

Sodium channel opening as a precursor to inactivation

A route to the inactivated state

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Received April 21, 1988/Accepted in revised form September 21, 1988

Abstract. The time constant of the process producing the delay in Na inactivation development as determined by the two pulse method (τ_{delay}) was extracted and compared to that of the slowest Na activation process (τ_3) for the I_{Na} during the conditioning pulse of that same determination. τ_{delay} and two pulse inactivation (τ_c) values were computer generated using a non-linear least squares algorithm. τ_3 and single pulse inactivation (τ_h) values were independently generated for each determination also with the aid of the computer using the same non-linear least squares algorithm. In one determination at 2 mV, τ_c was 4.68 and τ_{delay} 0.494 ms while τ_h was 4.70 and τ_3 0.491 ms for a τ_c/τ_h of 0.996 and a $\tau_{\text{delay}}/\tau_3$ of 1.006. Mean $\tau_{\text{delay}}/\tau_3$ from five determinations in four axons, both Cs and K perfused, and spanning a potential range of -27 to 2 mV was 1.068. The precursor process to inactivation is channel opening. Some fraction of channels presumably inactivate via another route where prior channel opening is not required.

Key words: Sodium currents, inactivation, kinetics, channel gating

Introduction

A clear delay preceding the development of Na inactivation has been reported in well controlled experiments in the giant axons of *Myxicola* (Goldman and Schauf 1972; Goldman and Kenyon 1982) and crayfish (Bean 1981), and small delays are also evident in the data of Gillespie and Meves (1980) in squid. The delay cannot be attributed to distortion of the time course of inactivation determined with the two pulse method by the development of activation during the conditioning pulse, voltage clamp errors associated with the current magnitude, or contamination by K currents. For some fraction of the channels, inactivation develops only subsequent to a precursor process, i.e. for these channels prior to inactivation gate closing there must first be another process which enables closure. Other frac-

tions of the channel population may reach the inactivated state by a route that requires either no or a different immediately preceding process (Aldrich and Stevens 1983).

Some information is available as to the identity of the precursor process in *Myxicola*. In an extensive series of determinations Goldman and Kenyon (1982) found that the operationally defined inactivation delay (time at which the unconditioned peak current value intersected the fitted τ_c exponential in a two pulse inactivation determination) was about proportional to the time to peak Na conductance (g_{Na}) during the conditioning pulse. If the precursor process is channel opening then both the inactivation delay and the time to peak g_{Na} would be approximate measures of the activation rate, and proportionality between these measures would be expected. These results suggest that the potential dependency of the time constant of the delay process and that of the activation process are similar. In addition, in 10%–15% of their experiments, amounting to seven determinations, Goldman and Kenyon were able to extract the time constant of the delay process, τ_{delay} . The measured $\tau_{\text{delay}}(V)$ values agreed well with activation kinetics, $\tau_m(V)$, previously reported in *Myxicola* (Goldman and Schauf 1973), suggesting that the precursor process to inactivation is channel opening. However, it is not clear from these results whether inactivation delay and activation kinetics agree quite closely, or are only rather similar. I present here new determinations of inactivation delay and activation time constants obtained from the same axons under the same conditions. Kinetics for the two processes are essentially indistinguishable. A preliminary report of these findings has been presented (Goldman 1987).

Methods

Myxicola infundibulum were obtained from the Laboratory of the Marine Biological Association of the United

Kingdom (Citadel Hill, Plymouth, United Kingdom). Methods for preparing and electrically recording from the giant axons were as described by Binstock and Goldman (1969). Normal artificial sea water (ASW) had the following composition: 440 mM Na, 10 mM Ca, 50 mM Mg, 560 mM Cl, 5 mM *Tris* (tris[hydroxymethyl] aminomethane; Sigma Chemical, St. Louis, MO), pH 8.0 ± 0.1 . Temperature was $5 \pm 0.5^\circ\text{C}$. Potentials are reported as absolute membrane potential (inside minus outside) and have been corrected for liquid junction potentials as described by Ebert and Goldman (1975).

Internal perfusion was initiated with the KCl-axoplasm dispersal method of Goldman and Kenyon (1979). Standard (K) internal perfusate had the following composition: 410 mM K, 50 mM F, 360 mM glutamate, 1 mM HEPES (*N*-2-hydroxyethylpiperazine *N'*-2-ethane sulfonic acid; Calbiochem-Behring, San Diego, CA), 4 mM EGTA (ethyleneglycol-*bis*-(β -amino-ethyl ether) *N,N'*-tetra acetic acid; Sigma Chemical Co.), 145 mM sucrose, pH 7.30 ± 0.05 . Cs perfusate was identical except that K glutamate and KF were replaced equivalent per equivalent with Cs glutamate and CsF. All internal perfusates had an osmotic pressure within 5% of that of ASW as determined by osmometer (osmette A; Precision Systems, Sudbury, MA). Internal perfusates were freshly prepared every few days.

All voltage clamp observations were made using compensated feedback to reduce errors produced by the series resistance, R_s , as described by Goldman (1986). To reduce R_s errors further, observations were made in bathing media with the Na concentration reduced by substitution with *TRIS* to 1/2 of that in ASW. With the current densities encountered in these measurements, displacements in membrane potential produced by any residual, uncompensated R_s , will generally be well less than 1 mV.

Pulses sent to the voltage clamp were formed by a PDP 11/34 computer (Digital Equipment, Maynard, MA). Membrane currents were filtered at 100 kHz with a four pole Bessel filter (model 4302; ITHACO, Ithaca, NY) digitized at a 20 μs sampling interval with a 12-bit analog to digital converter (model ADC-EH 12 B 3; Dattel Systems, Canton, MA) and stored on floppy disks for later analysis. Na currents were extracted by repeating the entire voltage clamp protocol in 1 μM tetrodotoxin (TTX; Calbiochem-Behring) and subtracting the two sets of records with the aid of the PDP-11/34. All experiments were done in the presence of 2 mM 3,4-diaminopyridine (Aldrich Chemical, Milwaukee, WI) in the external bathing medium to suppress the K conductance. 3,4-diaminopyridine solutions were freshly prepared each day and the pH carefully checked.

Two pulse inactivation determinations were made as described by Goldman and Kenyon (1982), an included a 5 ms gap (step back to the holding potential) between conditioning and test pulses. This gap width was sufficient to allow Na tail currents to decay to zero well before the start of the test pulse (Goldman 1986). Each conditioning pulse, gap, test pulse sequence was both preceded and followed by an unconditioned test pulse. The means of the peak I_{Na} values from these bracketing unconditioned test pulses were used to normalize each conditioned determination. Holding potential was always near -100 mV. All axons were held at the holding potential for 2 min before the start of each voltage clamp run, and 15 s were allowed between each voltage clamp pulse to minimize any effects of slow inactivation (Rudy 1981).

Peak currents were read with the aid of the PDP 11/34 after digitally filtering as described by Goldman and Chandler (1986). Time constants as determined both with one and two pulse methods were also determined with the aid of the PDP 11/34 using pattern-search, a non-linear least squares minimization procedure (see Colquhoun 1971). As I have configured it, the pattern-search procedure will fit optionally from one to four exponentials to any selected region of a current record or any selected region of a two pulse inactivation determination, providing an objective determination of time constants. These are simultaneous multi-exponential fits, and so computing time for single pulse current records increases substantially with each additional exponential fitted.

Results

In all cases multi-exponential fits to two pulse inactivation time courses and those to the I_{Na} during the conditioning pulses were generated independently by the PDP 11/34.

Both the results of the two pulse determinations and the single pulse current time course, $I_{\text{Na}}(t)$, are described as a sum of exponentials. For a complete description of $I_{\text{Na}}(t)$ a minimum of three relaxations are needed: an inactivation time constant (τ_h) and at least two activation time constants as there is a delay in the rise of g_{Na} . There is some experimental evidence for yet an additional activation process in *Myxicola* (Goldman and Hahn 1978). A possible description for the entire $I_{\text{Na}}(t)$, then, is

$$I_{\text{Na}}(t) = -I_1 \exp(-t/\tau_1) + I_2 \exp(-t/\tau_2) - I_3 \exp(-t/\tau_3) + I_h \exp(-t/\tau_h) + I_{\text{Na}\infty} \quad (1)$$

In none of the two pulse determinations analyzed here was there more than one inactivation delay process detected, and these results have all been described as the sum of just two exponentials, with time constants

τ_C and τ_{delay} , plus a steady state term. Accordingly, for $I_{\text{Na}}(t)$ no attempt was made to accurately define more than two time constants, τ_h and τ_3 . Those governing the delay in the rise of g_{Na} are either partially or totally unresolved. Single pulse fits were never extended to times shorter than 100 μs into the voltage step, and in no case were fits attempted with more than three exponentials.

Figure 1 presents two Na current records from different axons spanning the range of potentials examined. The top trace was obtained from a Cs perfused axon at -27 mV. Superimposed on the current trace (thin line) is a three exponential fit constructed with τ_h and two activation time constants, τ_3 and τ_2 . The fitted curve and the current record agree quite closely. The rise in g_{Na} is determined nearly entirely by a single exponential process (τ_3). A two exponential fit to this same record now starting at 300 μs into the voltage step produced a τ_3 only 1% faster and an identical τ_h . The lower trace is a current record at 2 mV also from a Cs perfused axon and again with a superimposed three exponential fit. A two exponential fit starting at 300 μs into the voltage step yielded a τ_3 3% faster and again an identical τ_h . The faster activation time constants (τ_2) in the three exponential fits can only be approximate values. These fits were started at 100 μs into the step. The accuracy of the τ_2 values depends both on their relative magnitude and on the relative magnitudes of the coefficients on these exponential terms as well as the relative magnitudes and relative magnitudes of the coefficients on the exponentials of any additional activation relaxations, which are totally unresolved in these computations. The point of the three exponential fits is to demonstrate that τ_3 can be determined reliably without accurately resolving the activation delay process(es).

The two pulse inactivation time course for the same determination as the lower trace of Fig. 1 is shown in Fig. 2. For this experiment both the test pulse and the conditioning pulse were at 2 mV. Open circles indicate peak values of I_{Na} during the test pulse, normalized to that with no conditioning pulse, as a function of conditioning pulse duration. The solid curve is a two exponential fit to these data, and the dotted curve indicates the extrapolation of the slower fitted τ_C exponential to zero time. The inset presents the first two ms of this determination on an expanded time scale. The delay in inactivation development is quite prominent. Following very brief conditioning pulses I_{Na} during the test pulse actually increases somewhat, indicating that states preceding the conducting state have not all returned to their resting occupancies by the end of the gap. In fact, such depolarizing conditioning potentials do shift the $I_{\text{Na}}(t)$ to the left along the time axis (Hahin and Goldman 1978). A three exponential fit to these same data did not

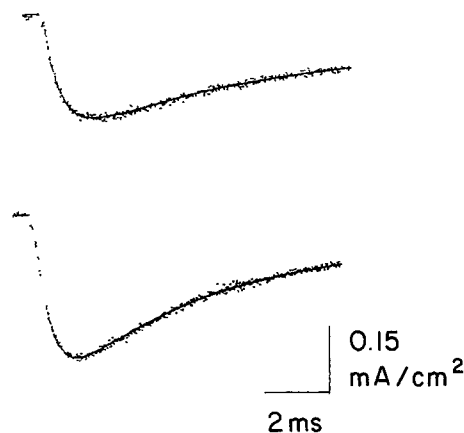


Fig. 1. Na current records from two different Cs perfused axons. The zero current level is indicated by the horizontal line at the left in each case. *Top:* Trace at a step to -27 mV from a holding potential of -102 mV. Superimposed thin line indicates a three exponential fit to the record using a τ_3 of 0.581 ms and a τ_h of 6.70 ms. Axon 4MP25. *Bottom:* Trace at a step to 2 mV from a holding potential of -103 mV. Superimposed three exponential fit computed with a τ_3 of 0.491 and a τ_h of 4.70 ms. Axon 4MP16. Inactivation was well described by single exponential in both cases. Measured current at 30 ms for the record at 2 mV was 0.068 mA/cm² while the computed value was 0.071 mA/cm². For the record at -27 mV the corresponding values at 30 ms were 0.049 and 0.055 mA/cm². Scale: 0.15 mA/cm², 2 ms

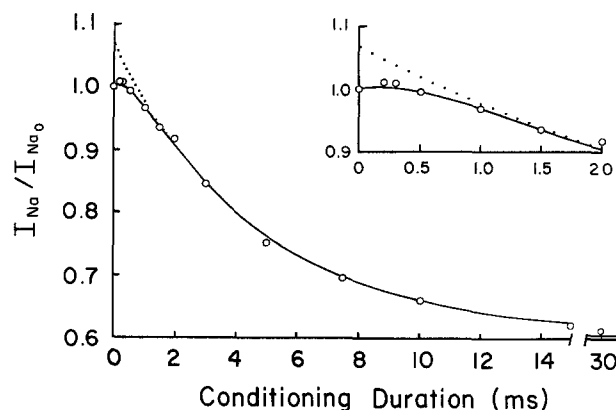


Fig. 2. Two pulse inactivation time course for the same determination as that for the lower trace of Fig. 1. Points indicate peak I_{Na} during a 2 mV test pulse, relative to that with no conditioning pulse, as a function of the duration of a 2 mV conditioning pulse. There was always a 5 ms step back to the holding potential between conditioning and test pulses. *Solid curve* is a two exponential fit computed with a τ_{delay} of 0.494 and a τ_C of 4.68 ms. *Dotted line* indicates the extrapolation of the τ_C relaxation to zero time. *Inset:* First 2 ms of the determination on an expanded time scale

produce any indication of a process that was itself a precursor to this delay process.

For the two exponential fits of Fig. 2 τ_C and τ_{delay} are 4.68 and 0.494 ms respectively, while for the time course of the I_{Na} during the conditioning pulse of this same experiment the three exponential fit illustrated in

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