

Figure S5. The coupled motion between adjacent protomers indicated by the structural comparison of Na_vRh and Na_vAb. **(A)** There would be a structural clash between VSDs and the pore domain of Na_vAb if its VSDs were to adopt the same conformation and position as in Na_vRh. Structures of Na_vRh and Na_vAb are superimposed relative to the pore domains. The pore helix bundle of one subunit from Na_vAb is shown and colored silver; the same segments of Na_vRh as in Figure S4A are shown in a side view. The inset illustrates the steric clash between Na_vRh -S4 (blue) and Na_vAb-S5 of the adjacent protomer (silver cartoon with orange side chains) when the pore domains are superimposed. **(B)** Concerted shift of the S4-5 helices and S5 segments between adjacent protomers. Note that the N-terminal tip of the S4-5 helix is shifted toward the pore by more than 6 Å (Figure 1D); yet there is little change in

the distance between the base of the S5 segment in one protomer (yellow) and the N-terminal tip of the S4-5 helix in the adjacent protomer (blue) in the two structures. Refer to *Supplementary Movies S5-6* for animations.