Some Kinetic and Steady-State Properties of Sodium Channels after Removal of Inactivation

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ABSTRACT To study the kinetic and steady-state properties of voltage-dependent sodium conductance activation, squid giant axons were perfused internally with either pronase or N-bromoacetamide and voltage clamped. Parameters of activation, τ_m and $g_{\text{Na}}(V)$, and deactivation, τ_{Na} , were measured and compared with those obtained from control axons under the assumption that $g_{Na} \propto m^3 h$ of the Hodgkin-Huxley scheme. $\tau_m(V)$ values obtained from the turn-on of I_{Na} agree well with control axons and previous determinations by others. $\tau_{\text{Na}}(V)$ values derived from Na tail currents were also unchanged by pronase treatment and matched fairly well previously published values. $\tau_m(V)$ obtained from 3 \times $\tau_{\text{Na}}(V)$ were much larger than $\tau_m(V)$ obtained from I_{Na} turn-on at the same potentials, resulting in a discontinuous distribution. Steady-state In $(g_{\text{Na}}/g_{\text{Na}}^{\text{max}} - g_{\text{Na}})$ vs. voltage was not linear and had a limiting logarithmic slope of 5.3 mV/e-fold g_{Na} . Voltage step procedures that induce a second turn-on of I_{Na} during various stages of the deactivation (Na tail current) process reveal quasiexponential activation at early stages that becomes increasingly sigmoid as deactivation progresses. For moderate depolarizations, primary and secondary activation kinetics are superimposable. These data suggest that, although m^3 can describe the shape of I_{Na} turn-on, it cannot quantitatively account for the kinetics of g_{Na} after repolarization. Kinetic schemes for g_{Na} in which substantial deactivation occurs by a unique pathway between conducting and resting states are shown to be unlikely. It appears that the rate-limiting step in linear kinetic models of activation may be between a terminal conducting state and the adjacent nonconducting intermediate.

INTRODUCTION

The events responsible for the voltage-dependent opening and closing of sodium channels in the excitable membranes of nerve and muscle cells have been classified into experimentally observable categories of activation and inactivation behavior. The nature of the relationship between activation and inactivation properties of sodium channels has received much attention, lately centering primarily on the question of whether the two phenomena represent independent mechanisms (e.g., Hodgkin and Huxley [1952 b]) or are coupled

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to each other in some manner (Goldman, 1976; Bezanilla and Armstrong, 1977; Gillespie and Meres, 1980).

The discovery of asymmetry currents (gating currents) related to the voltage-dependent gating of sodium channels allowed comparisons between the kinetic and steady-state properites of these gating currents and both the activation and inactivation properties of the sodium conductance (Armstrong and Bezanilla, 1977; Keynes and Rojas, 1974 and 1976; Schauf et al, 1977; Neumcke et al., 1976). The failure, on the one hand, to observe asymmetry currents exhibiting kinetics corresponding to the time-course of sodium inactivation, while, on the other, gating currents are "immobilized" by procedures that promote sodium conductance inactivation has led to the postulation of activation-inactivation coupling in which most of the apparent voltage dependence of inactivation is derived from the activation process (Armstrong and Bezanilla, 1977). Attempts to quantitatively correlate the voltage dependence and time dependence of gating current with the predictions of the Hodgkin-Huxley activation parameter, m, have met with moderate (Keynes and Rojas, 1976) or negligible (Neumcke et al., 1976; Bullock and Schauf, 1978) success.

Analyses of sodium conductance activation and associated intramembrane charge movements are complicated by the presence of the temporarily overlapping inactivation process(es). Previous descriptions of sodium conductance activation have, therefore, necessarily relied upon assumptions concerning the relation between activation and inactivation. The selective destruction of sodium inactivation in squid axons internally perfused with either pronase (Armstrong et al., 1973) or N-bromoacetamide (Oxford et al., 1978) offers the opportunity to "study Na activation directly, without interference from inactivation" (Armstrong et al., 1973). Thus far the results of such an investigation have not appeared. Rather, the use of pronase has been limited to characterization of both natural (Bezanilla and Armstrong, 1977) and pharmacologically induced (Yeh and Narahashi, 1977; Cahalan, 1978) inactivation of open sodium channels.

The aim of the present study is to compare direct experimental measurements of the voltage dependence and time dependence of sodium conductance activation after removal of inactivation with activation behavior expected from the assumption of independent processes (Hodgkin and Huxley, 1952 b). Also, more detailed information on activation in the voltage region close to the action potential threshold will be provided, and experimental protocols will be introduced that perhaps provide more insight into the nature of the transitions between closed and open channel states than can be obtained from simple changes in conditioning and test-pulse magnitudes. A preliminary account of portions of this work has appeared (Oxford and Yeh, 1977).

METHODS

Experiments were performed on single giant axons isolated from *Lol,go pealei* at the Marine Biological Laboratory, Woods Hole, Mass. Axons were cleaned of most adhering tissue, and the axoplasm was squeezed out with a tiny rubber roller and

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replaced with an artificial internal solution introduced via a micrometer syringe. Axons were then mounted in a Plexiglas chamber, continuously perfused both internally and externally, and voltage clamped with conventional axial wire techniques as previously described (Wu and Narahashi, 1973; Oxford et al., 1978).

Certain technical problems arise, or are amplified, in experiments in which the membrane sodium conductance is allowed to remain high for relatively long periods. In this regard, several details of the apparatus and procedures are particularly relevant to the data presented in Results and are mentioned here briefly: (a) The diameter of the axial Pt-Ir wire that injects the voltage control current was increased to 100 or 125 μ m to reduce polarization. (b) The Pt-plated silver electrode assemblies for measuring membrane current and electrically guarding that measurement from nonradial current flow were positioned on two sides of the axon as close as possible to the membrane. At very small depolarizations from the holding potential (-80 mV) , the small inward transient current was occasionally observed to follow an irregular, notched time-course when only a single guard electrode assembly was employed. Such patterns were never observed with double electrode assemblies. (c) The two ends of each axon located outside the electrically guarded region of the chamber were suspended in air. During experiments in which sodium inactivation was removed, the ends were moistened with tetrodotoxin solution (TTX, $1 \mu M$) to prevent the generation of prolonged action potentials in the unguarded, unclamped regions. In the absence of this precaution, electrical activity in the end regions results in substantial increases in holding current that can last for several tens of seconds. The increases were most often triggered by very small step depolarizations of the space-clamped region of membrane and were eliminated by \overline{TX} treatment. (d) All experiments described here were performed on axons bathed in artificial seawater containing onethird to one-fourth normal sodium concentration to reduce artifacts arising from residual series resistance (see below) and to further minimize the "end effects" mentioned above. (e) Voltage offsets arising from the various liquid junction potentials in the measuring system were eliminated for each axon by nulling the potential measured between the internal and external voltage electrode when the former was placed in a pool containing the particular internal perfusion solution, which was connected via a 3 M KC1 agar bridge to the external solution pool containing the latter electrode. The inverse of the corresponding junction potential between the internal and external solution was measured just before the axial electrode assembly was inserted in the axon and after withdrawal of the electrode at the end of the experiment. The "drift" in potential during the course of an experiment was never more than 4.3 mV, and typically less than 1 mV .

The response time of the voltage-clamp circuit was $2 \mu s$ (10-90% of a voltage step). Leakage and capacitative currents were reduced by an analogue transient bridge (see Armstrong and Bezanilla [1975]). Series resistance was determined by current-step methods and optimally compensated by positive feedback under voltage clamp. The compensation was usually enhanced by feeding the output of the transient bridge to the feedback potentiometer rather than the direct output of the current-to-voltage converter. This procedure greatly reduces the contribution of the early capacitative transient to the feedback, thus reducing ringing in the membrane voltage step and allowing higher feedback gain to achieve satisfactory compensation. A compromise resulting from this procedure is that series resistance compensation is less than adequate for the first $10-30 \mu s$ of the voltage step; however, conductance changes are normally difficult to quantify at such early times, rendering such data questionable in any event.

Experimental Solutions

Axons were externally perfused with a potassium-free, low-sodium artificial seawater containing ions in the following concentration (mM) : Na⁺, 100-150; tetramethylammonium (TMA^+) , 350-300; Ca^{++} , 50; Cl^- , 566; HEPES buffer, 5. External pH was adjusted to 8.0, and the osmolarity was $\sim 1,010$ mosM. At the beginning of each experiment axons were internally perfused with a standard internal solution (SIS) that consisted of (mM): Na⁺, 50; K⁺, 350; glutamate⁻, 320; F⁻, 50; phosphate buffer, 15. Sucrose was added to adjust the osmolarity to \sim 1,040 mosM, and the pH was adjusted to 7.3. During data collection, axons were internally perfused with a K-free internal solution in which $250-275$ mM Cs replaced the potassium and the osmolarity was maintained by the addition of sucrose.

N-Bromoacetamide (NBA) was obtained from Sigma Chemical Co., St. Louis, Mo., and pronase from Calbiochem, San Diego, Calif. Temperature in all experiments was maintained at $6-10 \pm 0.1$ °C by a Peltier device and electronic feedback circuitry.

Data Acquisition and Analysis

Membrane currents were photographed from oscilloscope traces using multiple vertical gains and time bases to optimize the graphical analysis of different kinetic components. In some experiments, currents resulting from voltage steps of equal amplitude and opposite polarity were averaged with an 8-bit signal averager kindly loaned by Dr. Howard Wang. In later experiments, currents were recorded with a 12 bit analogue-to-digital transient averager with dual sample rate capabilities similar to those of a design described elsewhere (Bezanilla and Armstrong, 1977). Currents could be sampled at a maximum rate of $10 \mu s$ /point. The averager was interfaced directly to a PDP 11/03 computer (Digital Equipment Corp., Maynard, Mass.) for data storage and analysis.

Activation kinetics were derived from sodium currents with intact inactivation behavior according to the assumptions of the Hodgkin-Huxley scheme for the sodium conductance. Details are presented in Results. After removal of inactivation with pronase or NBA, activation kinetics were derived by curve fitting procedures detailed in Results.

For convenience the following terminology is adopted here: "Activation" refers to the turn-on of sodium conductance as channels open after a step depolarization (more positive internal potential). "Inactivation" refers to the decay of sodium current (or conductance) as a step depolarization is maintained. "Deactivation" describes the decay of sodium current after a step repolarization (more negative internal potential), often referred to as a "Na tail." The time constants characteristic of activation, inactivation, and deactivation of the sodium conductance are τ_m , τ_h , and τ_{Na} , respectively. None of these terms is meant to imply any particular mechanism with which they may have been associated in previous literature. Their use here is simply to provide a familiar reference terminology.

RESULTS

Activation Kinetics of Na Channels with Intact Inactivation

The extraction of sodium conductance activation kinetics from sodium current records exhibiting both activation and inactivation behavior is clearly model dependent with regard to the inactivation process. Models in which activationinactivation coupling are proposed (e.g., Bezanilla and Armstrong [1977]) as yet lack a quantitative description of the voltage dependence of such coupling that is independent of knowledge about activation behavior alone. Thus, for convenience as well as for comparison with previous measurements, sodium activation is here assumed to obey kinetics of the type described by the Hodgkin-Huxley m parameter. The time-course of activation was determined from records of sodium current turn-on at various step depolarizations from a holding potential of -80 mV according to

$$
I_{\mathbf{Na}}(t) = I_{\mathbf{Na}}^0(1 - e^{-t/\tau_m})^a \cdot [f + (1 - f)e^{-t/\tau_h}], \tag{1}
$$

where $a = 3$ and f is the ratio of steady-state residual I_{Na} (commonly observed in perfused squid axons) to I_{Na}^0 (see also Eq. 3 of Cahalan and Almers, 1979). I_{Na}^0 and τ_h were determined by fitting an exponential to the late phase of inactivation of $I_{\text{Na}}(t)$ and extrapolating to the beginning of the voltage step

FIGURE 1. Comparison of experimental I_{Na} kinetics with I_{Na} computed from *m3h.* Dotted lines represent tracings of sodium-current records at the indicated potentials taken at random for axons with inactivation present *(top record)* or after NBA treatment *(bottom record).* Solid lines are solutions to Eq. 1 with the following parameter values: *(top record)* $f = 1, \tau_m = 468 \mu s, I_{Na}^0 = 0.75 \text{ mA}$ */* cm²; *(bottom record)* $f = 0.08$, $\tau_m = 253 \,\mu s$, $\tau_h = 2.33 \text{ ms}$, $I_{\text{Na}}^0 = 1.28 \text{ mA/cm}^2$.

 $(t = 0)$. The activation time constant, τ_m , was then determined by one of two methods. (a) The exponential fit to the inactivation time-course was vertically scaled by one-half, replotted (semilogarithmic) along with $I_{\text{Na}}(t)$, and the time coordinate of their intersection was used to compute τ_m after the method of Hille (1967). (b) Alternatively, $I_{\text{Na}}(t)$ was divided by $f + (1 - f) \exp(-t/\tau_h)$ to yield $I_{\text{Na}}'(t)$. τ_m was then determined from the slope of a straight line fit to a plot of $\ln (1 - (I_{Na}(t)/I_{Na}^0)^{1/3})$ vs. time (Keynes and Rojas, 1976). Both methods yielded comparable values of τ_m at the same membrane potential. The values extracted for I_{Na}^0 , τ_m , and τ_h as well as the choice of $a = 3$ were routinely tested by computing $I_{\text{Na}}(t)$ from Eq. 1 and visually comparing the fit with the raw data (see Fig. 1).

Deviations of the computed $I_{\text{Na}}(t)$ from the raw data at the beginning of the voltage step probably represent the contribution of asymmetrical charge movement (gating current), which can easily be seen in a single sweep when the transient bridge is properly adjusted or in averaged records from voltage steps of equal amplitude but opposite polarity. Furthermore, the possibility that activation and inactivation are coupled can alter this early time-course and render the quality of fit meaningless. The presence of a small delay in the onset of activation was detected (e.g., Keynes and Rojas [1976]), but, at the present holding potential of -80 mV, the maximum delay was 70 μ s (test pulse, 40 mV; temperature, 6°C) resulting in an error of only $\sim 15\%$ in τ_m . Preliminary observations of the test-pulse dependence of this delay agree with the data of Keynes and Rojas (1976) and result in an approximately constant error in τ_m as a function of voltage if the delay is neglected. The magnitude of this error is negligible with regard to the objectives of the present paper; however, further investigation of these delays will yield important clues concerning details of the activation kinetic scheme.

Values of τ_m determined as described above are shown for several different membrane potentials in Fig. 2 A. The data points (filled circles) are scaled to a temperature of 6.3 \degree C by assuming a Q_{10} of 3.0. The dashed line represents $\tau_m = 1/(\alpha_m + \beta_m)$ of the Hodgkin-Huxley model as derived from their original measurements. Fairly good agreement is obtained for potentials more positive than -20 mV, whereas τ_m values are slightly larger than expected for more negative potentials.

Na Activation Kinetics with Inactivation Removed

Internal perfusion with either pronase (0.2 mg/ml) or NBA $(0.2-1 \text{ mM})$ selectively removes the inactivation behavior of the sodium conductance (Armstrong et al., 1973; Oxford et al., 1978). The extent of removal was monitored at two voltages during either treatment and could be reasonably controlled by changing to normal internal solution or lowering the temperature to slow the rate of reaction. Both treatments are usually quite difficult to carry to completion. The peak sodium conductance is almost always reduced in these cases and its sensitivity to these manipulations varies greatly among axons. Removal of >90% of the inactivation behavior while retaining sufficiently large currents (in reduced external sodium) was the objective for each axon. After removal of inactivation I_{Na}^0 was set equal to the steady-state level of $I_{\text{Na}}(t)$; thus, $f= 1$ and Eq. 1 reduce to

$$
I_{\mathbf{Na}}(t) = I_{\mathbf{Na}}^0 (1 - e^{-t/\tau_m})^a.
$$
 (2)

This expression adequately fits the data points after determination of τ_m , as indicated above (Fig. 1, *top).* At very low depolarizations a somewhat greater lag in the activation of the conductance could be observed. This could correspond either to an increase in the value of a or to an additional delay, as mentioned above. The small amplitude of current at these potentials precludes a choice between the two possibilities inasmuch as both gave approximately equal fits to the records. The values of τ_m chosen in the two cases were slightly

FIGURE 2. Time constants for sodium activation and deactivation as a function of membrane potential. τ_m values (filled circles) and τ_{Na} values (open circles) **are shown for axons before (A) and after (B) pronase treatment. Data points from NBA-treated axons are represented by triangles in B. All data points are** scaled to 6.3°C, assuming a Q10 of 3. Dashed lines computed from the Hodgkin-Huxley model as $(\alpha_m + \beta_m)^{-1}$ at 6.3°C. Equations for the voltage dependence of α_m and β_m were taken from Palti (1971).

different, however, and so I report here those determined by assuming $a = 3$ **and a small additional delay.**

 τ_m values obtained from axons after removal of inactivation are shown in **Fig. 2 B at several membrane potentials. Data points after either pronase** treatment (filled circles) or NBA treatment (filled triangles) agree well with the values from axons with intact inactivation behavior and with $1/(\alpha_m +$ β_m) of the Hodgkin-Huxley model. The close correspondence between τ_m values before and after removal of inactivation indicates the relative insensitivity of such measurements to the relationship between activation and inactivation mechanisms embodied in contemporary models of the sodium conductance (see also Fig. 9).

Deactivation Kinetics before and after Inactivation Removal

Deactivation kinetics were obtained from sodium tail currents after repolarization from a depolarizing prepulse that activated the sodium conductance to various degrees. The prepulse and postpulse potentials, as well as the prepulse duration, were variable. The most common experiment involved depolarizing to the reversal potential (E_{Na}) and repolarizing to various potentials at the peak of the sodium conductance. This value of prepulse minimized polarization of the current electrode immediately before the tail current.

Deactivation proceeded as a single exponential at potentials more negative than \sim -40 mV. Tail currents at -80 mV that follow prepulses of various durations are shown in Fig. 3 A for a control axon and in Fig. 3 B for another axon after NBA treatment. These currents were well fit by single exponentials, and the corresponding time constants for each are given in the legend. Fig. 3 C shows tail currents at three different potentials obtained digitally and signal averaged (three sweeps each). Single exponentials fit to each record are superimposed on the currents. At more positive potentials, a second, slower component was observed. This component was removed after internal perfusion with pronase or NBA, suggesting that it represents inactivation kinetics (Fig. 3 D). At potentials at which a significant second slow exponential could be distinguished, the time constant τ_{Na} was determined by first fitting an exponential to the slow component and then peeling this fit away from the total current to reveal the faster exponential (Fig. $3 D$).

In Fig. 2 the time constants for deactivation (τ_{Na}) are plotted as a function of voltage from axons with intact inactivation (2λ) , open circles), after pronase perfusion (2 B, open circles), and after NBA perfusion (2 B, open triangles). Again, the time constants determined in the presence of inactivation agree well with those obtained after removal of inactivation. The τ_{Na} values coincide well with the τ_m values derived from sodium conductance turn-on to form nearly symmetrical bell-shaped distributions. This finding is very surprising, however, in view of the prediction of the Hodgkin-Huxley model that $\tau_m =$ $3\tau_{\text{Na}}$! Multiplying the averaged τ_{Na} values from all experiments by three as dictated by the Hodgkin-Huxley formulation yields the values of τ_m derived from deactivation shown in Fig. 4 A. The result is a discontinuous τ_m vs. E_m curve, with τ_m values at negative potentials much larger than expected from the Hodgkin-Huxley model.

The observation of such markedly slower turn-off of g_{Na} curiously is not unique to the present data. In Fig. 4 B, the average τ_{Na} and τ_m values from Fig. 2 A are replotted along with a variety of τ_{Na} values taken from the

FIGURE 3. Sodium tail currents upon repolarization from E_{Na} . (A) Axon 061978a, 8°C. Prepulse, $+25$ mV; postpulse, -80 mV. τ_{Na} values for $a-j$ are consecutively (μ s) 128, 215, 235, 225, 243, 238, 224, 235, and 219. (B) Axon 061678a. 5.1°C. Prepulse, $+20$ mV; postpulse, -80 mV. τ_{Na} values for $k-q$ are consecutively (μ s) 225, 265, 280, 300, 320, 315, and 321. (C) Axon 79JUL23a. 12°C. Na-free internal perfusate containing 300 mM Cs. Prepulse, $+80$ mV and 0.5 ms. Smooth curves are exponentials with time constants of 84 μ s (-80 mV), 124 μ s (-60 mV), and 218 μ s (-40 mV). (D) Axon 072376b. 9°C. Prepulse, +80 mV ; postpulse, -30 mV. Semilogarithmic plot of sodium tail current before (filled circles) and after (filled triangles) pronase treatment. Open circles are the differences between the filled circles and the straight line fit to the late portion of the control time-course. The time constant for this fit is 2.35 ms. The straight line fit to the open circles yields $\tau_{\text{Na}} = 510 \,\mu\text{s}$. Data points in pronase have been vertically scaled to visually overlap the open circles.

literature and symbolized as letters. Details of the sources for these points are provided in the legend. The average data reported in this paper agree quite well with the previously reported values. Curiously, even the average value of τ_{Na} from the original experiments of Hodgkin and Huxley (1952 b) (assuming

FIGURE 4. Mean τ_m and τ_{Na} values as a function of membrane potential. (A) Mean values of τ_m from Fig. 2 A (filled circles) and B (filled triangles) and mean values of τ_{Na} from Fig. 2 A (open circles) and B (open triangles). Open symbols are scaled \times 3 and replotted as half-filled symbols. \times represents the average of τ_{Na} values at -20 mV from two pronase-treated axons. (B) Mean values of τ_m and τ_{Na} symbolized as in A. Letters repressent τ_{Na} values adjusted to 6.3°C (Q_{10} = 3) obtained from the literature as follows: (A) average value from Table 1 of Hodgkin and Huxley (1952 a), assuming a resting membrane potential of -58 mV; (B) τ_{Na} in 22 mM external Ca, assuming a resting potential of -58 mV from page 231 of Frankenhaeuser. and Hodgkin (1957); (C) τ_{Na} at -60 mV from Table III of Armstrong et al. (1973); (D) τ_{Na} at -80 mV from Fig. 10 B of Meves (1974); (E) average τ_{Na} at -70 mV after 1-ms prepulses, from Table IV of Armstrong and Bezanilla (1977); (F) average τ_{Na} at -70 mV from Table 1 of Bezanilla and Armstrong (1975 a); (G) present study, τ_{Na} at -60 mV for axon 79JUL23a; (H) τ_{Na} at -76 mV computed from $(\tau_m)^{-1}$ given on page 684 of Chandler and Meves (1970) ; (J) values from Table III of Yeh and Narahashi (1977); (K) τ_{Na} from Fig. 7 a of Armstrong and Gilly (1979); (L) τ_{Na} from Fig. 9 of Cahalan and Almers (1979). The solid line represents a least-squares fit of the equation $\tau = (A \exp^{BV} + C \exp^{DV})^{-1}$, where V is the membrane potential and $A = 0.0095~\mu s$, $B = 0.080~\text{mV}^{-1}$, $C = 0.0001~\mu s^{-1}$, and $D = -0.080$ mV⁻¹.

a resting membrane potential of -58 mV) agrees well with the values reported here. The diversity of laboratories represented by these data and that both *Loligo pealei* and *Loligo forbesi* were used suggest that neither systematic error nor species differences appreciably contaminate the values. Furthermore, the precautions outlined in the Methods section minimized errors arising from series resistance properties in the present experiments. Accordingly, changing the magnitude of g_{Na} as much as fourfold, by either changing the external sodium concentration or the value of the prepulse, did not significantly affect the measured τ_{Na} values at a given potential.

Attempts were made to extend τ_{Na} measurements after pronase treatment to more positive voltages to compare with τ_m , which becomes increasingly smaller with depolarization. The expected small differences in steady-state activation (m) between prepulse and postpulse levels in this voltage range yielded only small, slow tail currents during transition to relatively large steady-state currents. Such small changes could be appreciably contaminated

FIGURE 5. Normalized steady-state sodium conductance $(g_{\text{Na}}/g_{\text{Na}}^{\text{max}})$ as a function of membrane potential before (filled circles) and after pronase treatment (open circles). The solid line represents $m³$ of the Hodgkin-Huxley model. The dotted line is a least-squares fit of a Boltzmann distribution to the data points below -10 mV with a midpoint potential of -23.4 mV and a slope of 7.38 mV/e -fold g_{Na} . The dashed line results from a simulation of g_{Na} via Scheme 3 using Hodgkin-Huxley rate constants, as done previously (Cahalan and Almers, 1979).

by any residual inactivation gating not removed by pronase or NBA, making estimates of τ_{Na} ambiguous in this range. Despite this uncertainty, it appeared as though τ_{Na} values continued to increase (see \times in Fig. 4 A) at more positive potentials, whereas τ_m measured from g_{Na} turn-on decreased. The large possible errors involved in such measurements will have to be further dealt with before this trend is established.

Steady-State Properties of Sodium Activation

Steady-state sodium conductances were determined from values of I_{Na}^0 obtained at various potentials as described above. g_{Na}^{V} was calculated by dividing I_{Na}^0 by the driving force on sodium $(E_m - E_{\text{Na}})$ and normalized with respect to the maximum value for each axon. In Fig. 5, $g_{\text{Na}}^0/g_{\text{Na}}^{\text{max}}$ is plotted as a function of E_m for three axons before and after treatment with pronase. The steady-state voltage dependence is not significantly different in the two cases and corresponds closely to that observed after NBA treatment (Oxford et al.,

1978). At first glance, the Hodgkin-Huxley m^3 parameter appears to describe the data adequately (solid line). However, the scatter is such that it is not possible to discriminate between this fit and others, such as a least-squares fit of a Boltzmann distribution of two-state gating particles to the data below -10 mV (dotted line) or a curve predicted from a sequential kinetic activation scheme terminated by an inactivation step (dashed line).

It has been pointed out that some sequential models can be discriminated from two-state constant dipole moment gating models by careful examination

FIGURE 6. $g_{\text{Na}} (g_{\text{Na}}^{\text{max}} - g_{\text{Na}})^{-1}$ vs. membrane potential. Data points from axons before (filled circles) and after (open circles) pronase treatment. Solid line is a least-squares fit to the data points for potentials more negative than -35 mV.

of the voltage dependence of activation in the region of negative membrane potential where channels just begin to activate (Almers, 1978). By replotting the data in Fig. 5 with additional data as $g_{\text{Na}}/(g_{\text{Na}}^{\text{max}} - g_{\text{Na}})$ vs. E_m (Fig. 6), the data points above \sim -30 mV deviate from a least-squares line fit to the points for voltages more negative than -35 mV. The straight line is expected from a gating model in which each channel exists in only one of two states, open or closed, with a constant dipole moment involved in the transition between states. The discrepancy between the data points and the line at more positive potentials is consistent with the prediction of a linear sequential activation scheme as depicted below (Almers, 1978).

$$
A \underset{k_{ba}}{\overset{k_{ab}}{\rightleftharpoons}} B \underset{k_{cb}}{\overset{k_{bc}}{\rightleftharpoons}} C \underset{k_{dc}}{\overset{k_{cd}}{\rightleftharpoons}} D^*
$$
 (Scheme 1)

A is the initial resting state, B and C are nonconducting intermediates, and D* is the conducting state. The fit at lower potentials reveals a limiting logarithmic potential sensitivity of 5.3 mV/e-fold g_{Na} which corresponds to a lower limit of approximately five equivalent electronic charges required to open a single sodium channel at potentials near the resting membrane potential.

Are Activation and Deactivation Separate Mechanisms?

Comparisons between the kinetics of sodium tail currents and "OFF gating currents" upon repolarization have led to conflicting observations. The Hodgkin-Huxley model suggests that OFF gating current (proportional to *dm/dt)* should decay with a time constant (τ_{off}) equal to τ_m , thus predicting a ratio of $\tau_{\text{off}}/\tau_{\text{Na}}$ of 3. In squid axons observed ratios have ranged between values only slightly greater than 1 (Bezanilla and Armstrong, 1975 a and 1975 b ; Armstrong and Gilly, 1979) and values close to 3.5 (Keynes and Rojas, 1976). The different ratios may be partially attributable to a dependence of τ_{off} on prepulse potential at a fixed pulse duration (Keynes and Rojas, 1976) and a dependence of the ratio on holding potential (Keynes and Rojas, 1976; Schauf et al., 1977). In any event, in the physiological voltage range (i.e., that encompassed by a normal action potential) the time-courses of gating-current and sodium-current deactivation in squid axons are quite similar, at variance with the Hodgkin-Huxley prediction.

To account for τ_{off}/τ_{Na} ratios close to 1, Bezanilla and Armstrong (1975 a and 1975 b) initially proposed that sodium channels might activate by a linear sequence of transitions through several closed states to the open conformation, but that channel closure (deactivation) proceeded by a direct route between the open state and the initial resting state.

$$
A \rightleftharpoons B \rightleftharpoons C \rightleftharpoons D^*
$$
\n
$$
\begin{array}{c|c}\n & k_{ad} \\
\hline\n & k_{da} \\
\hline\n\end{array}
$$
\n(Scheme)

If k_{ad} is made negligible at all membrane potentials while k_{da} is larger than k_{dc} at very negative voltages yet smaller than k_{cd} at more positive potentials, the scheme could qualitatively account for both sodium activation and deactivation as well as for making τ_{off} and τ_{Na} identical. Comparison of Schemes 1 and 2 reveals an important difference in their predictions of what happens during the activation process.

If g_{Na} turn-off occurs by a reverse sequence of transitions to the activation process (Scheme 1), then a brief period can exist during the deactivation of

 (2)

 g_{Na} when a closed intermediate state (e.g., C) is the most populated. During this period, early in the time-course of a sodium tail current, a return of the membrane potential to the original test-pulse level would result in a nearly exponential increase in g_{Na} inasmuch as most of the channels have only to make the transition $C \rightarrow D^*$. As the second depolarizing step is further delayed, states A and B become increasingly populated, and the conductance activates with a more sigmoid time-course. If deactivation proceeds to its steady state at the holding potential, another step depolarization would result in a g_{Na} time-course identical to the original activation (Fig. 7 A).

FIGURE 7. Discriminative model predictions for secondary sodium conductance activation. Computer simulations of I_{Na} response to a voltage step to 0 mV from a holding potential of -80 mV followed by a transient repolarization to -60 mV. The duration of the brief repolarization is progressively increased from 100 μ s to 1.8 ms. (A) Prediction of Scheme 1, where $k_{ab} = k_{bc} = k_{cd} = \alpha_m$ (Palti, 1971), and $k_{ba} = k_{cb} = k_{dc} = \beta_m/3$ (Palti, 1971). (B) Prediction of Scheme 2 with rate constants the same as above except $k_{ad} = 0.003 \text{ ms}^{-1}$, $k_{da} = \beta_m/3$, and $k_{\text{dc}} = 0.003 \text{ ms}^{-1}$.

In the alternative proposal (Scheme 2) g_{Na} deactivation does not lead to a significant change in the population of closed intermediate states. Rather, the initial resting state (A) is filled directly from the conducting state (D^*) . Thus, if a depolarizing pulse is reapplied during any stage of g_{Na} turn-off, the activation kinetics will always be sigmoid inasmuch as most of the channels are distributed between states A and D^* and further increases in the population of D^* must occur via the two intermediates in the sequence (Fig. 7 B).

I attempted experimental discrimination between these two schemes by applying twin depolarizations to perfused squid axons and varying the interval between steps. Typical results are shown in Fig. 8 A and B for two axons before pronase treatment. When the interval between pulses was very short (i.e., when only a small fraction of open channels had closed) the second activation of I_{Na} proceeded without significant delay. As the interval was progressively increased, the turn-on kinetics developed a sigmoid appearance similar to the prediction of Scheme 1. In all such determinations a secondary activation initiated early in the deactivation process proceeded approximately as a single exponential, regardless of test voltage. Comparable observations were made after pronase treatment.

An alternative protocol to demonstrate the same phenomenon involves a brief fixed-duration repolarization to different membrane potentials near the time of peak g_{Na} . Since the rate of deactivation is highly voltage dependent,

FIouRr 8. Secondary activation of sodium conductance in squid axons. Sodium current responses to the indicated voltage protocols for axons perfused externally with one-half normal Na concentration and internally with 250 mM Cs. A and B from axon 072577a; C is from axon 072577b. 5°C.

the second depolarization would occur during different stages of the turn-off process, depending on the speed of deactivation. For repolarizations to potentials increasingly more positive than the normal holding potential, more of the channels would be expected to populate state C of Scheme 1; thus, the subsequent g_{Na} turn-on would be less and less sigmoid. This pattern is verified in Fig. 8 C.

DISCUSSION

The goal of the present work is to provide kinetic and steady-state information about the activation of squid axon sodium channels from direct measurements of sodium currents that fail to inactivate and to compare this data with that previously obtained under the assumption that activation and inactivation are independent. Although the comparison yields some clear differences, they are, with only one exception, very small, suggesting that previous experimental determinations of activation parameters under pharmacological influence or some previous neural modeling attempts utilizing the Hodgkin-Huxley scheme for g_{Na} have not been in significant error, resulting from uncertainty about the voltage dependence and time dependence of activation.

The voltage dependence of g_{Na} measured from steady-state sodium currents agrees well with that determined from extrapolation of current-time curves with normal inactivation in the same axons. The limiting logarithmic voltage sensitivity of 5.3 mV/e-fold g_{Na} allows us to revise the lower limit on the number of equivalent charges that must be transferred across the membrane's electric field to open each sodium channel from six (Almers, 1978) to approximately five. It is interesting to note that Keynes and Rojas (1976) also found a shallower voltage dependence for g_{Na} (6.5 mV/e-fold g_{Na}) in Cs⁺-perfused axons from *Loligo forbesi* than had been originally observed by Hodgkin and Huxley (1952 b).

The voltage dependence of the rate of sodium activation after pronase or NBA treatment is reasonably close to that previously described for intact axons (Hodgkin and Huxley, 1952 b) and untreated $Cs⁺$ -perfused axons (Keynes and Rojas, 1976). Within the measurement resolution an exponential raised to the third power and delayed by 70 μ s or less is an adequate fit to the time-course of sodium conductance turn-on. This approximate fit should not be taken as evidence for a gating mechanism involving three homogenous two-state gating particles, a common interpretation of the Hodgkin-Huxley $m³$ parameter. The presence of the delay and its apparent voltage dependence suggests a more complicated scheme or a totally different approach and deserves further study. Furthermore, the sequential kinetic equivalent of $m³$ with forward rate constants in the ratio 3:2:1 is not a requirement, because a scheme with all forward rate constants set equal produces an approximate fit to sodium activation kinetics (Fig. 7; see also Bezanilla and Armstrong [1977]).

Kinetic schemes of g_{Na} in which inactivation is directly coupled to the conducting state (e.g., Bezanilla and Armstrong [1977] and Goldman and Hahin, [1978]) predict delayed inactivation kinetics as opposed to the purely exponential time-course of the h parameter. It might be expected that this difference should yield different kinetics for early g_{Na} activation in the presence of inactivation. Fig. 9 illustrates, however, that the predicted differences are actually quite small and also voltage dependent. The solid lines represent I_{Na} computed from the Hodgkin-Huxley model, with g_{Na} proportional to $m^3 h$. The dashed lines derive from a linear sequential scheme (equivalent to m^3) with the addition of a terminal inactivated state (E).

$$
A \rightleftharpoons B \rightleftharpoons C \rightleftharpoons D^* \overset{k_{de}}{\underset{k_{ed}}{\rightleftharpoons}} E \tag{Scheme 3}
$$

 g_{Na} is proportional to D^{*}, and the inactivation rate constants are made voltage independent and chosen to provide a good fit to the inactivation time-course of $m³$ h at -30 mV. Activation according to Scheme 3 is only slightly delayed relative to *m3h* and only over a particular voltage range. Thus, the presence or absence of coupling between activation and inactivation has a negligible influence on the voltage dependence of sodium activation kinetics determined in these experiments. Also, the time-course of inactivation is reasonably duplicated by Scheme 3, with voltage independent values of k_{de} and k_{ed} , as suggested previously (Bezanilla and Armstrong, 1977).

In pronase-treated axons, sodium tail currents (deactivation) fell approximately as single exponentials over the voltage range examined. A second slower component observed in untreated axons at potentials more positive than \sim -50 mV was removed by pronase perfusion, revealing the expected correlation with the time-course of inactivation (Schaufet al., 1977; Goldman and Hahin, 1978). Although the values of τ_{Na} obtained from fitting single exponentials to sodium tail currents are approximately three times larger than expected from the original determinations of β_m by Hodgkin and Huxley

FIGURE 9. Computed sodium currents for step depolarizations from -80 mV to the indicated voltages. Solid lines represent computations for $m³h$, assuming $E_{\text{Na}} = +55$ mV. Dashed lines were computed from Scheme 3 (time-course of D^{*}) with the following rate constants: $k_{ab} = 3\alpha_m$, $k_{bc} = 2\alpha_m$, $k_{cd} = \alpha_m$, $k_{dc} =$ β_m , $k_{cb} = 2/3\beta_m$, $k_{ba} = \beta_m/3$, $k_{de} = 1.5$ ms⁻¹, and $k_{ed} = 0.03$ ms⁻¹. Dashed lines were scaled to match the peak I_{Na} simulated with m^2h at each potential. Values for k_{de} and k_{ed} were chosen by trial and error to match visually the decay phase of I_{Na} from the Hodgkin-Huxley model at -30 mV.

 $(1952 a$ and $1952 b)$, they agree with previously published values from squid. Consequently, it would appear that a new formulation for the voltage dependence of β_m scaled down by a factor of approximately 3 would be appropriate. However, in view of the discontinuous τ_m vs. voltage distribution (Fig. 4 A) obtained when activation and deactivation measurements are combined, this change is not sufficient to bring the Hodgkin-Huxley scheme into agreement with the data. τ_m values derived from tail currents are clearly larger than those derived from I_{Na} turn-on at the same voltage. The controversial discrepancy between τ_{off} and $3\tau_{\text{Na}}$ in previous gating current measurements (Bezanilla and Armstrong 1975 a and 1975 b; Keynes and Rojas, 1976; Moore, 1978) prompted a search for alternatives to the Hodgkin-Huxley activation scheme. In this regard, it would appear that the discontinuity presently observed in the τ_m vs. voltage curve would render meaningless any attempt at correlating I_{Na} and I_{g} turn-off by factors of 3.

One possible interpretation of this finding in terms of Scheme 1 is that the corresponding rate constants for activation (or deactivation) are probably not in a ratio of 3:2:1 (or 1:2:3) as derived from m^3 . Sufficiently slow deactivation can be achieved by making the transition $D^* \rightarrow C$ rate limiting for deactivation while retaining reasonable kinetics of activation at the same potentials. At more positive potentials, as the experimental rate of activation becomes faster and the apparent rate of deactivation is slower, such adjustments are insufficient to explain the data. In view of the uncertainty of τ_{Na} values at potentials more positive than -30 mV, even after removal of inactivation, the validity and nature of this difference will have to await further experiments.

Recently sodium tail currents in voltage-clamped *Myxicola* axons have been reported to exhibit several exponential components (Schauf et al., 1977; Goldman and Hahin, 1978). These experiments were conducted on intact axons in which overlapping inactivation is expected to contribute to tailcurrent decay. Accordingly, Goldman and Hahin found a slow component of sodium tails with time constants approximating τ_h at relatively positive

FIGURE 10. Simulation of a slow component in sodium tail current by a coupled linear kinetic scheme for g_{Na} (Scheme 3). Rate constants were identical to those of Fig. 9 except k_{ed} , which was 0.03 ms⁻¹ for curve I and 3 ms⁻¹ for curve *II*

potentials. The slow component extended to more negative potentials, where it had little voltage dependence between -128 and -38 mV. Schauf et al. (1977) report a second slow component only for potentials more negative than \sim -50 mV, whereas at more positive potentials the current decays as a slow single exponential, a prediction just opposite to that of the Hodgkin-Huxley model. Again, the slow component is relatively independent of voltage. In a linear coupled scheme for g_{Na} (Scheme 3) a second slow exponential is also expected in sodium tail currents at negative voltages. This component arises from the coupling between states D^* and E rather than from a property of the activation sequence itself. The inactivated state (E) is loaded from the conducting state (D^*) during a depolarizing voltage step. Upon repolarization to negative potentials the inactivated state will depopulate through the conducting state, resulting in a current with a time constant similar to inactivation at that potential. When the rate constant for the transition $E \rightarrow$ D* is made large enough, tail currents similar to those reported by Schauf et al. (1977) can be simulated (Fig. 10). If the rate constants governing this transition are relatively voltage independent (at least at negative potentials),

as suggested by Bezanilla and Armstrong (1977), then the slow component of the tail current resulting from this step will also be unaffected by the potential of repolarization.

The magnitude (relative weighting factor) of the predicted slow component increases with increasing prepulse duration as the population of the inactivated state (E) increases. Similarly, Goldman and Hahin (1978) reported an increase in the relative contribution of the slow component of the tail current in *Myxicola* with increasing pulse length. They attribute this to an increased loading of the conducting state. Alternatively, the phenomenon could be related to the rate of inactivation. In this regard, it would be interesting to compare the time-course of increase in the slow I_{Na} tail component in *Myxicola* to that of activation and inactivation as has been done for the slow component of OFF-gating current in squid (Armstrong and Bezanilla, 1977). The analogy between the behavior of a linear coupled scheme for g_{Na} and the appearance of a slow exponential at large negative potentials in Na tail currents in *Myxicola* is not meant to imply that such a mechanism is actually operative in this case. Indeed, the time constants reported for the slow component are apparently much smaller than τ_h measured from reactivation protocols in this potential range (Schauf, 1976). Rather, the conclusion is that coupled sequential kinetic schemes are capable of generating similar behavior, and caution must be exercised in making inferences about the processes of sodium-channel activation and deactivation when inactivation is present.

In the present experiments a protocol involving relatively brief voltage perturbations applied during nonsteady-state conditions of g_{Na} was used to examine the kinetics of the transition from an intentionally populated intermediate to the conducting state $(C \rightarrow D^*$ of Scheme 1). The second depolarization applied during different stages of deactivation revealed that the secondary g_{Na} turn-on was nearly exponential. This suggests that at least part of the turn-off of g_{Na} occurs via the reverse of the sequence of activation and not by a separate pathway between conducting and initial resting states (Scheme 2). Such observations argue against cyclic models, such as have been proposed (Bezanilla and Armstrong, 1975 a and 1975 b; Keynes, 1975; Moore and Cox, 1976), in which deactivation occurs by a step(s) entirely separate from activation. If the quasi-exponential turn-on of I_{Na} during the second of two closely spaced depolarizations actually reflects a mechanism similar to Scheme 1, then its time constant will provide a reasonable estimate of the two rate constants between states C and D*. Better data at more potentials, however, are required before such estimates can be presented. One point is suggested by the available data. Shifting the records of secondary I_{Na} turn-on (0 mV) obtained at various states of deactivation along the time axis results in approximate superposition of the records. If during the early stages of deactivation the most populated closed state is the one adjacent to the conducting state and, thus, the principle source for open channels during subsequent activation, the last step $(C \rightarrow D^*)$ of the activation sequence is probably rate limiting. Were this transition significantly faster than the preceding steps of the sequence, the exponential turn-on would be measureably faster than the activation from the initial resting state $(A \rightarrow \rightarrow \rightarrow D^*)$.

Armstrong and Gilley (1979) have reached a similar conclusion from analysis of gating current records. It should be noted that the present observations were made over a limited voltage range and that the final step may not always be rate limiting. The use of similar protocols in examining g_{Na} and g_K will hopefully lead to better estimates for rate constants in kinetic schemes with several nonconducting states.

As noted above, several technical precautions were observed in the present experiments to minimize contributions to the kinetics of I_{Na} by errors involving series resistance, spatial voltage inhomogeneity, and electrode polarization. Despite these attempts to ensure that experimental records reflect the true behavior of g_{Na} , one further consideration precludes at this time a more quantitative analysis of the data in terms of the exact number of states in a linear kinetic scheme or rate constants governing their transitions. The ionic environment on both sides of the axon membrane has a profound influence upon the gating behavior of sodium channels. In addition to the well-known effects of external divalent cations upon the voltage sensitivity of g_{Na} (see Hille et al. [1975]), the commonly used changes in the internal monovalent cation composition can modify the kinetics of sodium channel gating (Schauf and Bullock, 1978; Oxford and Yeh, 1979). In view of the differences in solution compositions presently used by investigators examining voltage-dependent gating, a clearer picture of the contributions of ion-channel interactions to the observed kinetic behavior of ionic conductances is required to distinguish aspects of model schemes that reflect ion properties from those that represent inherent properties of the channel gating machinery. Such work is now in $progress.¹$

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