

# Pharmakodynamic Analysis of Electrophysiological Data for Voltage-Gated Sodium Channels

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### 0.1 Drug Binding to Inactivated States

Determination of the dissociation constant from a *drug-bound* to a *slow inactivated* state, as sketched in fig. 1 panel B and C. *Model B* is difficult to identify. In *model C*, drug-bound states (fast *ID* and slow *SD*) are lumped into a single state *D*. In this model the apparent dissociation constant from  $D \leftrightarrow S$  is identifiable via a concentration-response relationship as derived below.

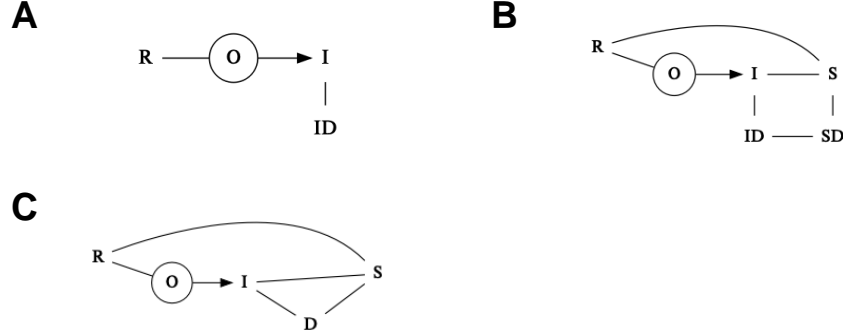


Figure 1: State models for drug-binding to Na<sub>V</sub> channels. A) Channel with only one fast inactivated state. B) Channel with a fast and slow inactivated state.

Only the fraction of channels that are in the *fast inactivated state* after the inactivation pulse are recovered into an activatable *resting state* *R* by the recovery pulse (20 ms duration). Subsequently only these mediate a current upon the test pulse. According to the model in fig. 1C, the following relationships for the amplitudes of control ( $I_C$ ) and drug-bound channels ( $I_D$ ) during the subsequent test pulse are proposed

$${}^0I_C = {}^0I_D = ([R] + [S] + [SD_n]) \cdot \bar{i}_{Na} \quad (1a)$$

$${}^\infty I_C = ([R] + [SD_n]) \cdot \bar{i}_{Na} \quad (1b)$$

$${}^\infty I_D = [R] \cdot \bar{i}_{Na} \quad (1c)$$

with the fractional current  $\bar{i}_{Na}$

$$\bar{i}_{Na} = \hat{g}_{Na} \cdot (V_P - E_{Na}) \quad (2)$$

Here,  ${}^0I_C$  and  ${}^0I_D$  represent control currents and currents for cells exposed to drug, respectively, at a inactivation impulse duration of  $t = 0s$ , or  ${}^\infty I_C$  and  ${}^\infty I_D$  after very long inactivation duration ( $t \rightarrow \infty s$ , i.e. steady-state). Symbols in square brackets indicate the “concentration” of channels in corresponding states after this infinitely long inactivation pulse. So,  $[R] + [S] + [SD_n]$  represents the total amount of channels. The dissociation constant for drug binding to channels in the *slow inactivated* conformation is described by



Therefore it holds that

$$K_S = \frac{[S] \cdot [D]^n}{[SD_n]} \quad (4)$$

According to Hill and Langmuir  $K_S$  determines the ratio of *drug-occupied* to *total amount of channels* in state  $S$  or alternatively *drug-free* to *total amount of channels*

$$\begin{aligned} y &= \frac{\text{drug-free channels in state } S}{\text{total number of channels in state } S} \\ y &= \frac{[S]}{[S] + [SD_n]} \end{aligned} \quad (5)$$

Rearranging eq. 4, substitution into eq. 5 and simplification gives the following Hill equation

$$y = \frac{K_S}{K_S + [D]^n} \quad (6)$$

The ratio  $y$  can be expressed from measured currents in eqq. 1 as follows

$$y = \frac{{}^0I_C - {}^\infty I_C}{{}^0I_D - {}^\infty I_D} \quad (7a)$$

$$= \frac{[R] + [S] + [SD_n] - [R] - [SD_n]}{[R] + [S] + [SD_n] - [R]} \quad (7b)$$

$$= \frac{[S]}{[S] + [SD_n]} \quad (7c)$$

From eq. 6 and eq. 7a

$$\frac{{}^0I_C - {}^\infty I_C}{{}^0I_D - {}^\infty I_D} = \frac{K_{bS}}{K_{bS} + [D]^n} \quad (8)$$

Therefore,  $K_{bS}$  could be estimated by fitting eq. 8 to currents measured according to the *block-entry protocol*.  $K_{bS}$  represents the *dissociation constant* as determined from a functional assay, which is equal to  $K_S$  if the expression in 7b were correct.