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Structural mechanisms of selectivity and gating in anion channelrhodopsins

Hideaki E. Kato^{1,2,13*}, Yoon Seok Kim^{3,4,5,13}, Joseph M. Paggi^{6,7}, Kathryn E. Evans^{3,4,5}, William E. Allen^{3,4,5}, Claire Richardson⁶, Keiichi Inoue^{2,10,11}, Shota Ito¹⁰, Charu Ramakrishnan^{3,4,5}, Lief E. Fenno^{3,4,5}, Keitaro Yamashita¹², Daniel Hilger¹, Soo Yeun Lee^{3,4,5}, Andre Berndt^{3,4,5}, Kang Shen^{8,9}, Hideki Kandori^{10,11}, Ron O. Dror^{6,7}, Brian K. Kobilka¹ & Karl Deisseroth^{3,4,5*}

¹Department of Molecular and Cellular Physiology, Stanford University School of Medicine, Stanford, CA, USA. ²PRESTO, Japan Science and Technology Agency, Kawaguchi, Japan. ³Department of Bioengineering, Stanford University, Stanford, CA, USA. ⁴Department of Psychiatry and Behavioral Sciences, Stanford University, Stanford, CA, USA. ⁵Howard Hughes Medical Institute, Stanford University, Stanford, CA, USA. ⁶Department of Computer Science, Stanford University, Stanford, CA, USA. ⁷Institute for Computational and Mathematical Engineering, Stanford University, Stanford, CA, USA. ⁸Department of Biology, Stanford University, Stanford, CA, USA. ⁹Howard Hughes Medical Institute, Stanford University, Stanford, CA, USA. ¹⁰Department of Life Science and Applied Chemistry, Nagoya Institute of Technology, Nagoya, Japan. ¹¹OptoBioTechnology Research Center, Nagoya Institute of Technology, Nagoya, Japan. ¹²RIKEN SPring-8 Center, Hyogo, Japan. ¹³These authors contributed equally: Hideaki E. Kato; Yoon Seok Kim. *e-mail: hekato@stanford.edu; deissero@stanford.edu

Supplementary Discussion

Distinct photocycles in iC⁺⁺ and GtACR1

Both iC⁺⁺ and GtACR1 work as light-gated anion channels, but their photocycles are remarkably different. While GtACR1 has 5 intermediate states (K, L, M, N, and O), iC⁺⁺ has only three photo-distinguishable intermediate states (K, M and O), and the lifetime of the K intermediate is unusually long. The K-intermediate state is generally characterized by the 13-*cis* retinal twisted around the C14-C15 bond. The twisted retinal becomes planar during the transition from K- to L- intermediate state, and the Schiff base proton is transferred from the Schiff base nitrogen to the proton acceptor during the transition from L- to M- intermediate state. Since iC⁺⁺ has an unusually long K-intermediate state and does not exhibit the L-intermediate, some of the mutations introduced to engineer iC⁺⁺ (i.e. T98S, E129Q, E162S, and/or N297Q) may change the local environment around the Schiff base and inhibit the efficient relaxation of the twist around the C14-C15 bond of 13-*cis* retinal. The proton would be directly transferred from the Schiff base nitrogen of twisted retinal to the proton acceptor (probably Asp292), and thereby iC⁺⁺ could skip the L-intermediate state.

Ion selectivity in ACRs

This study supports a framework in which both nACR and dACR follow the overall surface-electrostatic model for pore selectivity, in which the contribution of each individual amino acid position to anion selectivity can vary between ACRs. In iC⁺⁺, amino acids at the CCS (Q129 and Q297) position are particularly important for ion selectivity, and as shown in Fig. 3e, the Q129E/Q297E double mutation depolarizes V_{rev} by ~40 mV. An additional 8 mutations (T98S, E122N, E140S, V156R, E162S, V281R, T285N, and E312S) mainly introduced inside the ion-conducting pathway work cooperatively, creating an electropositive surface suitable for anion selectivity, and thus further hyperpolarize V_{rev} by 20 mV. This pattern also holds for another dACR (iChloC, in which the E-to-R mutation to E90 (Glu-129 in iC⁺⁺) decreases V_{rev} from ~+10 to ~-40 mV⁴). An additional four mutations, E83Q, E101S, D156N, and T159C (E122Q, E140S, D195N, and T198C in iC⁺⁺ numbering, respectively), of which the first two are introduced inside the ion-conducting pathway, further hyperpolarize V_{rev} to ~-65 mV⁵. The mutational tests for positions inside the nACR ion-conducting pathway tend to show more

variable contribution to anion selectivity; while Gln-46, another residue comprising the CCS in *GtACR1*, does contribute to anion selectivity (lowering V_{rev} by ~ 15 mV), the E-to-Q mutation of Glu-68 (Glu-129 in *iC++*) does not significantly change V_{rev} (Fig. 3e; this mutation however would be predicted to give rise to a more mild electrostatic effect than the E-to-R mutation in *iChloc* noted above), and positively-charged residues positioned at the nACR ion-conducting pathway vestibules appear important for tuning the electropositive surface suitable for anion conduction. Therefore, although both dACR and nACR follow the pore surface-electrostatic model, thus far the tested residues contributing most to anion selectivity in dACRs have been found positioned at the CCS or inside the ion-conducting pathway, and the tested residues found to contribute more to anion selectivity in nACRs have been positioned near the vestibules of the ion-conducting pathway. Structures for additional ChRs will be informative regarding possible evolutionary contributions to this pattern and to chloride/cation selectivity in general, and already our demonstrated ability to interconvert ChR ion selectivity (CCR \rightarrow ACR and now ACR \rightarrow CCR), guided by our crystal structures and structure-informed electrostatic model, may help in the screening of ChR sequences for informative new variants in nature.

Relationship between distribution of charged residues and photocurrent amplitude

Both *iC++* and *GtACR1* have several positively charged residues that govern exclusive anion selectivity, but their distributions are completely different, consistent with the overall surface-electrostatic model and not with a model involving singular high-affinity binding sites. These different distributions (in the setting of similar ion selectivity) not only support the selectivity model but also may be relevant to explaining differences in photocurrent amplitude, a distinct ChR property in which *iC++* and *GtACR1* differ markedly. Except for the Schiff base lysine, *iC++* has four positively charged residues along the pore (Lys-132, Arg-156, Arg-159, and Arg-281), but only one of these is conserved in *GtACR1*, and Lys-132, Arg-156, and Arg-281 are replaced by Thr-71, Pro-91, and Glu-223, respectively. Instead, *GtACR1* has 12 positively charged residues at and near the pore vestibules, which create an electropositive surface suitable for anion selectivity. Because of its lysine and arginine residues inside the ion-conducting pathway, the surface of pore in *iC++* is predicted to be more “sticky” for substrate anions. As proposed for potassium and calcium channels previously^{48,49}, ions tightly binding to the inner surface of the pore will exhibit lower mobility. Therefore the lower conductance of *iC++* relative

to *GtACR1* may be understood from the structures and a sticky-pore model, and it will be of interest to see if this pattern can be extended to CCRs.