatively small, this current would be expected to result in a significant and maintained hyperpolarization of the resting potential (20– 25 mV) in the chondrocyte under pathophysiological temperature; indeed, the chondrocyte resting membrane potential apparently can be hyperpolarized to approximately -92 mV at

It is important to evaluate and understand overall homeostatic control of intracellular electrolytes and related changes

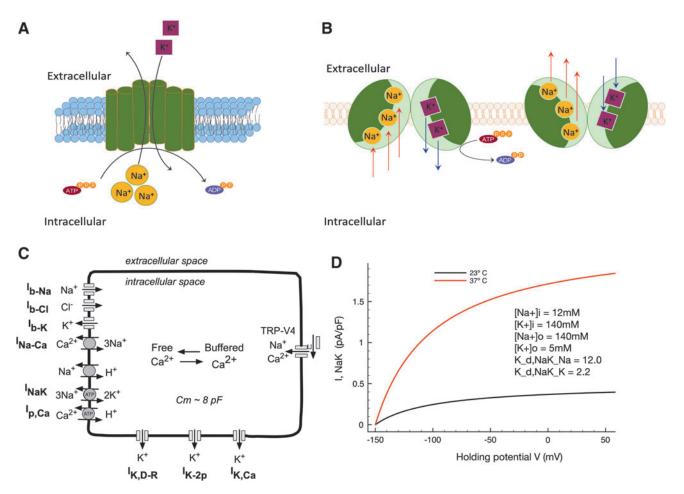
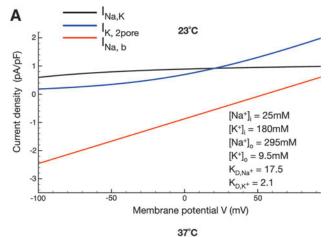


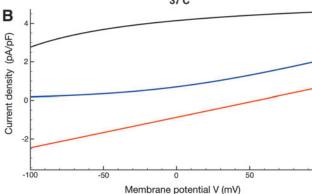
FIG. 1. Modeling of the chondrocyte resting membrane potential and demonstration of the effects of temperature on the electrogenic current generated by the Na⁺/K⁺ pump in a human chondrocyte preparation. (**A, B**) Present an illustration of the structure and function of the electrogenic sodium-potassium ATPase whose role in chondrocyte electrophysiology is investigated here; each pump turnover results in 3 Na⁺ ions' expulsion from and 2 K⁺ ions' inclusion into the cell, generating a net outward current. (**C**) Shows an illustrated schematic of the mathematical model of the chondrocyte resting membrane potential used here, as previously published.³⁸ (**D**) Explores the steady-state voltage dependence of the electrogenic pump Na⁺/K⁺ pump current density, given through our previously published model.³⁸ Strong temperature dependence is revealed, and the steady-state current at healthy joint temperature (23°C, black trace), as well as at pathophysiological temperature (37°C, red trace), is illustrated. Both I-V curves also show the curvilinear waveform that is due to the intrinsic voltage dependence of the Na⁺/K⁺ pump in mammalian cells. Intracellular and extracellular ion concentrations as initial modeling conditions are assumed (as previously) to be those measured in synovial fluid (as shown in inset).

in osmotic strength when studying active transport mechanisms in very small cells such as the chondrocyte. 10,62 Accordingly, we have tracked and illustrated time-dependent changes in intracellular Na⁺ and Ca²⁺, as well as extracellular K⁺, in the in silico conditions in which our baseline mathematical model operates. Figure 3A confirms the steady-state changes in the Na⁺/K⁺ pump current due to a step change in temperature; four sets of time-dependent results are shown in Figure 3B. The upper left plot shows the time-dependent change in chondrocyte membrane potential that results from a step change in temperature (and hence pump turnover rate) from 23°C to 37°C. Corresponding changes in intracellular Na⁺ and K⁺ confirm that there are no significant changes from the starting conditions or baseline model values. Somewhat similarly, although intracellular Ca⁺ increases approximately twofold soon after the temperature change, this stabilizes very near the 0.2 mM value still characteristic of most healthy mammalian cells. In summary, therefore, although the elec-

trogenic current can contribute to a very substantial hyperpolarization of the resting membrane potential, its activity does not significantly alter intracellular electrolyte homeostasis in this model. This is important since the electrochemical gradient for Na⁺ is a primary variable in regulating chondrocyte Na⁺/H⁺ ion exchange and hence intracellular pH and in modulating Na⁺/Ca²⁺ exchange in chondrocytes⁶³ as it does in cardiomyocytes.⁶⁴ It is known that changes in intracellular Ca²⁺ can alter Na⁺/K⁺ pump activity in most mammalian cells.⁶⁵ In addition, the ability of the Na⁺/K⁺ pump to stabilize intracellular Na⁺ and thus contribute to medium- and long-term cellular volume regulation is well known.⁶⁶

In the mammalian articular joint, electrolyte levels and particularly those for *intracellular* Na⁺ and K⁺ may deviate significantly from those in standard plasma or intracellular levels found in other tissues, for example, skin or muscle. ^{10,67} These differences and the resulting changes in Na⁺/K⁺ pump activity and baseline chondrocyte conductances were a focus





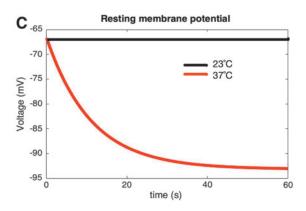


FIG. 2. Temperature-dependent contribution of the Na⁺/K⁺ pump electrogenic current to the chondrocyte resting membrane potential. The superimposed I-V curves in (A) were generated using our published model of the chondrocyte resting potential as shown, assuming a temperature of 23°C and measured and estimated ionic concentrations within the chondrocyte matrix. The analogous superimposed I-V curves shown in (B) were generated using the same model after the Na⁺/K⁺ pump current was adjusted to pathophysiological temperature (37°C) based on a Q₁₀ of 3.0 and the channelmediated background currents were adjusted using a Q₁₀ of 1.2. The two records of membrane potential in (C) illustrate the significant but relatively small effect of the electrogenic current due to the Na⁺/K⁺ pump at 23°C contrasted with the much larger hyperpolarization generated by this pump current at 37°C. As shown, the additional Na⁺/K⁺ pump current simulated at pathophysiological temperature in the chondrocyte matrix hyperpolarizes the chondrocyte resting membrane potential by $\sim 25 \text{ mV}$ (approximately -67 mV at 23°C and approximately -92 mV at 37°C).

of our original paper³⁸ and have been studied previously in physiological and pathophysiological settings. 10,14,16,43,44 These considerations, when combined with the demonstration that the healthy bovine and human chondrocytes express a number of different isoforms of the α , β , and γ subunits of the Na⁺/K⁺ pump protein complex, ^{39,41} raise important questions concerning the relative sizes of the electrogenic current and related changes in membrane potential that would be expected due to predominant expression of specific combinations of the α , β , and γ subunits. 42,45 We have approached this by simulating several of the combinations of expression of α and β subunits that are specified in a comprehensive review of the molecular physiology of the Na⁺/K⁺ pump⁶⁸ and more recent studies of the kinetic transitions of the movement of Na⁺ and K⁺ ions through the pump.⁶⁹ Results expressed in terms of temperature-dependent development of steady-state electrogenic currents for five subsets of data are shown in Figure 4. The most common physiological condition, assuming an exclusive $\alpha 1$, $\beta 1$, subunit composition with the affinity constants as shown in the inset, is shown in black. Note that the maximal current is $\sim 4.5 \text{ pA/pF}$ and that the steady-state membrane potential is approximately -92 mV at 37°C (membrane potential traces not shown). The red trace illustrates an identical calculation done under the assumption of a predominant $\alpha 2$, $\beta 1$, subunit expression. The corresponding higher affinity for the intracellular Na⁺ site on the Na⁺/K⁺ pump predictably results in a somewhat larger steady state electrogenic current (nearly 5.0 pA/pF) and larger hyperpolarization of the resting membrane potential to approximately -95 mV. In contrast, the dark green trace simulates the effect of predominant expression of the $\alpha 3$, $\beta 1$, subunits; based on the markedly decreased affinity for intracellular Na⁺ of this combination, the maximal electrogenic current reaches ~ 2.5 pA/pF and the chondrocyte resting membrane potential stabilizes at -78 mV.

Discussion

Main findings

This computational work, based on our published model of the healthy adult chondrocyte membrane potential, 38 confirms that the electrogenic current generated by the Na $^+/K^+$ pump is strongly temperature dependent. The resulting net outward current is relatively small. However, it can have substantial hyperpolarizing influences on the resting membrane potential of the chondrocyte due mainly to the very high input resistance, ~ 10 Giga-ohms 14,55,56 of this cell. In both physiological and pathophysiological settings it is likely that the Na $^+/K^+$ pump in the chondrocyte is strongly activated due to the relatively high intracellular Na $^+$ levels ($\sim 20 \ \text{mM}$ or more) in this cell type. 10,43,70

Previous findings

Hall et al. first reported evidence for energy-requiring active transport of K⁺ across the surface membrane of healthy adult mammalian chondrocytes. This initial observation was supported and put into a conventional cell physiology context by Mobasheri et al. 9 who adapted H-labeled ouabain binding methods to demonstrate substantial expression of the Na⁺/K⁺ pump alpha subunit in mammalian chondrocytes. Mobasheri et al. then confirmed and extended these

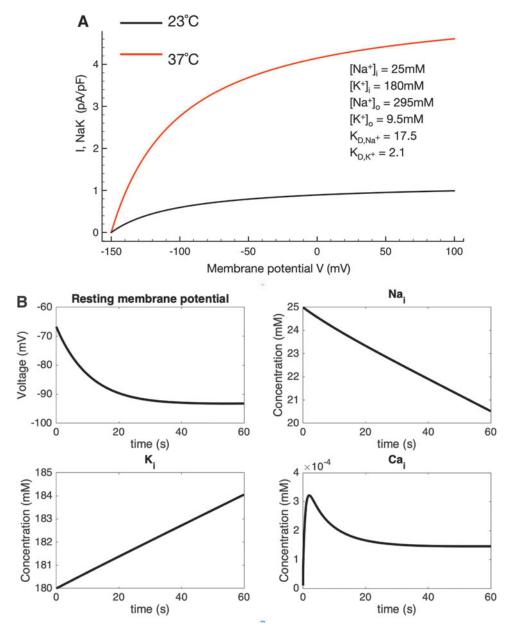


FIG. 3. Intracellular electrolyte homeostasis in the setting of altered Na $^+$ /K $^+$ pump activity/turnover rate due to a step change in temperature (23–37°C). As noted, the electrogenic Na $^+$ /K $^+$ pump current is changed significantly when intracellular and extracellular ion concentrations are altered to reflect the ionic milieu of the chondrocyte matrix found in vivo (shown in inset, A); compare to simulations performed assuming a baseline synovial fluid environment (Fig. 1D). In (A) we illustrate steady-state voltage dependence and the temperature dependence of the Na $^+$ /K $^+$ pump current. The steady-state current at 23°C (black trace), as well as at 37°C (red trace), is shown. The additional Na $^+$ /K $^+$ pump current simulated at a pathophysiological temperature hyperpolarizes the chondrocyte resting membrane potential by $\sim 25 \, \text{mV}$ at 37°C in the chondrocyte milieu. (B) Also shows the time-dependent hyperpolarization of chondrocyte membrane potential resulting from a step change in temperature from 23°C to 37°C, as well as the corresponding alterations in intracellular Na $^+$, K $^+$, and Ca $^{2+}$. In the cases of Na $^+$ and K $^+$ these changes are small: the shift in, for example, [Na $^+$]_i is about a 20% change over the time course measured and likely to be of small physiological relevance overall. The intracellular Ca $^{2+}$ level changes transiently (perhaps due to the intrinsic voltage dependence of the Na $^+$ /Ca $^{2+}$ exchanger or specific model-dependent features of intracellular Ca $^{2+}$ buffering) but quickly stabilizes near 0.2 μ M, an accepted level for a resting mammalian cell.

findings based on experimental ³H-labeled ouabain binding and confocal immunofluorescence microscopy work. ^{39,40} They demonstrated regulation of surface expression by changes in levels of intra- and extracellular Na⁺ in isolated cells ^{44,72} and in the ECM of articular cartilage from healthy

bovine joints⁴³ and also reported a scheme for overall Na⁺ regulation based on data that identified functional roles in the chondrocyte for Na⁺/Ca²⁺ exchange, Na⁺/H⁺ exchange, and antiporter exchange due to Na⁺/K⁺/Cl⁻ expression. ¹⁰ These articles and subsequent work have specified the essential role

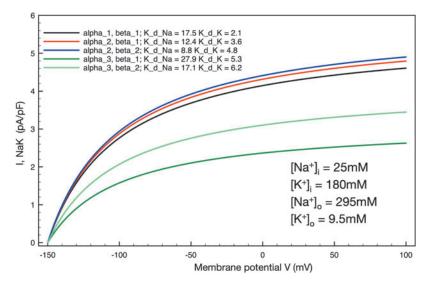


FIG. 4. Simulations of the effects on steady-state electrogenic Na⁺/K⁺ pump currents due to assumed changes in the α and β subunit composition of the Na⁺/K⁺ pump complex in an adult mammalian chondrocyte. Illustrates five different combinations of α and β subunits and the corresponding changes in maximum steady state electrogenic currents and alterations in chondrocyte resting membrane potential. The steady-state voltage dependence and overall Na⁺/K⁺ pump expression held constant, the altered expression of α and β subunits (as specified in the inset, upper left), were modeled by changing the Na⁺/K⁺ pump affinities for [Na⁺⁻]_i and [K⁺]_o known to be associated with these isozymes.⁶⁸ All ionic concentrations reflect the chondrocyte milieu, and all simulations were performed at 37°C to reflect a putative pathophysiological state, for example, OA. OA, osteoarthritis.

of the Na⁺/K⁺ pump and overall regulation of intracellular Na⁺ levels in volume regulation of the chondrocyte. It is also known that regulation of volume in the chondrocyte depends, in part, on the membrane potential of these cells. ^{20,73}

Physiological effects of the electrogenic Na⁺/K⁺ pump in chondrocytes

Insights gained from our mathematical modeling support the working hypothesis that the Na⁺/K⁺ pump can strongly regulate the resting membrane potential in chondrocytes from healthy and diseased adult articular cartilage. Specifically, it is plausible that due to its expression levels, turnover rates, 74 and intrinsic voltage dependence, 75,76 this pump mechanism produces a hyperpolarizing influence that can be as large as 30 mV. This hyperpolarization would be expected to modulate volume regulation; in addition, however, it is also likely to markedly alter the overall electrophysiological function or electrophysiological operating point of the chondrocyte.⁷⁷ This is because a number of the other ion channels that are expressed in the chondrocyte, for example, L-type Ca²⁺ channels, ^{15,27,78–80} delayed rectifier K⁺ channels, ⁵⁵, ⁵⁷ and 2-pore K⁺ channels ⁵⁶ exhibit strong intrinsic voltage dependence in the range -40 to -80 mV. Accordingly, the hyperpolarizing influence of the Na⁺/K⁺ pump significantly regulates the activation and/or deactivation of these (and perhaps other) ion channels in the chondrocyte.

We note, however, that the singular focus on the physiological and pathophysiological effects of the electrogenic current produced by the Na⁺/K⁺ pump in this study could be somewhat misleading. When the chondrocyte is stimulated by stretch, or activated by ligands such as histamine or ATP, agonist-induced ion fluxes through piezo, ⁸¹ Cl⁻, ⁷⁷ or TRP channels ^{14,77,82,83} will reduce the input resistance of the cell.

The resulting parallel conductance will partially "shunt" the influence of the electrogenic current generated by the Na⁺/K⁺ pump. In addition, in both health and disease, the chondrocyte exists and functions in a relatively hypoxic environment. ATP as the principal energy source of energy for chondrocytes and limit pump activity to a range that is less than the maximal currents shown in the Figure 1.

Concluding remarks

Although we acknowledge the possibility that the Na⁺/K⁺ pump function in mammalian chondrocytes may be modulated by altered expression levels of selected isoforms of the α , β , and γ subunits, our analysis is not sufficiently complete to fully examine the consequences as has been done in other tissues, for example, skeletal muscle.⁷⁴ It is known that the Na⁺/K⁺ pump can be strongly modulated by altered redox conditions 90 such as those that occur in sterile inflammation or "low grade inflammation" in the context of chondrocyte biology and OA. 91,92 Our work provides a basis for this type of analysis, but important pathophysiological effects such as this have not been studied. Finally, both classical findings and more recent detailed analyses have drawn attention to conditions under which changes in intracellular Ca^{2+} can markedly alter the function of the Na^+/K^+ pump. 93 Our model, at its present state of development, cannot be used to simulate these Ca2+-dependent effects due to the simplistic formulations now used for intracellular Ca²⁺ buffering and Na⁺/Ca²⁺ exchange; in the absence of mathematical descriptors for Ca²⁺ pumps; and Ca²⁺-sensitive channels localized to the endoplasmic reticulum. 94 These additions and other improvements will be needed before the mathematical modeling approach used in this study can be extended to

analysis of ion homeostasis and the chondrocyte channelome ^{16,22} in a more physiological context, specifically in chondron units, which represent the chondrocyte and its immediate pericellular environment. ⁹⁵ Further model development is also needed before our simulations can provide insights into the altered articular joint electrolyte homeostasis resulting from disease-producing point mutations in one or more of the Na⁺/K⁺ pump subunits or accessory proteins. ⁶⁹

From a tissue engineering perspective, there is ongoing interest in the Na⁺/K⁺ pump, ion transport, and the modulation of intracellular Na⁺ by pharmacological agents such as ouabain and bumetanide as in vitro treatments for altering intracellular ion concentrations as a viable method for manipulating ECM synthesis by chondrocytes and enhancing the mechanical properties of engineered articular cartilage.⁹⁶

Finally, in the context of drug development and screening for diseases such as OA, which is known to be characterized by cellular senescence, or "chondrosenescence," ^{29,97} the Na⁺/K⁺ pump has already been identified as a candidate target for modulation by cardiac glycosides, re-entering the limelight as classical cardiotonic drugs reinvented as senolytic compounds. ⁹⁸ Thus, the modeling approaches described in this article can support drug development for OA and related osteoarticular disorders, as well as understanding of the basic cellular physiology and pathophysiology of joint tissues.

Authors' Contributions

W.R.G. drafted the article. M.M.M. performed the mathematical modeling. M.M.M., P.M.-V., and A.M. edited the article and contributed data. All authors made significant intellectual contribution to the concept proposed in this article. All coauthors reviewed and approved the article before submission and confirm that the article has been submitted solely to this journal and is not published, in press, or submitted elsewhere.

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Author Disclosure Statement

No competing financial interests exist. The authors do not have any financial conflicts of interest in relation to this work.

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Supplementary Material

Supplementary Data Supplementary Figure S1

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