

1    **Optical electrophysiology for probing function and pharmacology of voltage-gated ion**  
2    **channels**

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10

11    **Abstract**

12    Voltage-gated ion channels mediate electrical dynamics in excitable tissues and are an important class of  
13    drug targets. Channels can gate in sub-millisecond timescales, show complex manifolds of  
14    conformational states, and often show state-dependent pharmacology. Mechanistic studies of ion  
15    channels typically involve sophisticated voltage-clamp protocols applied through manual or automated  
16    electrophysiology. Here, we develop all-optical electrophysiology techniques to study activity-  
17    dependent modulation of ion channels, in a format compatible with high-throughput screening. Using  
18    optical electrophysiology, we recapitulate many voltage-clamp protocols and apply to Na<sub>v</sub>1.7, a channel  
19    implicated in pain. Optical measurements reveal that a sustained depolarization strongly potentiates  
20    the inhibitory effect of PF-04856264, a Na<sub>v</sub>1.7-specific blocker. In a pilot screen, we stratify a library of  
21    320 FDA-approved compounds by binding mechanism and kinetics, and find close concordance with  
22    patch clamp measurements. Optical electrophysiology provides a favorable tradeoff between  
23    throughput and information content for studies of Na<sub>v</sub> channels, and possibly other voltage-gated  
24    channels.

25

26 **Introduction**

27 To gain detailed mechanistic insight into ion channel function and pharmacology, one often studies  
28 single channels, heterologously expressed, under voltage-clamp protocols.<sup>1-3</sup> Carefully designed  
29 sequences of voltage steps prepare channels in select conformational states.<sup>4,5</sup> Distinct sub-states often  
30 have widely divergent affinities and kinetics of interaction with drugs. Knowledge of this state-  
31 dependent behavior is critical in developing models of channel function and in predicting how drugs will  
32 function *in vivo*. State-dependent dynamical measurements are typically the domain of manual or  
33 automated patch clamp. Functionally equivalent optical assays would open the prospect of high  
34 throughput screens with sophisticated state-dependent selection criteria; and might enable  
35 measurements in cell types or environments (e.g. in a tissue or whole animal) that are challenging to  
36 access with conventional methods.

37 Tools for optical electrophysiology—simultaneous optical perturbation and optical readout of  
38 membrane potential—have been making inroads into neuroscience<sup>6,7</sup>, with a primary emphasis on  
39 spatially resolved measurements *in vivo*<sup>8,9</sup> or in complex cell cultures<sup>10</sup>. Voltage-sensitive dyes (VSDs)  
40 have been applied in a wide range of physiological assays *in vitro*<sup>11</sup> and *in vivo*,<sup>12</sup> but existing red-shifted  
41 VSDs are still excited by the wavelengths used to stimulate optogenetic actuators, leading to optical  
42 crosstalk<sup>13,14</sup>, or are not commercially available<sup>15</sup>. A combination of a blue-shifted channelrhodopsin  
43 (CheRiff) and a red-shifted voltage indicator protein (QuasAr2) recently achieved spectrally orthogonal  
44 optical stimulation and readout.<sup>10</sup>

45 Optical electrophysiology measurements are typically semi-quantitative, at best, while ion channel  
46 assays require accurate perturbations to voltage and measurements of current. The optical techniques  
47 face several challenges: expression levels of optogenetic actuators and reporters vary from cell to cell;  
48 channelrhodopsins function as a conductance, not a voltage clamp; and fluorescence can only be used  
49 to measure membrane voltage, not current.<sup>12,16</sup> Thus it is not obvious whether one can apply optical  
50 electrophysiology as a functional surrogate for standard voltage-clamp protocols.

51 Here we address this challenge by developing optical assays of the state-dependent electrophysiology  
52 and pharmacology of voltage-gated sodium ( $Na_v$ ) channels. We begin with electrophysiologically inert  
53 HEK cells. We then stably express four transgenic constructs: an inward rectifier potassium channel and  
54 a voltage-gated sodium channel imbue the HEK cells with the ability to produce regenerative electrical  
55 spikes.<sup>17-19</sup> A channelrhodopsin variant, CheRiff, triggers these spikes upon exposure to flashes of blue

56 light. An Archaerhodopsin variant, QuasAr2, enables fluorescent readout of membrane voltage via red  
57 excitation and near-infrared fluorescence. The QuasAr2 reporter has a ~1 ms response time and a linear  
58 response between -100 to +100 mV, providing a direct correlation of fluorescence and voltage<sup>10</sup>.

59 Brief flashes of blue light trigger sodium channel-mediated action potentials, which manifest as flashes  
60 of near infrared fluorescence. Steady state illumination with blue light induces steady state changes in  
61 voltage, and thereby changes in the distribution of  $\text{Na}_v$  channels among substates. We develop stimulus  
62 and data analysis protocols that are robust to sources of cellular variation, and we compare our results  
63 to measurements by manual patch clamp. While whole-cell voltage clamp remains the gold standard for  
64 absolute accuracy, optical electrophysiology provides a favorable tradeoff between accuracy and  
65 throughput.

66  $\text{Na}_v$  channels mediate the rising phase of the action potential and play significant physiological functions  
67 in excitable tissues. There are nine subtypes of  $\text{Na}_v$  channels in the human genome.  $\text{Na}_v$  channel  
68 dysfunction has been implicated in many human diseases. For example, loss-of-function mutations in  
69  $\text{Na}_v1.1$  can cause Dravet syndrome and in  $\text{Na}_v1.5$  can cause Brugada syndrome<sup>20</sup>. The  $\text{Na}_v1.7$  sodium  
70 channel plays an important role in mediating pain sensation. Homozygous loss of function leads to  
71 congenital insensitivity to pain<sup>21</sup>, gain of function mutations lead to spontaneous severe pain, called  
72 erythermalgia<sup>22</sup>, and nucleotide polymorphisms modify sensitivity to pain in the general population<sup>23</sup>.  
73 While recent results have suggested that the connection of  $\text{Na}_v1.7$  to pain may involve other signaling  
74 pathways as well<sup>24</sup>, there remains strong interest in finding selective blockers of this channel. Recent  
75 structural work has mapped an isoform-specific binding site for  $\text{Na}_v1.7$ -specific blockers<sup>25</sup>, opening the  
76 possibility to develop new blockers via structure-guided design. Here we apply the Optopatch spiking  
77 HEK cell platform to study  $\text{Na}_v1.7$ , and we demonstrate its applicability to  $\text{Na}_v1.5$  also.

78

## 79 **Results**

### 80 **Construction and characterization of $\text{Na}_v1.7$ Optopatch Spiking ( $\text{Na}_v1.7$ -OS) HEK cells**

81 We engineered a monoclonal HEK293 cell line stably expressing human  $\text{Na}_v1.7$  and the Optopatch  
82 constructs (see Methods). Both QuasAr2-mOrange2 and CheRiff-eGFP showed good membrane  
83 trafficking (Figure 1B). We used manual whole-cell patch clamp measurements to characterize the  
84 performance of each component. Under whole-cell voltage clamp ( $V_m = -60$  mV) CheRiff was activated

85 by 488 nm light with an EPD50 (effective power density for 50% activation) of 20 mW/cm<sup>2</sup> and a  
86 saturating steady-state photocurrent density of  $13.0 \pm 1.2$  pA/pF (mean  $\pm$  s.e.m.,  $n = 5$  cells, Figures 2A,  
87 2B). As with channelrhodopsin 2, CheRiff showed inward rectification<sup>26</sup> with a reversal potential of  
88 +4 mV, consistent with non-selective cation conductivity. Under voltage steps from a holding potential  
89 of -100 mV, Na<sub>v</sub>1.7 mediated robust inward currents with fast activation and inactivation kinetics within  
90 10 ms and a peak current density of  $-61.4 \pm 13.6$  pA/pF at -20 mV (mean  $\pm$  SD,  $n = 11$  cells, Figure 2C).

91 We found that stable expression of K<sub>ir</sub>2.1 interfered with cell growth, so we expressed this channel via  
92 transient transfection. We call the quadruply expressing cells Na<sub>v</sub>1.7 Optopatch Spiking (Na<sub>v</sub>1.7-OS) HEK  
93 cells (Figure 1A). In a bath solution containing 2 mM K<sup>+</sup>, Na<sub>v</sub>1.7-OS HEK cells had a resting potential of -  
94  $97.2 \pm 2.2$  mV (mean  $\pm$  s.e.m.,  $n = 7$  cell clusters), sufficient to prime most of the Na<sub>v</sub>1.7 channels for  
95 activation. Upon voltage steps from -100 mV, K<sub>ir</sub>2.1 showed inward rectifying behavior (Figure 2D).

96 Using manual patch clamp, we quantified the effect of CheRiff activation on membrane voltage. Brief  
97 optical stimuli (20 ms, 50 mW/cm<sup>2</sup>) reliably triggered single spikes (Figure 2-figure supplement 1A), with  
98 a firing threshold of  $-48.0 \pm 1.2$  mV, peak depolarization of  $+30.1 \pm 3.7$  mV, and spike width at half-  
99 maximum repolarization (APD<sub>50</sub>) of  $33.5 \pm 3.3$  ms (mean  $\pm$  s.e.m.,  $n = 5$  cells). Under steady-state blue  
100 illumination, cells asymptotically approached a steady state depolarization that increased monotonically  
101 with stimulus intensity (Figure 2-figure supplement 1B, 1C), reaching an asymptotic value of  $-25.8 \pm$   
102  $6.2$  mV (mean  $\pm$  SD,  $n = 4$  cells) under intense illumination.

103 We then performed simultaneous recordings of membrane voltage and QuasAr2 fluorescence under  
104 optical CheRiff stimulation (Figure 2E). Cells were illuminated with continuous red light (640 nm,  
105 400 W/cm<sup>2</sup>) to excite QuasAr2 fluorescence. Pulses of blue light (500 ms on, 1.5 s off, stepwise  
106 increasing intensity from 1.1 to 26.0 mW/cm<sup>2</sup>) were applied to activate CheRiff. Stimuli of intensity  
107 15 mW/cm<sup>2</sup> or greater induced action potentials. The fluorescence traces faithfully reproduced both  
108 the action potential waveforms and the subthreshold depolarizations.

109 Optopatch measurements report membrane voltage, while patch clamp measurements typically control  
110 voltage and measure current. We thus used manual patch clamp measurements to determine the  
111 relation between voltage spike height measured in current clamp, and peak Na<sub>v</sub>1.7 current measured in  
112 voltage clamp. We used the state-dependent binding of amitriptyline to induce varying degrees of  
113 channel block, and then either applied a current pulse and measured the voltage response, or a voltage  
114 step and measured the current response (Methods, Figure 2-figure supplement 2A, 2B). Figure 2-figure

115 supplement 2C shows that the voltage spike amplitude was smoothly and monotonically related to the  
116  $\text{Na}_v1.7$  current. Thus optical measurements of spike height are a quantitative probe of  $\text{Na}_v$  current-  
117 carrying capacity. High-magnification fluorescence measurements showed that each individual cell gave  
118 a graded spike amplitude as a function of  $\text{Na}_v1.7$  capacity (Figure 2-figure supplement 3A), with an 8%  
119 standard deviation in spike height at 50% channel block ( $n = 5$  cells, Figure 2-figure supplement 3B).

120 Finally, we tested for photothermal or photochemical damage from the intense red illumination used for  
121 QuasAr2 imaging. In a 35 mm dish containing 2 mL of imaging buffer, continuous red illumination at  
122  $400 \text{ W/cm}^2$  (315 mW total power) for 10 min induced a temperature rise  $< 0.3 \text{ }^\circ\text{C}$  (Figure 2-figure  
123 supplement 4A). In a single well of a 384-well plate, containing 36  $\mu\text{L}$  of imaging buffer, the  
124 temperature rise was  $2.5 \pm 0.4 \text{ }^\circ\text{C}$  (mean  $\pm$  s.e.m.,  $n = 4$  replicates) in 35 s and  $9.8 \pm 0.6 \text{ }^\circ\text{C}$  (mean  $\pm$  s.e.m.,  
125  $n = 4$  replicates) in 10 min. While a  $10 \text{ }^\circ\text{C}$  rise is within the physiological range for measurements  
126 starting at room temperature ( $23 \text{ }^\circ\text{C}$ ), we kept all measurement protocols shorter than 35 s to avoid  
127 possibility of thermal artifacts.

128 Under these same conditions ( $400 \text{ W/cm}^2$ , 384 well plate, 36  $\mu\text{L}$  buffer) the cells continued to produce  
129 optically evoked spikes for 10 min, with little change in spike waveform (Figure 2-figure supplement 4B).  
130 The baseline QuasAr2 fluorescence dropped by 12% in this interval (Figure 2-figure supplement 4B) and  
131 the spike amplitude dropped from  $3.5 \pm 0.04\% \Delta F/F$  to  $2.3 \pm 0.06\% \Delta F/F$  (mean  $\pm$  SD,  $n = 6$  spikes, Figure  
132 2-figure supplement 4C). The signal-to-noise ratio (SNR, spike height/baseline noise) dropped from  $107$   
133  $\pm 2$  to  $73 \pm 2.6$  (mean  $\pm$  SD,  $n = 6$  spikes, Figure 2-figure supplement 4C). We explored the dependence  
134 of SNR on red illumination intensity (Figure 2-figure supplement 4D, 4E). At  $400 \text{ W/cm}^2$  the SNR was  
135  $99.8 \pm 3.3$  (mean  $\pm$  SD,  $n = 8$  spikes) at a 100 Hz frame rate. At  $6.3 \text{ W/cm}^2$  (5 mW) the SNR was  $9.5 \pm 1.1$   
136 (mean  $\pm$  SD,  $n = 8$  spikes) at the same frame rate. Thus spiking HEK cells can be imaged under a wide  
137 range of conditions, without photochemical or photothermal toxicity.

138

### 139 **Optically probing $\text{Na}_v1.7$ pharmacology with $\text{Na}_v1.7$ -OS HEK cells**

140 Sodium channel blockers are expected to change the firing properties of  $\text{Na}_v1.7$ -OS HEK cells. Most  
141 clinically used sodium channel blockers (e.g. lidocaine) show use-dependent or state-dependent action.  
142 We stimulated  $\text{Na}_v1.7$ -OS HEK cells with bursts of blue light (20 ms duration, six pulses) at 2, 4, and 8 Hz.  
143 The optically evoked action potentials were recorded by QuasAr2 fluorescence, averaging over  $\sim 150$

144 cells. In untreated cells, each stimulus evoked an action potential. After addition of lidocaine (200  $\mu$ M),  
145 cells continued to spike faithfully at 2 Hz. At 4 Hz and 8 Hz, cells spiked in response to the first stimulus,  
146 but failed for subsequent stimuli, a hallmark of activity-dependent  $\text{Na}_v$  block (Figure 3 A).

147 Sodium channel blockers often show complex state-dependent binding affinities and kinetics. Voltage  
148 clamp protocols have been developed to prepare specific states to probe these mechanisms. Most  
149 voltage-clamp protocols comprise a prepulse, an optional recovery interval, and a test pulse. The  
150 voltage during each interval can be selected to populate different states. We sought to recreate these  
151 protocols by programming the duration and intensity of the blue light pulses.

152 First we varied the duration of the 488 nm optical prepulse from 20 ms to 500 ms, to probe state-  
153 dependent binding. The intensity was 50 mW/cm<sup>2</sup>, which correspond to a depolarization to  $\sim$ 30 mV.  
154 The cells were then given 200 ms recovery with no optical stimulus. The recovery interval was selected  
155 to allow drug-unbound channels to reprime. The test pulse (30 ms, 50 mW/cm<sup>2</sup>) probed residual  
156 excitability. We used the amplitude of the fluorescence spike during the test pulse as a proxy for the  
157 degree of remaining  $\text{Na}_v1.7$  current. Simultaneous fluorescence and manual patch clamp  
158 measurements showed close correspondence of the optical and electrical signals (Figure 3B). After  
159 adjusting the fluorescence data for scale and offset relative to the voltage recording (as in Ref.<sup>27</sup>), the  
160 residual variations in fluorescence had an amplitude equivalent to 2.8 mV in a 200 Hz bandwidth.

161 We then applied the measurements using optical stimulation and recording alone, without patch clamp.  
162 Amitriptyline, a tricyclic antidepressant, showed strong state-dependent binding with degree of channel  
163 block dependent on prepulse duration (Figure 3-figure supplement 1A). The IC50 values of amitriptyline  
164 varied from  $11.7 \pm 1.6 \mu\text{M}$  at 20 ms prepulse to  $1.6 \pm 0.1 \mu\text{M}$  at 500 ms prepulse (standard error of fit to  
165 Hill equation,  $n = 3$ -5 wells per data-point, Figure 3C). This state-dependent block is consistent with  
166 previous patch clamp results<sup>28</sup>. In contrast, TTX showed very modest state dependence (Figure 3-figure  
167 supplement 1B). The optically recorded IC50 values of TTX were  $126 \pm 13 \text{ nM}$  at 20 ms prepulse and  $62 \pm 5.8 \text{ nM}$  at 500 ms prepulse (standard error of fit to Hill equation,  $n = 4$ -6 wells per data-point, Figure  
168 3D), consistent with prior findings that TTX has a slightly increased affinity for the inactivated  
169 channel<sup>29,30</sup>. Some sodium channel blockers can slow channel repriming after inactivation<sup>31</sup>. We  
170 examined this effect by varying the recovery period. Cells were exposed to a prepulse with fixed  
171 duration of 500 ms, a variable recovery period from 40 ms to 5120 ms, and a test pulse of 30 ms. Cells  
172 treated with DMSO vehicle showed nearly complete recovery within 40 ms. Carbamazepine, a

174 commonly used drug for the treatment of seizure and neurological pain, blocked recovery at 40 ms, but  
175 not at 80 ms. Amitriptyline had an even more dramatic effect, slowing the half-recovery time to  $280 \pm$   
176 36 ms, when tested at 3  $\mu$ M (mean  $\pm$  s.e.m.,  $n = 9$  wells per data-point). This result implies that  
177 amitriptyline has slow dissociation from the channel at resting membrane potential of -97 mV (Figure 3E,  
178 F).

179 Traditional voltage clamp protocols are flexible and precise in both time and voltage, while optical  
180 control of voltage is only semi-quantitative. Nonetheless, we explored stimulus intensity-dependent  
181 protocols, using the relation between steady state depolarization and illumination intensity (Figure 2,  
182 S1A) as a guide. We developed a protocol to probe separately voltage dependent activation and fast  
183 inactivation of  $\text{Na}_v1.7$ . Cells were exposed to a prepulse of 1 s duration with variable intensity from 1.7  
184 to 14 mW/cm<sup>2</sup>, corresponding to depolarizations of -84 to -47 mV. The blue light intensity was then  
185 stepwise increased to 14 mW/cm<sup>2</sup>, with no intervening recovery period. We quantified the amplitude of  
186 the fluorescence spike at the onset of the prepulse and the test pulse. The former probed channel  
187 activation, and the latter probed fast inactivation.

188 In conventional voltage clamp measurements, the region of overlap between activation and inactivation  
189 is called the “window current” and is important in governing cellular excitability. Mutations that  
190 increase the window current have been associated with pain disorders<sup>32</sup> and cardiac arrhythmias<sup>33</sup>.  
191 Compounds that decrease the window current by left-shifting inactivation or right-shifting activation  
192 have therapeutic potential<sup>32,34</sup>. We examined the effect of carbamazepine on voltage-dependent  
193 activation and fast inactivation. Consistent with observations from traditional electrophysiology<sup>32,34</sup>,  
194 carbamazepine reduced the overlap between activation and inactivation by leftward shifting the fast  
195 inactivation curve without altering the activation curve (Figure 3G, H). While optical electrophysiology is  
196 not able to quantify the window current, it can identify the qualitative mechanistic feature, the sign, and  
197 the approximate magnitude of the effect.

198 **Characterization of PF-04856264, a  $\text{Na}_v1.7$  specific inhibitor**

199 Recently, a subtype specific drug binding pocket was identified in the voltage sensor of Domain IV of  
200  $\text{Na}_v1.7$ <sup>35</sup>. Significant effort has gone into developing subtype-specific blockers, due to their potential  
201 analgesic applications. One such compound, PF-04856264, selectively blocks  $\text{Na}_v1.7$  in a state-  
202 dependent manner, with reported IC<sub>50</sub> of 28 nM when the holding potential is -70 mV<sup>35</sup>. The  
203 mechanistic details of the interaction of this compound with the channel have not been characterized.

204 We varied the precondition pulse length and found that even at 2 s of precondition pulse, PF-04856264  
205 failed to inhibit the channel at 100 nM (Figure 4A).

206 We hypothesized that the discrepancy between our measurements and literature results was due to the  
207 difference between the resting potential of our cells (~-97 mV) and the holding potential in the prior  
208 work (-70 mV). To control the resting potential of the  $\text{Na}_v1.7$ -OS HEK cells, we varied the extracellular  $\text{K}^+$   
209 concentration and observed approximately Nernstian behavior, as expected for a leak current  
210 dominated by  $\text{K}_{ir}2.1$  (Figure 4B). We elevated bath potassium from 2 mM to 8 mM, which decreased the  
211 magnitude of the resting potential to ~-70 mV. Under this condition and a 2 s prepulse, PF-04856264  
212 inhibited  $\text{Na}_v1.7$  mediated spikes with  $\text{IC}_{50}$  at 43 nM. We further investigated use dependent inhibition  
213 of  $\text{Na}_v1.7$  by PF-04856264 with 8 mM bath potassium, and observed both tonic and use dependent  
214 inhibition (Figure 4C). Our results show that binding of PF-04856264 is strongly dependent on the  
215 resting voltage even at potentials where  $\text{Na}_v1.7$  activation is minimal. Due to slow binding kinetics,  
216 sustained baseline depolarization is more effective than strong but brief depolarization at inducing  
217 binding.

## 218 **High throughput screening of $\text{Na}_v1.7$ inhibitors**

219  $\text{Na}_v1.7$  is widely considered to be a promising target for analgesic drugs<sup>36</sup>, so we sought to develop a  
220 high throughput screen based on Optopatch measurements in  $\text{Na}_v1.7$ -OS HEK cells. We used the ability  
221 to optically stimulate and record to screen for activity-dependent modulators of  $\text{Na}_v1.7$ . The platform  
222 was based around a commercial inverted microscope (Olympus IX-71) with an automated scanning  
223 stage and an air objective. Optogenetic stimulation and fluorescence imaging were performed through  
224 the objective. Fluorescence from a region 320 by 166  $\mu\text{m}$ , comprising approximately 150 cells, was  
225 binned on the detector and digitized at 100 Hz. We programmed the system to record sequentially from  
226 each well in a glass-bottomed 384 well plate.

227 We tested a library of 320 FDA approved drugs. Each well was treated with a single compound at 10  $\mu\text{M}$   
228 concentration. Amitriptyline (10  $\mu\text{M}$ ) and DMSO (0.1%) were used as positive and negative controls,  
229 respectively. Sixteen control wells (8 positive, 8 negative) were placed at the beginning and end of the  
230 plate. Cells were incubated with compound for 20 min. Each well was then stimulated with 8 pulses of  
231 blue light, 20 ms per pulse, 10 Hz, and the binned fluorescence was recorded. Automated scanning of  
232 the whole plate required 20 min.

233 Amitriptyline and DMSO showed robustly distinct firing patterns. DMSO wells showed consistent firing  
234 throughout the stimulus train. Amitriptyline wells showed rapid activity-dependent decrease in spike  
235 amplitude (Figure 5A). We also observed that some compounds induced more complex spiking patterns,  
236 either suppressing alternate spikes, or leading to erratic firing responses. We designed two simple  
237 parameters to capture the main features of use-dependent block, an important attribute of sodium  
238 channel blockers<sup>35,37,38</sup>. Let  $S_i$  be the height of the  $i^{\text{th}}$  spike ( $i$  runs from 1 to 8), and let  $\tilde{S}_i \equiv S_i / S_1$  be the  
239 height of the  $i^{\text{th}}$  spike divided by the height of the first spike (hence  $\tilde{S}_1 = 1$ ). We calculated the use  
240 dependence index as  $\Gamma = 1 - \langle \tilde{S}_i \rangle_{2-8}$ , where the subscripts indicate the range of spikes averaged. We  
241 also calculated a measure of recovery from inactivation via the standard deviation in the spike  
242 amplitude,  $\sigma = \langle (\tilde{S}_i - \langle \tilde{S}_i \rangle)^2 \rangle^{1/2}$ . This parameter was large for wells that showed alternating or erratic  
243 spike patterns. Thus every well was represented by a point on a two-dimensional ( $\sigma, \Gamma$ ) graph.  
244 Compound names and screening results are available in **Figure 5 Source Data File**. Remarkably, in this  
245 blinded screen, Doxepin was classified as functionally adjacent to the amitriptyline controls. Doxepin is  
246 a tricyclic antidepressant with structure and pharmacology very similar to those of amitriptyline (Figure  
247 5B).

248 Using  $\sigma$  alone to distinguish positive and negative controls, the  $Z'$  factor<sup>39</sup> for the assay was 0.57, within  
249 the range appropriate for a high-throughput screen<sup>39</sup>. We identified compounds for which  $\sigma$  was greater  
250 than 5 standard deviations from the average of negative controls. The hit rate was 12.2% by this  
251 measure, consistent with the notion that voltage gated sodium channels are promiscuous binders  
252 (Figure 5B)<sup>37</sup>.

253 The “hit” compounds showed diversity in their spike patterns. For amitriptyline, doxepin, and  
254 trifluoperazine, spike amplitude decayed monotonically throughout the pulse sequence. For other  
255 compounds, e.g. isradipine and iloperidone, spike amplitude alternated between even and odd stimuli, a  
256 pattern we called “alternans” (Figure 5C). We hypothesized that the alternans compounds had a more  
257 transient inhibitory effect on channel repriming, compared to the amitriptyline-like compounds.

258 To test this hypothesis and to relate the parameters measured by the screen to more conventional  
259 electrophysiological parameters, we performed voltage clamp recordings in the presence of several hits  
260 from the screen. A 200 ms prepulse and a 100 ms test pulse were separated by a recovery interval  
261 varying from 1 ms to 256 ms. We measured the ratio of the peak inward sodium currents at the test  
262 and pre-pulse. Control cells showed half-maximal recovery in 4.5 ms. Cells treated with isradipine or

263 iloperidone (alternans compounds), showed half-maximal recovery in 20 ms. Cells treated with  
264 amitriptyline or trifluoperazine (strong blockers) showed less than 50% recovery in 256 ms (Figure 5D, E).  
265 Thus compounds that clustered nearby in the optically measured ( $\sigma$ ,  $I$ ) graph, also showed similar  
266 effects by conventional patch clamp measurements.

267 For a compound to be a safe analgesic, it should not interact significantly with other  $\text{Na}_v$  channels,  
268 particularly the cardiac  $\text{Na}_v1.5$  channel. We created a  $\text{Na}_v1.5$ -OS HEK cell line analogous to the  $\text{Na}_v1.7$ -  
269 OS HEK cell line described above, and re-tested some hits from the screen against  $\text{Na}_v1.5$  (Figure 5-  
270 figure supplement 1). Unsurprisingly, on a ( $\sigma$ ,  $I$ ) plot, all hits arranged in a similar pattern for  $\text{Na}_v1.5$   
271 and  $\text{Na}_v1.7$ , indicating no subtype selectivity ( $\text{Na}_v1.7$ -selective compounds are exceedingly rare). In  
272 contrast, both PF-04856264 (1  $\mu\text{M}$ ) and TTX (1  $\mu\text{M}$ ) clustered with the negative controls when tested on  
273  $\text{Na}_v1.5$ , consistent with the known fact that neither of these compounds blocks  $\text{Na}_v1.5$  at the  
274 concentrations tested<sup>35,37</sup> (Figure 5F).

275 Kir2.1 is not considered as a promiscuous drug binder<sup>40</sup> and we are not aware of any compounds that  
276 block the conductance of channelrhodopsin or interfere with the voltage-dependent fluorescence of  
277 QuasAr2. Nonetheless, it is a formal possibility that false-positive readings could arise from compounds  
278 that modulate these other components. We developed a simple optical test for off-target effects and  
279 applied it to a panel of seven compounds selected for diverse mechanisms of  $\text{Na}_v1.7$  block. First, we  
280 added TTX (1  $\mu\text{M}$ ) to block  $\text{Na}_v1.7$ . We then applied steps of blue light (500 ms, 3.2 – 56 mW/cm<sup>2</sup>) and  
281 monitored the QuasAr2 fluorescence, which reported optically induced depolarizations without  
282 regenerative spikes (Figure 5-figure supplement 2A). We then performed the same measurement in the  
283 presence of TTX + test compound. Any drug interaction with Kir2.1, CheRiff, or QuasAr2 would alter the  
284 fluorescence response. Six of the seven tested compounds had minimal effect (within 7% of TTX,  $n = 9$ -  
285 10 wells per compound). Carbamazepine induced a slight decrease in fluorescence signal (20% smaller  
286 than TTX alone, Figure 5-figure supplement 2B).

287 We further verified the optical tests using manual patch clamp measurements. None of the eight tested  
288 compounds (seven drugs and TTX) affected CheRiff photocurrents (Figure 5-figure supplement 2C, 2D).  
289 Seven of the compounds had no effect on QuasAr2 fluorescence or voltage sensitivity relative to buffer  
290 control. Carbamazepine reduced QuasAr2 voltage sensitivity by 17%, consistent with the all-optical  
291 assay (Figure 5-figure supplement 2E, 2F). This off-target effect of carbamazepine had no effect on our  
292 optical assays of activity-dependent block (Figure 3F and Figure 3G), because fluorescence spike heights

293 were normalized to the height of the highest spike. A slight decrease in overall fluorescence signal is  
294 cancelled in this analysis.

295 Finally, we explored whether the OS-HEK cells could be used to screen for modulators of other channel  
296 classes. Kv4.3 is an A-type fast-activating voltage-gated potassium channel, active in the heart and  
297 central nervous system. Conventional ionic flux based optical approaches to screening for modulators  
298 of Kv4.3 are extremely challenging because the channel is inactivated at the resting potential of HEK cells.  
299 We stably expressed Kv4.3, Nav1.5 and Optopatch constructs in HEK cells. Voltage-clamp experiments  
300 revealed robust expression of Kv4.3, with a maximum current density of 218 pA/pF at +40 mV (Figure  
301 6A). Kv4.3 has very fast activation kinetics with a time constant  $\tau_{act} = 0.69$  ms at +40 mV. The inactivation  
302 of Kv4.3 can be best described as a double exponential decay<sup>41</sup>, with  $\tau_{fast} = 51$  ms and  $\tau_{slow} = 352$  ms at  
303 +40 mV (Figure 6A). We then transiently expressed Kir2.1 to prime Nav1.5 and Kv4.3 and called these  
304 cells Nav1.5-Kv4.3-OS cells.

305 When stimulated by pulses of blue light (100 ms, 50 mW/cm<sup>2</sup>), the presence of Kv4.3 led to a dramatic  
306 change in the optically induced and optically recorded action potential waveform, featuring a transient  
307 fast repolarization almost reaching resting potential before a recovery toward plateau potential (Figure  
308 6B). We then tested the effect of heteropoda toxin 2 (HpTx2), a potent and specific blocker of channels  
309 in the Kv4 family<sup>42</sup>. HpTx2 increased the action potential amplitude, consistent with its inhibition on the  
310 fast inactivated Kv4.3 peak current; HpTx2 also increased the plateau potential amplitude when  
311 compared at the end of the 100 ms light pulse, which can be explained by its inhibitory effect on the  
312 slow inactivated Kv4.3 current (Figure 6C). HpTx2 showed dose-dependent blockade, with an IC50 of  
313 252 nM, consistent with literature results (Figure 6D)<sup>43</sup>. However, the high rate at which test compounds  
314 blocked the Nav channel precluded use of these cells in high-throughput screening applications.  
315 Screening would require use of a Nav channel or a Nav channel mutant which is resistant to most drugs.

316

## 317 **Discussion**

318 Despite variable expression levels of optogenetic actuator and voltage indicator, we have shown that  
319 Optopatch assays can probe state-dependent pharmacology of Nav channel modulators, and can  
320 accurately report binding affinities and kinetics. Key to achieving this accuracy were (1) performing

321 measurements averaged over large numbers of cells, and (2) developing stimulus and analysis protocols  
322 that were insensitive to modest variations in expression levels of the optogenetic components.

323 Optical flux-based assays have been widely used in ion channel screens<sup>44</sup>. However, these assays  
324 typically only probe steady-state channel behavior. Flux-based assays are widely used, however,  
325 because they offer high throughput and high reproducibility. Recent advances in automated  
326 electrophysiology<sup>45</sup> enable control of membrane voltage in heterologous expression systems.  
327 Automated electrophysiology offers the advantage of direct control of voltage and measurement of  
328 current. However these techniques have lower throughput and higher cost than optical assays, only  
329 work on certain cell types, and can be challenging to optimize. Patch clamp measurements also involve  
330 a perturbation to the integrity of the cell membrane, which can lead to changes in cytoplasmic  
331 composition and artifacts from mechanosensitive channels<sup>46</sup>. The Optopatch assays developed here  
332 provide detailed and quantitative mechanistic information; are compatible with high-throughput  
333 screening; and are non-invasive. What are the limitations on throughput of optical electrophysiology  
334 screens? Here we performed serial measurements, one well at a time. At a measurement time of ~3  
335 s/well, a 384-well plate was scanned in ~20 min. There are no fundamental principles that prevent  
336 scaling to more densely packed wells (e.g. 1534 well plates), or to parallelizing the measurements.  
337 While optopatch measurements require high intensity red illumination, high SNR can be obtained at  
338 lower intensities than the 400 W/cm<sup>2</sup> we used here (Figure 2-figure supplement 4). Specialized  
339 instrumentation has been developed for sensitive fluorescence recording from multi-well plates<sup>47</sup>, and  
340 with such instrumentation one could achieve throughputs compatible with primary screening. Given  
341 greater parallelism of measurement, one could also implement more complex stimulus protocols such  
342 as we developed here, while maintaining adequate throughput.

343 Illumination intensities for imaging Arch-based GEVs are typically 10 to 100-fold greater than are used  
344 for imaging GFP-based GEVs. Thus it is natural to worry about phototoxicity from the red laser. A  
345 recent study explored phototoxicity in cultured mammalian U2OS cells.<sup>48</sup> Illumination at 200 W/cm<sup>2</sup>,  $\lambda$  =  
346 488 nm for 240 s led to 100% of the cells being either dead or “frozen”; while illumination at  
347 5,900 W/cm<sup>2</sup>,  $\lambda$  = 640 nm for 240 s led to undetectable cell death. Our observation of good cell viability  
348 at 400 W/cm<sup>2</sup>,  $\lambda$  = 640 nm is consistent with these literature results.

349 In principle, the screening approaches described here could be adapted to work with a red-shifted  
350 voltage-sensitive dye. Fluorescence signals would be more homogeneous than with a genetically

351 expressed indicator; one could more readily switch between cell lines; and there is a possibility that the  
352 imaging could be performed at lower illumination intensity, on conventional equipment. However,  
353 existing red-shifted dyes, e.g. PGH1<sup>49</sup> and Di-2-ANBDQ<sub>2</sub>PQ<sup>50</sup> still retain considerable excitation at the blue  
354 wavelengths used for channelrhodopsin activation, and these dyes are not at present commercially  
355 available.

356 Finally, we consider the diversity of channels for which Optopatch-style screens may be feasible. Here  
357 we demonstrated assays for  $Na_v1.7$ ,  $Na_v1.5$ , and  $K_v4.3$ . We previously demonstrated spiking HEK cells  
358 expressing  $Na_v1.3$ <sup>19</sup> and Hsu *et al.* demonstrated spiking CHO cells expressing  $Na_v1.2$ . HEK cell lines  
359 expressing  $Na_v1.1$  through  $Na_v1.8$  are commercially available, and a method for heterologous  
360 expression of chimeric  $Na_v1.9$  was recently demonstrated.<sup>51</sup> Voltage-gated  $Ca^{2+}$  channels can also  
361 mediate regenerative spiking and thus are also plausible targets for the assay. Fast and repetitive  
362 optogenetic activation of  $Ca_v3.2$  by channelrhodopsin2 has been achieved in HEK293T cells<sup>52</sup>. Recently,  
363 the state dependent inhibition of  $Ca_v1.3$  has been studied by channelrhodopsin stimulation protocols<sup>53</sup>.  
364 In principle, optogenetic activation could be applicable to other types of  $Ca_v$  channels. Delayed rectifier  
365 potassium channels such as hERG and  $K_v7$  may also be amenable to optical interrogation if co-expressed  
366 with an inactivation deficient  $Na_v$  channel. Modulation of the potassium current would manifest as a  
367 change in the action potential duration<sup>54</sup>.

368

### 369 **Acknowledgments**

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371 technical assistance, and Owen McManus for helpful discussions. This work was supported by the  
372 Howard Hughes Medical Institute, and US National Institutes of Health (NIH) grant 1-R01-EB012498-01.

373

### 374 **Competing Financial Interests**

375 AEC is a founder of Q-State Biosciences

376

377 **Figure legends**

378 **Figure 1.  $\text{Na}_v1.7$  Optopatch Spiking ( $\text{Na}_v1.7$ -OS) HEK cells.** (A) Genes expressed heterologously in  
379  $\text{Na}_v1.7$ -OS HEK cells.  $\text{K}_{ir}2.1$  maintains a hyperpolarized resting potential close to the  $\text{K}^+$  reversal potential.  
380  $\text{Na}_v1.7$  imparts electrical excitability. CheRiff depolarizes the cells upon optical excitation and can  
381 trigger a  $\text{Na}_v1.7$ -mediated action potential. QuasAr2 is excited by red light and emits near infrared  
382 fluorescence in a voltage-dependent manner. (B) Epifluorescence images of QuasAr2 and CheRiff-eGFP  
383 expressed in  $\text{Na}_v1.7$ -OS HEK cells. Scale bar 10  $\mu\text{m}$ .

384

385 **Figure 2. Biophysical characterization of  $\text{Na}_v1.7$ -OS HEK cells.** (A) CheRiff current in a  $\text{Na}_v1.7$ -  
386 Optopatch HEK cell. Membrane potential was held at -80 mV and then stepped for 2 s to -80 to +40 mV  
387 in 20 mV increments. During each depolarization, the cell was exposed to 5 pulses of blue light, 100 ms  
388 duration, with increasing intensity (1.7, 18, 50, 79, 93 mW/cm<sup>2</sup>). The horizontal dashed line indicates  
389 zero current. (B) I-V relation of CheRiff, under different light intensities. Currents were measured  
390 relative to baseline without blue light. Inset: Steady state photocurrent density as a function of blue  
391 light intensity, with a holding potential of -60 mV. (C) Peak  $\text{Na}_v1.7$  current densities as a function of  
392 depolarization potential. Membrane potential was held at -100 mV and then stepped for 100 ms to -  
393 90 mV to + 30 mV in 10 mV increments. These measurements were performed prior to transient  
394 expression of  $\text{K}_{ir}2.1$ . Inset: currents in the 10 ms interval following each voltage step. (D) I-V  
395 relationship of  $\text{K}_{ir}2.1$  expressed in  $\text{Na}_v1.7$ -OS HEK cells. Membrane potential was held at -100 mV and  
396 stepped for 500 ms to -130 mV to +30 mV in 10 mV increments. Inset: representative  $\text{K}_{ir}2.1$  current  
397 recording. Red line indicates the time point (4 ms after voltage step) at which the current was  
398 quantified. (E) Simultaneous voltage and QuasAr2 fluorescence recording from  $\text{Na}_v1.7$ -OS HEK cells. The  
399 cell was exposed to a series of blue laser pulses, 500 ms duration, with increasing intensities (1.1, 2.3,  
400 4.3, 7.0, 11, 15, 20, 26 mW/cm<sup>2</sup>) and QuasAr2 fluorescence was monitored with 640 nm excitation,  
401 400 W/cm<sup>2</sup>. Inset: overlay of the voltage and fluorescence recordings from the most intense blue pulse  
402 (26 mW/cm<sup>2</sup>).

403

404 **Figure 2-figure supplement 1. Current clamp recording of light triggered action potentials in  $\text{Na}_v1.7$ -  
405 OS HEK cells.** (A) An action potential recorded via manual patch clamp from a  $\text{Na}_v1.7$ -OS HEK cell cluster  
406 stimulated by 20 ms blue light pulse at 50 mW/ cm<sup>2</sup>. The dashed line indicates the firing threshold. (B)  
407 Plateau potential induced by different intensities of blue light stimulation. Current clamp recordings

408 were performed on  $\text{Na}_v1.7$ -OS HEK cell clusters stimulated with 500 ms blue light ranging from 1.1 to 84  
409  $\text{mW/cm}^2$ . (C) Membrane potential at 400 ms after onset of blue light stimulus as a function of the blue  
410 light intensity.

411

412 **Figure 2-figure supplement 2. Relationship between  $\text{Na}_v1.7$  current density and spike height.** (A)  
413 Combined current clamp and voltage clamp protocol in the presence of 3  $\mu\text{M}$  amitriptyline to prepare  
414 cells with varying  $\text{Na}_v1.7$  capacities. Initially, a current clamp protocol was applied in which a  
415 depolarizing pulse led to amitriptyline binding and complete channel block. After a variable recovery  
416 period,  $\Delta t$ , a test pulse of current induced a voltage spike whose amplitude was recorded. The voltage  
417 during the prepulse and recovery periods was then replayed in voltage clamp mode to induce an  
418 identical level of channel block. A step depolarization to -20 mV induced a spike in  $\text{Na}_v1.7$  current  
419 whose amplitude was recorded. For details see Methods. (B) Upper trace: current clamp recording  
420 showing voltage during 500 ms prepulse intervals, variable recovery period, and test pulses (asterisks).  
421 Bottom left: magnified view of voltage during test pulses. Bottom right: magnified view of  $\text{Na}_v1.7$   
422 current during test pulse under voltage clamp. (C) Voltage spike amplitude as a function of  $\text{Na}_v1.7$   
423 current amplitude for paired current-clamp and voltage-clamp recordings. Error bars represent s.e.m. of  
424  $n = 8$  cells.

425

426 **Figure 2-figure supplement 3. Cell-to-cell variability in Optopatch measurements.** (A)  $\text{Na}_v1.7$ -OS HEK  
427 cells in a confluent monolayer were imaged in the presence of 3  $\mu\text{M}$  amitriptyline with a 60 $\times$  oil  
428 immersion objective (numerical aperture 1.45). Cells were illuminated with a red laser at 400  $\text{W/cm}^2$ . A  
429 blue prepulse (500 ms, 50  $\text{mW/cm}^2$ ) depolarized membrane potential and allowed drug binding. A  
430 variable recovery period (40 ms to 5120 ms) led to partial unbinding. A blue test pulse (20 ms,  
431 50  $\text{mW/cm}^2$ ) probed the voltage spike induced by residual  $\text{Na}_v1.7$  capacity. Traces show fluorescence of  
432 QuasAr2 from five different single cells recorded in parallel. (B) Ratio of spike amplitude in test pulse to  
433 spike amplitude at the longest recovery time (5120 ms), as a function of recovery time for the five single  
434 cells shown in (A)

435

436 **Figure 2-figure supplement 4. Effects of intense red laser illumination.** (A) Heating by red laser  
437 illumination (635 nm, 400  $\text{W/cm}^2$ ). Temperatures were recorded with a small thermocouple not directly

438 illuminated by the laser. Measurements were performed in a single well of a 384-well plate with 36  $\mu$ L  
439 buffer, and in a 35 mm dish with 2 mL buffer (error bars represent s.e.m.,  $n$  = 4 replicates). (B) Test for  
440 photobleaching and phototoxicity.  $\text{Na}_v1.7$ -OS HEK cell were illuminated with red laser (635 nm,  
441 400 W/cm<sup>2</sup>) continuously for 10 min. Every one minute, the cells were stimulated with blue light (6  
442 pulses of 20 ms, 5 Hz, 50 mW/cm<sup>2</sup>). The first, sixth and the tenth spike trains are shown above. (C) SNR  
443 and spike height ( $\Delta F/F_0$ ) for each spike train in (B) as a function of red laser illumination time. Error bars  
444 represent s.d. of  $n$  = 6 spikes. (D) Effect of red laser intensity on SNR of QuasAr2 fluorescence spikes.  
445 Cells were illuminated with red laser at varying intensities and stimulated with pulses of blue light (8  
446 pulses of 20 ms, 10 Hz, 50 mW/cm<sup>2</sup>). (E) SNR of spikes recorded in (D) as a function of red laser intensity.  
447 Error bars represent s.d. of  $n$  = 8 spikes.

448

449 **Figure 3. Mechanistic studies of  $\text{Na}_v1.7$  blockers.** (A) Approximately 150  $\text{Na}_v1.7$ -OS HEK cells were  
450 stimulated with pulses of blue light (20 ms, 50 mW/cm<sup>2</sup>) at increasing frequencies (2 Hz, 4 Hz, 8 Hz) and  
451 their total QuasAr2 fluorescence was recorded with 635 nm excitation, 400 W/cm<sup>2</sup>. In control cells  
452 (blue), the fluorescence indicated spiking in response to each stimulus. After exposure to 200  $\mu$ M  
453 lidocaine (red), cells showed activity-dependent block at 4 Hz and 8 Hz, but not at 2 Hz (red arrows). (B)  
454 Simultaneous current clamp and QuasAr2 fluorescence recordings from a  $\text{Na}_v1.7$ -OS HEK cell cluster (4  
455 cells) stimulated with prepulses of varying length (20, 100, 200 and 500 ms; 50 mW/cm<sup>2</sup>) followed by  
456 200 ms recovery and a test pulse (30 ms, 50 mW/cm<sup>2</sup>). (C,D) Application of the protocol in (B) to dose-  
457 response curves for (C) amitriptyline or (D) TTX. Test pulse spike amplitude was normalized to its value  
458 in the presence of the lowest tested concentration of drug ( $n$  = 3-5 wells for amitriptyline per data-point;  
459  $n$  = 4-6 wells for TTX per data-point; ~150 cells per well). (E) Optical assay of  $\text{Na}_v1.7$  recovery from fast  
460 inactivation. A 500 ms prepulse (50 mW/cm<sup>2</sup>) populated the fast inactivated state and allowed drug  
461 binding. A variable recovery period (40 ms to 5120 ms) was followed by a 20 ms test pulse (50 mW/cm<sup>2</sup>).  
462 Traces show fluorescence of QuasAr2 for control cells and after addition of either 100  $\mu$ M  
463 carbamazepine or 3  $\mu$ M amitriptyline. (F) Ratio of spike amplitude in test pulse to spike amplitude at  
464 the longest recovery time (5120 ms), as a function of recovery time. Carbamazepine modestly slowed  
465 recovery and amitriptyline dramatically slowed recovery ( $n$  = 9 - 11 wells per curve, ~150 cells per well).  
466 (G) Optical protocol to measure voltage-dependent  $\text{Na}_v1.7$  activation and inactivation. Cells were  
467 stimulated with 1000 ms prepulse with increasing intensity (1.7, 3.6, 6.3, 9.8, 14 mW/cm<sup>2</sup>), immediately  
468 followed by a test pulse (150 ms, 14 mW/cm<sup>2</sup>). Traces show representative fluorescence recordings of

469 control and 100  $\mu$ M carbamazepine. (H) Effect of carbamazepine on activation and inactivation curves.  
470 Spike amplitudes were normalized to the maximum spike amplitude in the trace and were then plotted  
471 against prepulse intensity ( $n = 3$  wells per curve). Carbamazepine left-shifted the inactivation curve,  
472 decreasing the optically measured overlap between activation and inactivation.

473

474 **Figure 3-figure supplement 1. Use-dependent inhibition of  $\text{Na}_v1.7$  by amitriptyline and TTX.** Data  
475 used to produce Figures 3C, D. Plots show the fluorescence response evoked by the test pulse with  
476 variable duration prepulses. Each trace is the average of  $n = 3 - 5$  wells for amitriptyline or  $n = 4 - 6$   
477 wells for TTX.

478

479 **Figure 4. Effect of PF-04856264, a subtype-specific blocker, on  $\text{Na}_v1.7$ -OS HEK cells.** (A) Dose-response  
480 curves for PF-04856264 when stimulated with prepulses of different durations and with different bath  
481  $\text{K}^+$  concentrations ( $n = 4$  wells for each concentration). The optical protocol was as in Figure 3B, with  
482 prepulse duration specified in figure legends. (B) Comparison between membrane voltage predicted by  
483 the Nernst Equation (assuming pure  $\text{K}^+$  conductance) and recorded by manual patch clamp, as a function  
484 of bath  $[\text{K}^+]$  ( $n = 4-7$  cell clusters per data point). (C) Use-dependent inhibition of spiking in  $\text{Na}_v1.7$ -OS  
485 HEK cells by PF-04856264, at 8 mM external  $\text{K}^+$ . Cells were stimulated with eight pulses of blue light  
486 (20 ms, 50 mW/cm<sup>2</sup>) at 5 Hz and 10 Hz and QuasAr2 fluorescence was monitored with 635 nm excitation,  
487 400 W/cm<sup>2</sup>. After photobleaching correction, the QuasAr2 fluorescence in the absence or in the  
488 presence of 100 nM PF-04856264, was normalized to peak amplitude of the first spike at 5 Hz in the  
489 absence of the drug. Each trace was averaged from 4 wells. Inset: structure of PF-04856264.

490

491 **Figure 5. High throughput screening of a FDA-approved drug library in  $\text{Na}_v1.7$ -OS HEK cells.**

492 (A) QuasAr2 fluorescence from positive (amitriptyline) and negative (DMSO) control wells. Cells were  
493 stimulated with eight pulses of blue light (20 ms, 50 mW/cm<sup>2</sup>) at 10 Hz, and QuasAr2 fluorescence was  
494 monitored with 635 nm excitation, 400 W/cm<sup>2</sup>. (B) Screen results. The response of each well was  
495 parameterized by its use dependence index and standard deviation in spike amplitude. Positive controls  
496 (red) and negative controls (green) were well separated. Selected hits were chosen for further analysis.  
497 Inset: structures of amitriptyline and doxepin. (C) QuasAr2 fluorescence traces of doxepin,  
498 trifluoperazine, isradipine and iloperidone recorded in the screen. (D) Validation of select hits by manual  
499 electrophysiology.  $\text{Na}_v1.7$ -Optopatch cells were held at -100 mV. A 200 ms prepulse to 0 mV allowed

500 drug binding. Recovery times at -100 mV ranged from 1 ms to 256 ms. A test pulse to 0 mV, 100 ms  
501 duration, probed the degree of channel recovery. Blue:  $\text{Na}_v1.7$  current during prepulse. Red:  $\text{Na}_v1.7$   
502 current during test pulse. (Prepulse and test pulse currents have been time-shifted and overlaid for easy  
503 comparison). Each compound was tested at 10  $\mu\text{M}$ . (E) Quantification of compound effects on  $\text{Na}_v1.7$   
504 recovery from inactivation. The plots show ratio of current amplitude at test pulse to prepulse, as a  
505 function of recovery period ( $n = 3$  cells for each compound,  $n = 12$  cells for control). (F) Characterization  
506 of select hits from (B) in  $\text{Na}_v1.5$ -OS HEK cells. Cells were stimulated with eight pulses of blue light (20 ms,  
507 50 mW/cm<sup>2</sup>) at 4 Hz. The 4 Hz stimulus was selected because action potential width of  $\text{Na}_v1.5$ -OS cells  
508 lasted longer than 200 ms under control conditions. Data analyzed and plotted as in (B) ( $n = 4$ -6 wells  
509 per drug). Drug concentrations were TTX: 1  $\mu\text{M}$ , PF-04856264: 1  $\mu\text{M}$ , amitriptyline: 10  $\mu\text{M}$ ,  
510 trifluoperazine: 10  $\mu\text{M}$ , isradipine: 10  $\mu\text{M}$ , iloperidone: 30  $\mu\text{M}$ ).

511

512 **Figure 5-figure supplement 1. Fluorescence traces from  $\text{Na}_v1.5$ -OS HEK cells with different drugs.**  
513  $\text{Na}_v1.5$ -OS HEK cells were stimulated with eight pulses of blue light (20 ms, 50 mW/cm<sup>2</sup>) at 4 Hz, and  
514 QuasAr2 fluorescence was monitored with 635 nm excitation, 400 W/cm<sup>2</sup>. TTX and PF-04856264 had  
515 little effect on channel function. Amitriptyline and trifluoperazine showed strong use-dependent block.  
516 Isradipine and iloperidone showed use-dependent block with fast recovery, leading to alternating  
517 response amplitudes.

518

519 **Figure 5-figure supplement 2. Characterization of off-target effects via optical and manual patch**  
520 **assays.** (A) Optical assay to detect perturbations to  $\text{K}_v2.1$ , CheRiff, or QuasAr2 in  $\text{Na}_v1.7$ -OS HEK cells.  
521  $\text{Na}_v1.7$  was blocked with 1  $\mu\text{M}$  TTX. Cells were stimulated with increasing intensities of blue light (500  
522 ms, 3.2, 12, 27, 41, 56 mW/cm<sup>2</sup>), and QuasAr2 fluorescence was monitored with 635 nm excitation,  
523 400 W/cm<sup>2</sup>. Fluorescence with a test compound (10  $\mu\text{M}$  amitriptyline, 100  $\mu\text{M}$  carbamazepine, 10  $\mu\text{M}$   
524 trifluoperazine, 1  $\mu\text{M}$  PF-04856264, 200  $\mu\text{M}$  lidocaine, 10  $\mu\text{M}$  isradipine, 10  $\mu\text{M}$  iloperidone) was  
525 compared to TTX alone. The fluorescence changes were normalized to that of TTX only treated cells  
526 under 56 mW/cm<sup>2</sup> stimulation. (B) Mean fluorescence changes in (A) during the blue stimuli as a  
527 function of stimulus intensity. Error bars represent s.e.m. of  $n = 9$ -10 wells. (C) Voltage clamp protocol  
528 to test for drug effects on CheRiff photocurrent. Cells were held at -80 mV and then stepped to -60 mV  
529 to +40 mV in 20 mV increments. During each step depolarization, a blue light pulse (100 ms, 0.5 W/cm<sup>2</sup>)

530 was applied to activate CheRiff current. Control trace and trace after carbamazepine treatment are  
531 shown as an example. (D) I-V relationship of CheRiff current under control condition (before drug  
532 treatment) and with test compounds at the same concentrations as in (A). To control for cell-to-cell  
533 variations in CheRiff expression, the current amplitudes were normalized to that of control at -80 mV.  
534 Error bars represent s.e.m.,  $n = 29$  cells for control,  $n = 3-4$  cells for each compound. (E) Voltage-clamp  
535 measurements to test for drug effects on QuasAr2 voltage sensitivity. QuasAr2 fluorescence was  
536 monitored with 640 nm excitation,  $400 \text{ W/cm}^2$  while membrane voltage was modulated as shown.  
537 Single cell recordings were performed before and after addition of drug or buffer control. Example  
538 traces show fluorescence before and after addition of buffer, lidocaine or carbamazepine. (F) Mean  
539 QuasAr2 fluorescence as a function of voltage in the presence of test compounds at the same  
540 concentrations as in (A). To control for cell-to-cell variation in QuasAr2 expression, the fluorescence  
541 changes were normalized to the pre-drug fluorescence at +40 mV. Error bars represent s.e.m.,  $n = 30$   
542 cells for control,  $n = 3-4$  cells for each compound.

543

544 **Figure 5-source data 1.** Spreadsheet containing compound names and screening results.

545

546 **Figure 6. Optopatch assay of Kv4.3 function.** (A) Voltage clamp recording of Kv4.3 current in Na<sub>v</sub>1.5-  
547 Kv4.3 Optopatch HEK cells. The bath contained 30  $\mu\text{M}$  TTX to block the Na<sub>v</sub>1.5 current. Cells were held at  
548 -70 mV and then subjected to 1 s steps to -60 mV to +40 mV in 10 mV increments. Peak Kv4.3 current  
549 densities were 218 pA/pF. (B) Na<sub>v</sub>1.5-Kv4.3-OS HEK cells were probed with simultaneous current clamp  
550 and QuasAr2 fluorescence. The cells were stimulated with a pulse of blue light (100 ms, 50 mW/cm<sup>2</sup>),  
551 and QuasAr2 fluorescence was monitored with 640 nm excitation,  $400 \text{ W/cm}^2$ . Kv activation led to a  
552 narrow action potential width, followed by Kv inactivation and a return to steady-state depolarization. (C)  
553 Average QuasAr2 fluorescence traces from Na<sub>v</sub>1.5-Kv4.3-OS HEK cells treated with HpTx-2 ( $n = 3-4$  wells  
554 for each concentration). (D) Dose-response curve of HpTx-2 on Na<sub>v</sub>1.5-Kv4.3-OS HEK cells. Drug effect  
555 was quantified by the fluorescence at the peak repolarization (~40 ms after onset of stimulus) relative to  
556 peak fluorescence intensity under 1200 nM HPTX2 treatment.

557 **Materials and methods**

558 **Genetic engineering of  $\text{Na}_v1.5$ -OS,  $\text{Na}_v1.5$ - $\text{K}_v4.3$ -OS and  $\text{Na}_v1.7$ -OS cells**

559 The pIRESpuro3- $\text{Na}_v1.5$  and pcDNA3- $\text{K}_v4.3$  plasmids were obtained from ChemCORE at Johns Hopkins  
560 University. The Optopatch construct contains coding sequences of CheRiff-eGFP and QuasAr2-  
561 mOrange2, separated by a P2A self-cleaving peptide sequence. The entire Optopatch construct was  
562 cloned into a modified FCK lentivirus vector (mFCK), in which the original CaMKII promoter was replaced  
563 by a CMV promoter. The Kir2.1 cDNA was amplified from Addgene plasmid 32669 (pENTR-L5-Kir2.1-  
564 mCherry-L2) and cloned into a pLX304 lentivirus vector that contained a blasticidin selection marker.  
565 The Kir2.1 cDNA was also cloned into pIREShyg vector using the Gibson assembly method. The  $\text{K}_v4.3$   
566 cDNA was amplified from pcDNA3- $\text{K}_v4.3$  plasmid and then cloned into pIREShyg vector using the Gibson  
567 assembly method<sup>55</sup>.

568 HEK293 cells were transected with pIRESpuro3- $\text{Na}_v1.5$  using Transit-293 Transfection Reagent  
569 (Mirus Bio) following manufacturer's instruction. After 48 hrs of transfection, puromycin was added to a  
570 final concentration of 2  $\mu\text{g}/\text{mL}$ . Cells were selected for 14 days to stabilize the expression of  $\text{Na}_v1.5$ .  
571 Surviving cells were subsequently transduced with low-titer mFCK-Optopatch lentivirus. After 10 days of  
572 infection, all the GFP positive cells were enriched by fluorescence activated cell sorting (FACS). This  
573 polyclonal  $\text{Na}_v1.5$ -Optopatch stable cell line was used to generate the  $\text{Na}_v1.5$ -OS and  $\text{Na}_v1.5$ - $\text{K}_v4.3$  OS  
574 cells.

575 To generate  $\text{Na}_v1.5$ -OS cells,  $\text{Na}_v1.5$ -Optopatch cells were transduced by pLX304-Kir2.1  
576 lentivirus. After 48 hrs of transduction,  $\text{K}_v2.1$  expressing cells were selected by 5  $\mu\text{g}/\text{mL}$  blasticidin. At  
577 the same time, 2  $\mu\text{g}/\text{mL}$  puromycin was also included to ensure the stable expression of  $\text{Na}_v1.5$ . Cells  
578 were cultured for 14 days and then single cells were dispersed in wells of a 48 well plate. Monoclonal  
579  $\text{Na}_v1.5$ -OS lines were screened via Optopatch measurements for robust generation of action potentials  
580 under blue laser stimulus, and corresponding QuasAr2 fluorescence transients with SNR greater than 30.

581 To generate  $\text{Na}_v1.5$ - $\text{K}_v4.3$ -OS cells,  $\text{Na}_v1.5$ -Optopatch cells were transiently transfected by  
582 pIREShyg- $\text{K}_v4.3$ . Two days after transfection, 200  $\mu\text{g}/\text{mL}$  hygromycin was used to establish the  $\text{Na}_v1.5$ -  
583 Optopatch- $\text{K}_v4.3$  monoclonal stable cell line. Each monoclonal cell line was optically evaluated for  
584 spiking and fast repolarization behavior after transient transfection of pIREShyg- $\text{K}_v2.1$  plasmid. The best  
585 monoclonal cell line ( $\text{Na}_v1.5$ -Optopatch- $\text{K}_v4.3$ ) was further expanded and  $\text{Na}_v1.5$ - $\text{K}_v4.3$ -OS cells can be  
586 reliably generated by transient transfection of Kir2.1 into this monoclonal cell line.

587 The  $\text{Na}_v1.7$ -OS HEK cells were generated based on a  $\text{Na}_v1.7$  stable cell line established by G418  
588 selection, a kind gift from Dr. Bruce Bean at Harvard University. This stable cell line was transduced with

589 Optopatch by mFCK-Optopatch lentivirus. After 10 days, GFP positive ( $\text{Na}_v1.7$ -Optoptach) cells were  
590 enriched by FACS. We attempted, unsuccessfully, to further stabilize  $\text{K}_{ir}2.1$  in these  $\text{Na}_v1.7$ -optoptach  
591 cells by using pLX304- $\text{K}_{ir}2.1$  lentivirus transduction. Surviving cells after blasticidin selection were not  
592 able to fire action potentials, likely due to poor expression level of  $\text{K}_{ir}2.1$ . Therefore, single cells of  
593  $\text{Na}_v1.7$ -optoptach cells were dispersed into a 48 well plate and each  $\text{Na}_v1.7$ -optoptach monoclonal cell  
594 line was evaluated by transient transfection of pIREShyg-Kir2.1 using lipofectamine 2000 (Invitrogen)  
595 following manufacturer's instruction. The best  $\text{Na}_v1.7$ -optoptach monoclonal line that produced robust  
596 spikes with corresponding high SNR QuasAr2 fluorescence was selected and further expanded. The  
597 transfected cells are called  $\text{Na}_v1.7$ -OS cells.

598 Cells tested negative for mycoplasma contamination. Absence of contamination from other cell  
599 lines was ensured by growing up cells from a single clone.

600

#### 601 **Cell culture**

602  $\text{Na}_v1.5$ -OS,  $\text{Na}_v1.5$ -Optopatch- $\text{K}_v4.3$  cells, and  $\text{Na}_v1.7$ -Optopatch HEK cell lines were maintained in  
603 Dulbecco's Modified Eagle Medium (DMEM) with 10% fetal bovine serum, penicillin (100 U/mL),  
604 streptomycin (100  $\mu\text{g}/\text{mL}$ ). For  $\text{Na}_v1.5$ -OS cells, 2  $\mu\text{g}/\text{mL}$  puromycin and 5  $\mu\text{g}/\text{mL}$  blasticidin were  
605 included in the medium to maintain expression of  $\text{Na}_v1.5$  and  $\text{K}_{ir}2.1$ . For  $\text{Na}_v1.5$ -Optopatch- $\text{K}_v4.3$  cells, 2  
606  $\mu\text{g}/\text{mL}$  puromycin and 200  $\mu\text{g}/\text{mL}$  hygromycin were included in the medium to maintain expression of  
607  $\text{Na}_v1.5$  and  $\text{K}_v4.3$ . For  $\text{Na}_v1.7$ -Optopatch cells, 500  $\mu\text{g}/\text{mL}$  of G418 was included in the medium to  
608 maintain  $\text{Na}_v1.7$  expression.

609

#### 610 **Electrophysiology in HEK cells**

611 Electrophysiology measurements were performed in a bath solution of Tyrode's, containing (in mM):  
612 125 NaCl, 2 KCl, 2  $\text{CaCl}_2$ , 1  $\text{MgCl}_2$ , 10 HEPES, 30 glucose. The pH was adjusted to 7.3 with NaOH and the  
613 osmolality was adjusted to 305-310 mOsm with sucrose. Filamented glass micropipettes (WPI) were  
614 pulled to a resistance of 4 – 7  $\text{M}\Omega$  and filled with internal solution containing 140 mM KCl, 1 mM  $\text{MgCl}_2$ ,  
615 10 mM EGTA, 10 mM HEPES, 3 mM Mg-ATP, pH adjusted to 7.3 with KOH. To record CheRiff and  
616  $\text{Na}_v1.7$  current,  $\text{Na}_v1.7$ -Optopatch HEK cells were replated onto 0.02 mg/mL poly-d-lysine coated glass-  
617 bottom dishes (In Vitro Scientific) at a density of ~10,000 cells/cm<sup>2</sup>. The patch clamp recording was  
618 performed 4 - 8 hours after re-plating when most cells had firmly attached to the glass and were still  
619 dispersed as single cells. The whole cell voltage clamp recordings were acquired using an Axopatch 200B  
620 amplifier (Molecular Devices), filtered at 5 kHz with the internal Bessel filter and digitized with a

621 National Instruments PCIE-6323 acquisition board at 10 kHz. The series resistance and membrane  
622 capacitance were compensated, and whole cell membrane capacitance was obtained by direct reading  
623 from the amplifier. CheRiff mediated current was triggered by illumination from a blue laser (488 nm,  
624 50 mW, Omicron PhoxX) that was sent through an acousto-optic modulator (AOM; Gooch and Housego  
625 48058-2.5-.55-5W) for rapid control over its intensity. The Kir2.1 current was recorded from Nav1.7-OS  
626 HEK cells by using the same configuration with 1  $\mu$ M of TTX in the bath solution to block Na<sub>v</sub>1.7 current.  
627 The K<sub>v</sub>4.3 current was recorded from Na<sub>v</sub>1.5-Optopatch-K<sub>v</sub>4.3 cells with 30  $\mu$ M of TTX in the bath  
628 solution to block Na<sub>v</sub>1.5 current.

629

630 To correlate Na<sub>v</sub>1.7 current density with voltage spike amplitude we performed alternate single-cell  
631 current clamp and voltage-clamp measurements in the presence of 3  $\mu$ M amitriptyline. We used cells  
632 not expressing K<sub>ir</sub>2.1 to avoid confound from K<sub>ir</sub> currents. Paired current- and voltage-clamp protocols  
633 were always performed on the same cell. For both protocols, an extended prepulse depolarization  
634 induced amitriptyline binding and complete channel block. A recovery interval at -100 mV of variable  
635 duration led to partial channel recovery. A test depolarizing pulse of either current or voltage then  
636 probed the response of the recovered channels.

637

638 In the current clamp protocol, holding current,  $i_h$ , was adjusted between -100 to -50 pA to attain a  
639 steady-state voltage of approximately -100 mV. The cell was then stimulated with a depolarizing current  
640 pulse of magnitude -0.5  $i_h$  for 500 ms. This current brought the steady-state voltage to ~0 mV. The  
641 current was then brought back to  $i_h$  for a recovery period of variable duration from 40 - 5120 ms. Finally,  
642 the cell was stimulated with a test current pulse of magnitude -0.5  $i_h$  for 20 ms to induce a voltage spike  
643 whose amplitude we recorded. Then the cell was switched to voltage-clamp mode. The holding  
644 potential was -100 mV. To match precisely the degree of channel block in the current-clamp and  
645 voltage-clamp protocols, the voltage prepulse and recovery waveforms were copied directly from the  
646 voltage recorded during the immediately preceding current-clamp protocol. The test pulse comprised a  
647 20 ms step depolarization to -20 mV. The inward Na<sub>v</sub>1.7 current at each test pulse was then measured.

648

649

650 **Simultaneous electrophysiology and Optopatch recording in HEK cells**

651 The Na<sub>v</sub>1.7-Optopatch monoclonal cell line was transfected with pIREShyg-Kir2.1 plasmid using  
652 lipofectamine 2000 following standard protocols. The resulting Nav1.7-OS HEK cells were recorded

653 48 hrs after transfection. The day before recording, cells were replated onto 35 mm glass-bottom dishes  
654 (In Vitro Scientific) at a density of ~10,000 cells/cm<sup>2</sup>. At the time when recording was performed, the  
655 cells formed small clusters comprising 3 - 4 cells. The whole cell current clamp recording was performed  
656 on these small clusters under the I-Clamp Normal configuration of the Axopatch 200B amplifier. The  
657 liquid junction potential was measured and corrected by the standard Neher method<sup>56</sup>.

658 Patch clamp and fluorescence imaging data were synchronized by clocking the camera with  
659 analog output from National Instruments PCIE-6323 acquisition board while using the same clock for  
660 driving patch clamp inputs and outputs. The imaging experiments were conducted on a home-built  
661 inverted fluorescence microscope<sup>10</sup>. Briefly, QuasAr2 was excited by combined illumination from two  
662 red lasers (640 nm, 140 mW, Coherent Obis 637-140 LX and 640 nm, 100 mW, Coherent CUBE 640-  
663 100C) via a polarizing beam splitter. The red beam was expanded and focused onto the back focal plane  
664 of a 60x oil-immersion objective (60x APO, NA 1.49, Olympus). CheRiff was activated by illumination  
665 from a blue laser (488 nm, 50 mW, Omicron PhoxX), which was modulated by an acousto-optic  
666 modulator receiving control signals from a National Instruments PCIE-6323 acquisition board. During a  
667 typical Optopatch experiment, both blue and red lasers were reflected into the sample plane by a quad-  
668 band dichroic mirror (Di01-R405/488/561/635-25x36, Semrock). The red laser intensity was maintained  
669 at 400 W/cm<sup>2</sup>, while the blue laser intensity was modulated via the AOM and ranged from 1-  
670 100 mW/cm<sup>2</sup>. A 710/100-nm bandpass filter (Chroma, HQ710/100) was used for QuasAr2 imaging, and  
671 a variable-zoom camera lens (Sigma 18-200 mm f/3.5-6.3 II DC) was used to image the sample onto an  
672 EMCCD camera (Andor iXon Ultra 897), with 512x 512pixels. The variable zoom enabled imaging at a  
673 range of magnifications while maintaining the high light-collection efficiency of the oil-immersion  
674 objectives. Data were acquired with a ROI of 128x128 pixels at 4 × 4-pixel binning to achieve a frame  
675 rate of 200 frames/s.

676

#### 677 **Optopatch measurements on pharmacology of Nav1.7-OS HEK cells**

678 The Nav1.7-Optopatch monoclonal cell line was transfected with pIREShyg-Kir2.1 plasmid using  
679 lipofectamine 2000. A glass-bottom 384-well plate (P384-1.5H-N, Cellvis) was treated with 0.02 mg/mL  
680 poly-d-lysine to promote cell adhesion. At 24 hrs after transfection, cells were replated onto the  
681 multiwell plate at a density of ~20,000 cells/well in 50 µL of culture medium. The imaging experiments  
682 were performed at 48 hrs after transfection when the cells formed a confluent monolayer. The cells  
683 were washed with Tyrode's solution once and then each well is filled with 30 µL of Tyrode's solution. For

684 drug additions, 6  $\mu$ L drug solution at 6x target concentration was added to each well. Cells incubated in  
685 drug for 20 min at room temperature before imaging.

686 Experiments were conducted on an inverted epi-fluorescence microscope (Olympus IX-71)  
687 equipped with an automated scanning stage (Ludl electronics MAC 6000). Briefly, illumination from a  
688 red laser (635 nm, 500 mW, Dragon Lasers MRL-635-500mW) was expanded and focused onto the back  
689 focal plane of a 20 $\times$  air objective (NA 0.75, Olympus -UPlanSApo 20 $\times$ /0.75). Illumination from a blue  
690 laser (473 nm, 50 mW, Dragon Lasers MBL-473-50mW) was sent through an acousto-optic tunable filter  
691 (AOTF; Gooch and Housego 48058) for rapid intensity modulation. The red illumination intensity at the  
692 sample was 400 W/cm<sup>2</sup>. QuasAr2 fluorescence was filtered by a 710/100-nm bandpass filter (Chroma,  
693 HQ710/100) and collected by an EMCCD camera (Andor iXon Ultra 897). Data were acquired with a  
694 full camera chip of 512 $\times$  512 pixels at 16  $\times$  16-pixel binning to achieve a frame rate of 100 frames/s.

695

#### 696 **High throughput screening on Nav1.7-OS HEK cells by Optopatch measurements**

697 Cells were plated in a 384 well plate as above. After the cells formed a confluent monolayer (48 hrs  
698 after transfection) the cells were washed with Tyrode's solution once and then each well was filled with  
699 20  $\mu$ L of Tyrode's solution. A compound library consisting of 320 FDA-approved drugs was purchased  
700 from Broad Institute at 10 mM stock concentration in DMSO and then diluted to 30  $\mu$ M in Tyrode's  
701 solution. 10  $\mu$ L of the diluted compounds were added to the cell plate (Well A3-P22) to achieve a final  
702 concentration of 10  $\mu$ M. Wells A2-H2 and A23-H23 were treated with 0.1% DMSO vehicle and used as  
703 negative controls. Wells I2-P2 and I23-P23 were treated with 10  $\mu$ M amitriptyline and used as positive  
704 controls. After 20 min of drug incubation, the 384-well plate was placed on the microscope stage and  
705 each well was imaged serially.

706 The scanning started at well A2 and ended in well P23 in a column-wise manner. Each well was  
707 exposed to eight pulses (20 ms) of blue laser (50 mW/cm<sup>2</sup>) at 10 Hz to stimulate the firing of Nav1.7-OS  
708 HEK cells. The QuasAr2 fluorescence from each well was collected as above. Data were saved as a single  
709 tiff stack at the end of scanning.

710

#### 711 **Imaging processing and data analysis**

712 Imaging data were stored as a tiff stack and loaded into ImageJ software. For data acquired at high  
713 magnification (60 $\times$ ), a rectangular ROI surrounding the cells of interest was manually selected.  
714 Background fluorescence was determined by measuring the mean intensity of a nearby cell free region  
715 and was subtracted from the cell fluorescence. For data acquired from cell monolayers under low

716 magnification (20 $\times$ ), a rectangular ROI (400  $\times$  208 pixels) covering the region with most intense laser  
717 illumination was selected. This ROI corresponds to a 320  $\mu\text{m} \times$  166  $\mu\text{m}$  area on the sample plane,  
718 containing approximately 150 cells. The mean intensity within this ROI was calculated for all frames of  
719 the tiff stack. To calculate  $\Delta F/F_0$ , background fluorescence was determined by measuring the intensity  
720 from a well plated with parental HEK cells without QuasAr2 expression. After background subtraction,  
721 the data were further analyzed to extract spike parameters. Briefly, intensity traces were corrected for  
722 photobleaching by dividing the raw intensity by a median filtered copy of the intensity. Spike amplitude  
723 was defined as the difference between the maximum point of an action potential and the baseline. The  
724 use dependence index was defined as the fractional reduction of the spike amplitude averaged from the  
725 second to the eighth stimulus, compared to the initial stimulus.

726 Dose-response curves were fitted with the Hill equation  $y = \text{START} + (\text{END} - \text{START}) / [1 + (\text{IC50}/S)^n]$ ,  
727 where START and END are the values of the parameter at minimum and maximum drug concentration,  
728 IC50 is the drug concentration at 50% maximum effect, S is the drug concentration, and n is a measure  
729 of cooperativity. The Z' factor for the screen was calculated as  $Z' = 1 - 3(\sigma_p + \sigma_n) / |(\mu_p - \mu_n)|$ , where  $\sigma_p$  is the  
730 standard deviation of the positive controls,  $\sigma_n$  is the standard deviation of the negative controls,  $\mu_p$  is the  
731 mean of the positive controls and  $\mu_n$  is the mean of the negative controls.

732 The activation time constant of K<sub>v</sub>4.3,  $\tau_{\text{act}}$ , was determined by fitting the activation current trace  
733 using the equation:  $i(t) = a(1 - e^{-t/\tau_{\text{act}}})^4 + b$ . The inactivation time constants,  $\tau_{\text{fast}}$  and  $\tau_{\text{slow}}$  of  
734 K<sub>v</sub>4.3 were determined by fitting the inactivation current trace using the equation:  $i(t) = a e^{-t/\tau_{\text{fast}}} +$   
735  $b e^{-t/\tau_{\text{slow}}} + c$ .

736

### 737 **Statistics**

738 Information on number of replicates for each experiment is given in figure legends. For manual patch  
739 clamp measurements, sample size was predetermined to be  $> 5$  cells, following standard practice. For  
740 optical electrophysiology measurements, sample size was predetermined to be  $> 100$  cells. These  
741 sample sizes were selected for feasibility of measurement. In the screen of the FDA library, one of the  
742 32 control wells showed an anomalous spiking pattern (visible in Fig. 5A) and was omitted from Fig. 5B.

743

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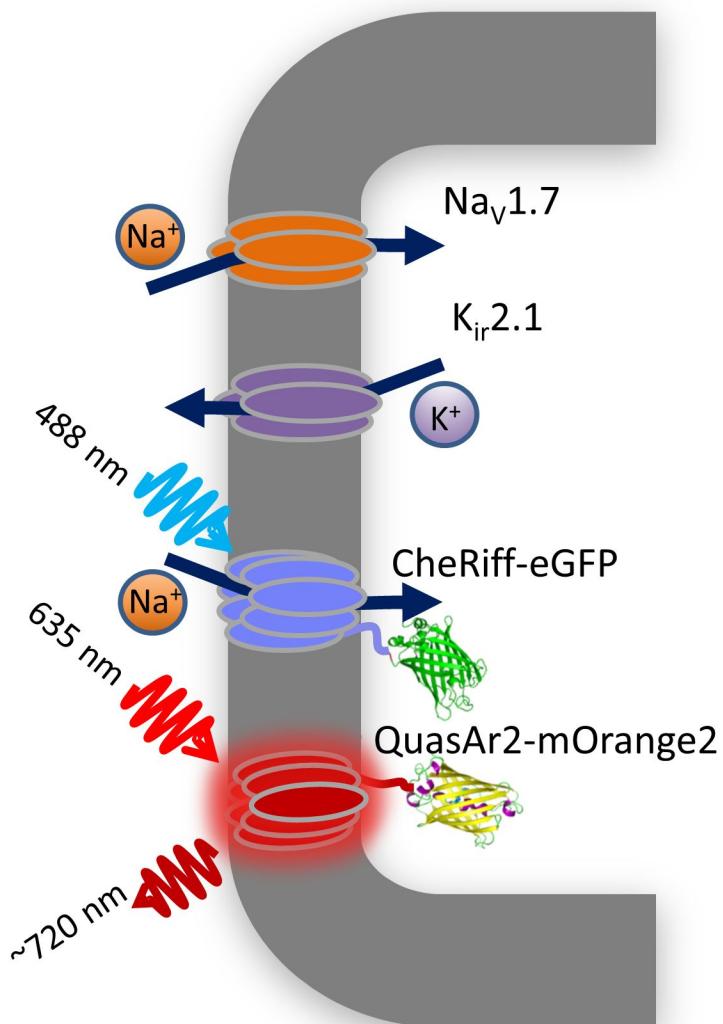
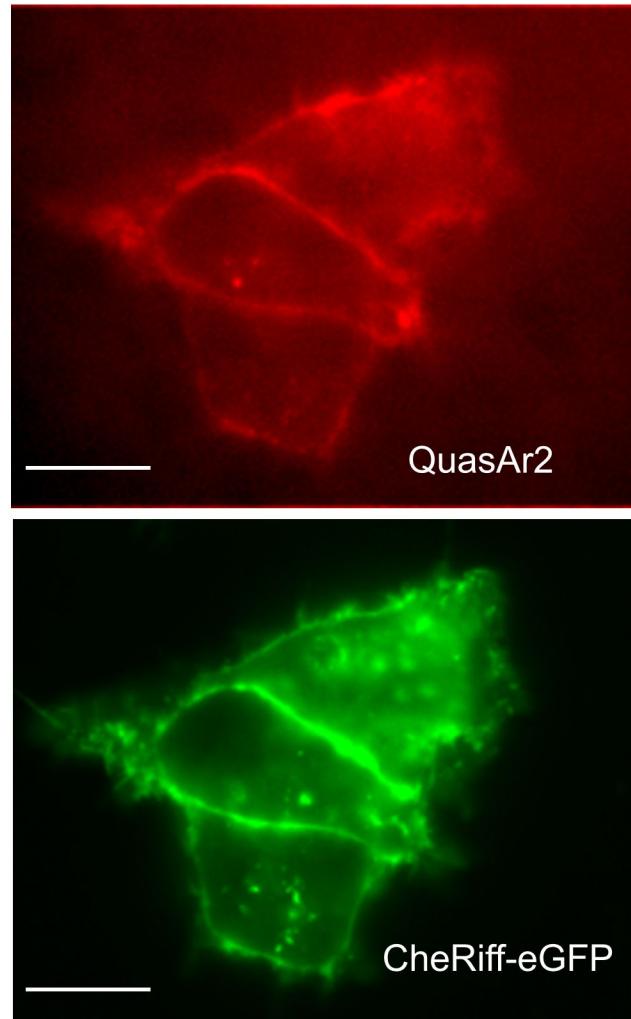
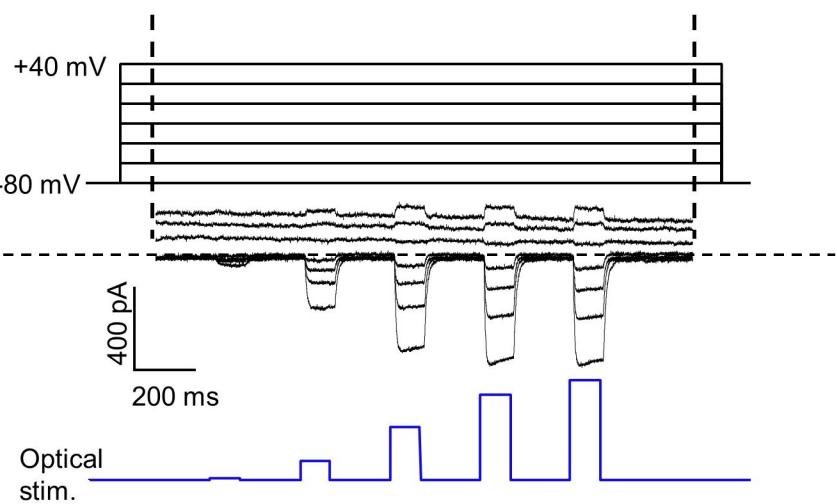
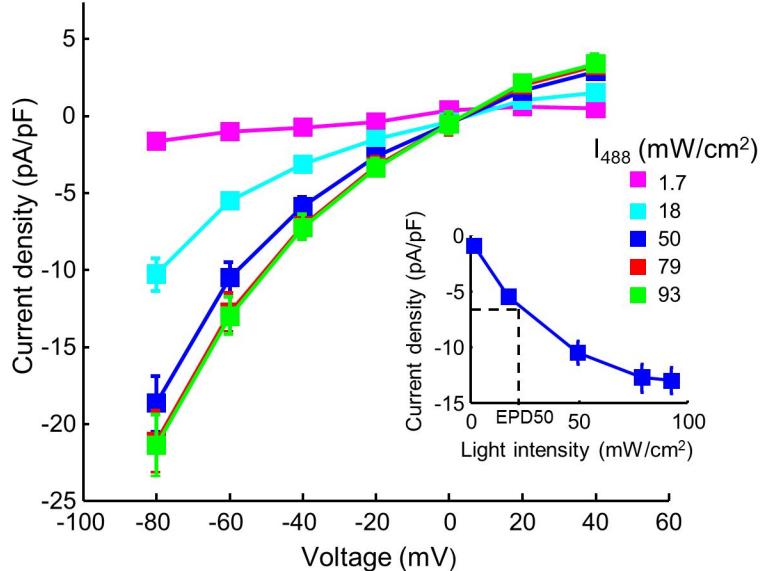
**A****B**

Figure 1

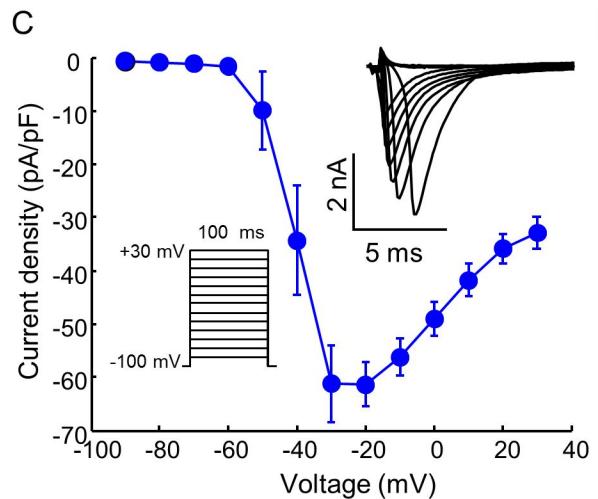
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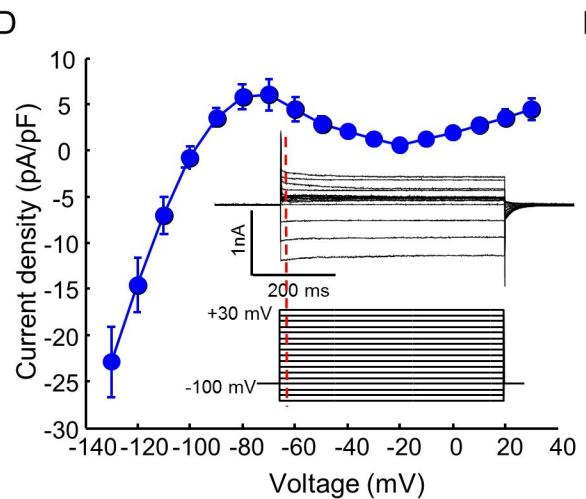
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C



D



E

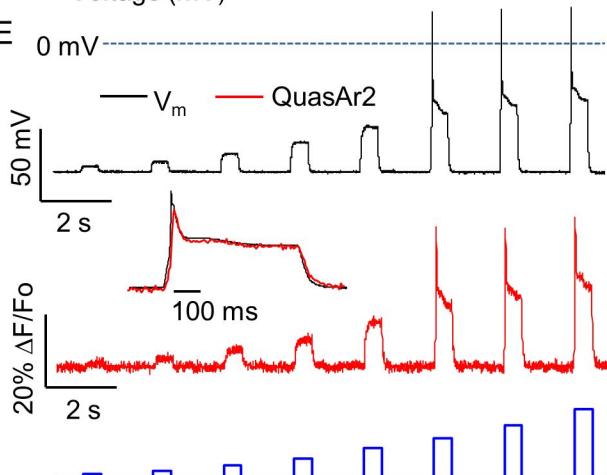


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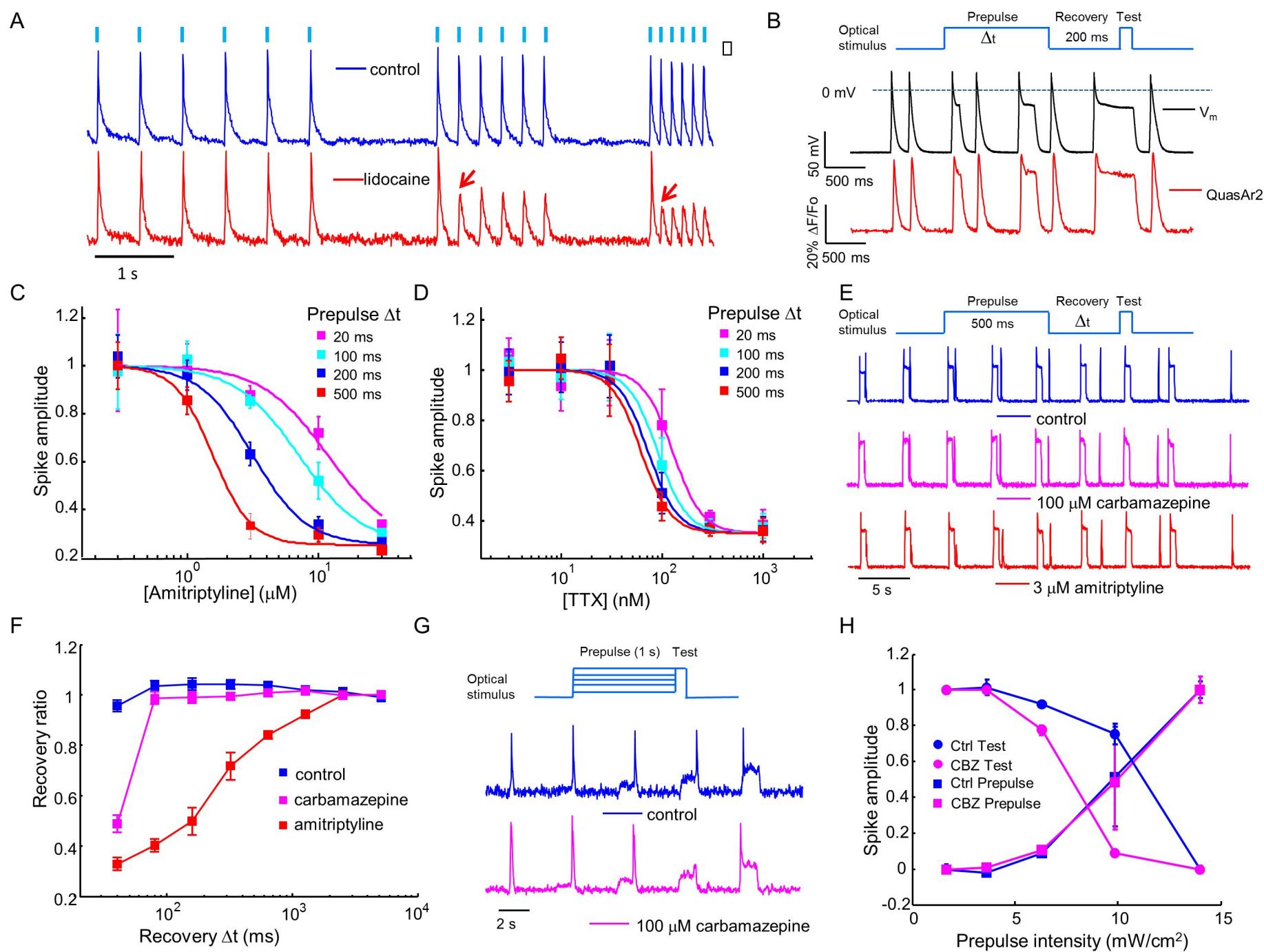


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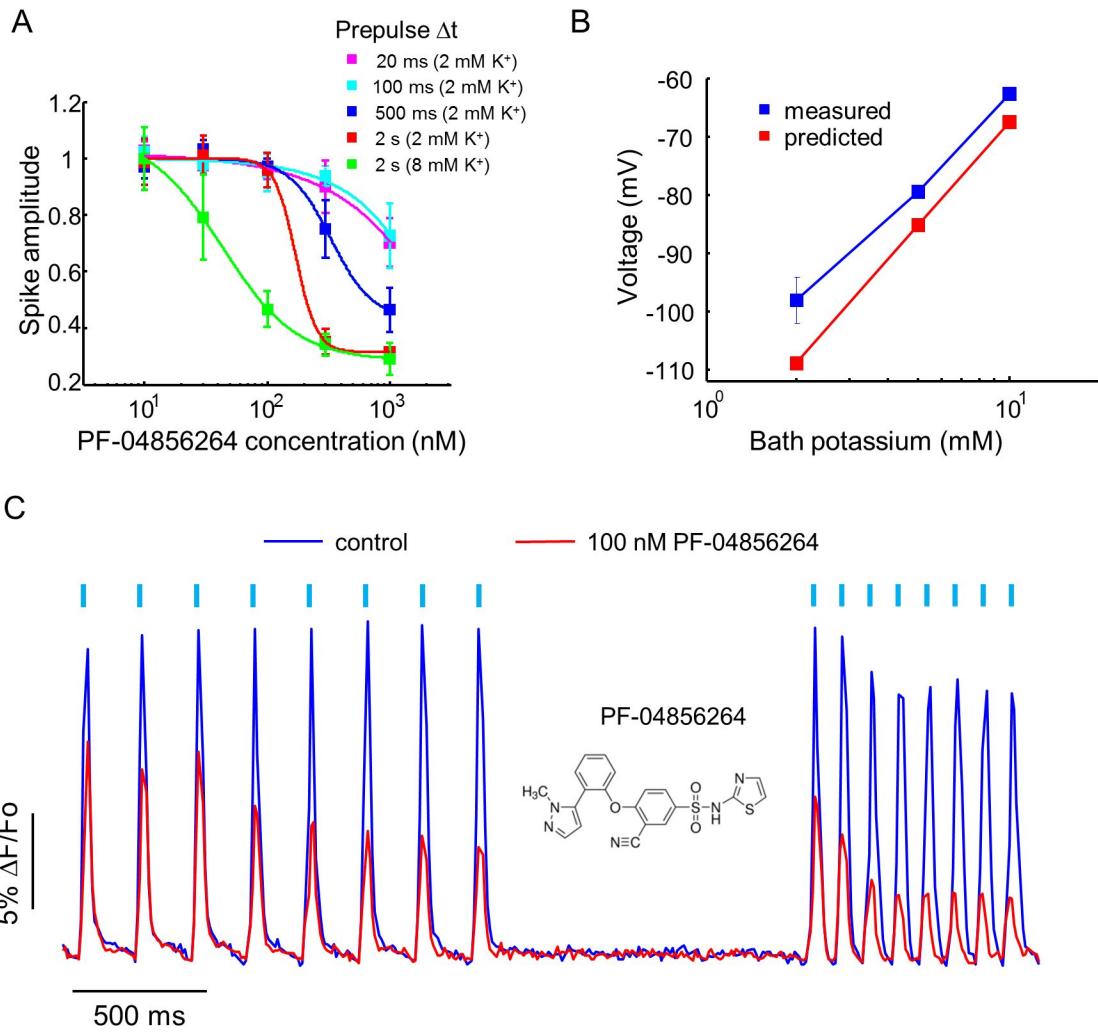
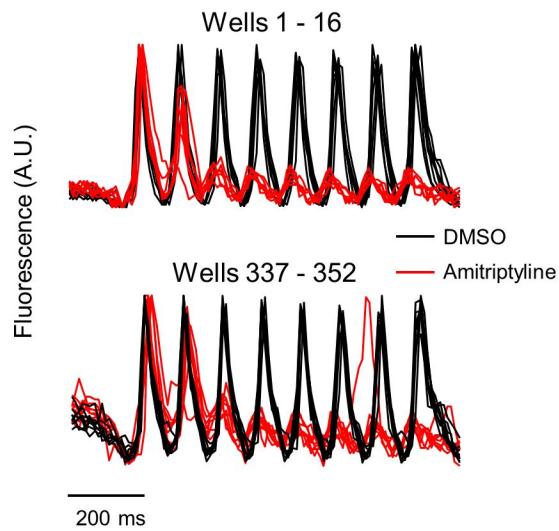
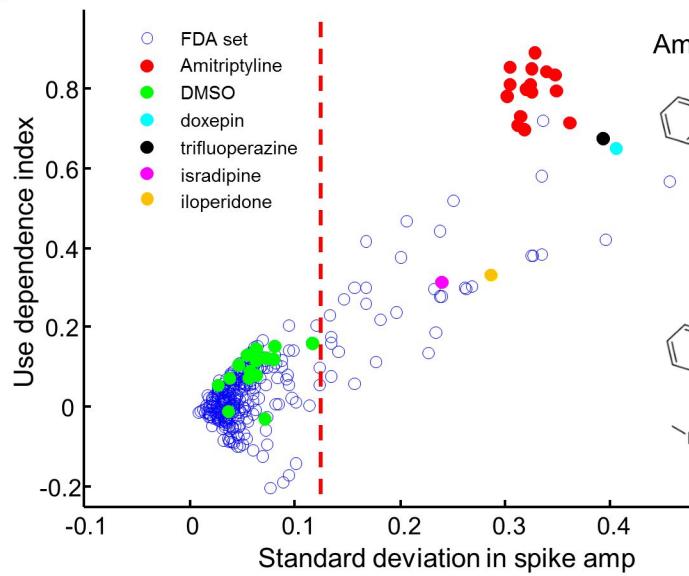


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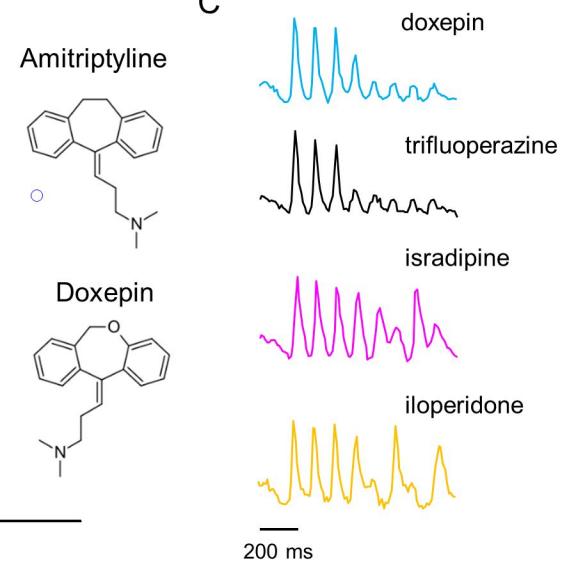
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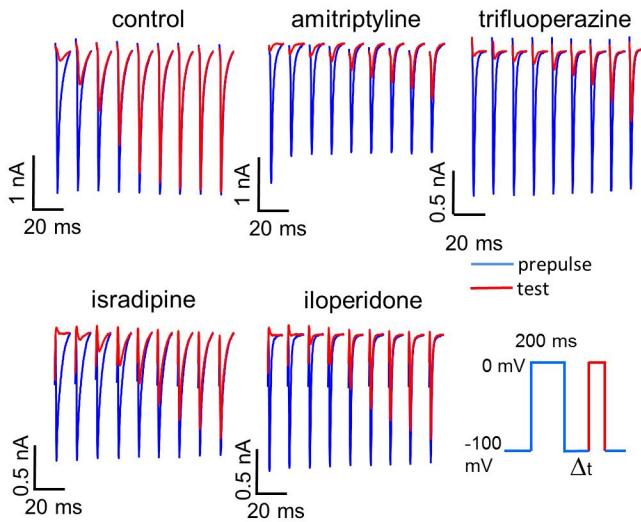
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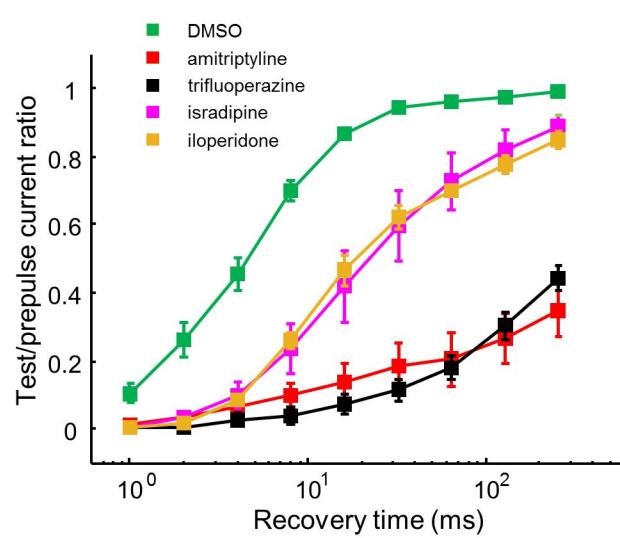
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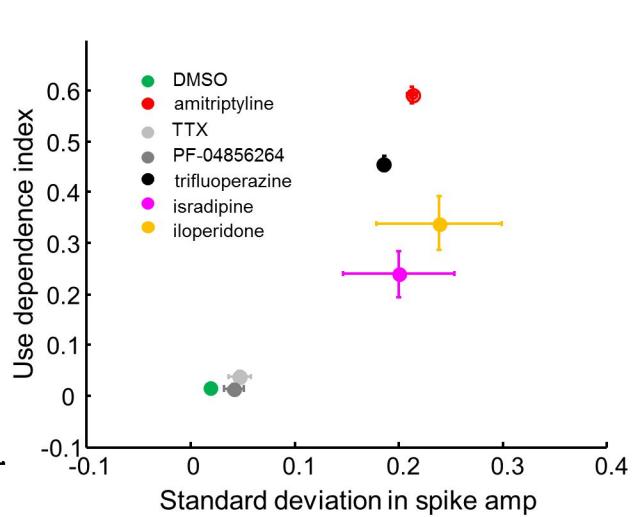


Figure 5

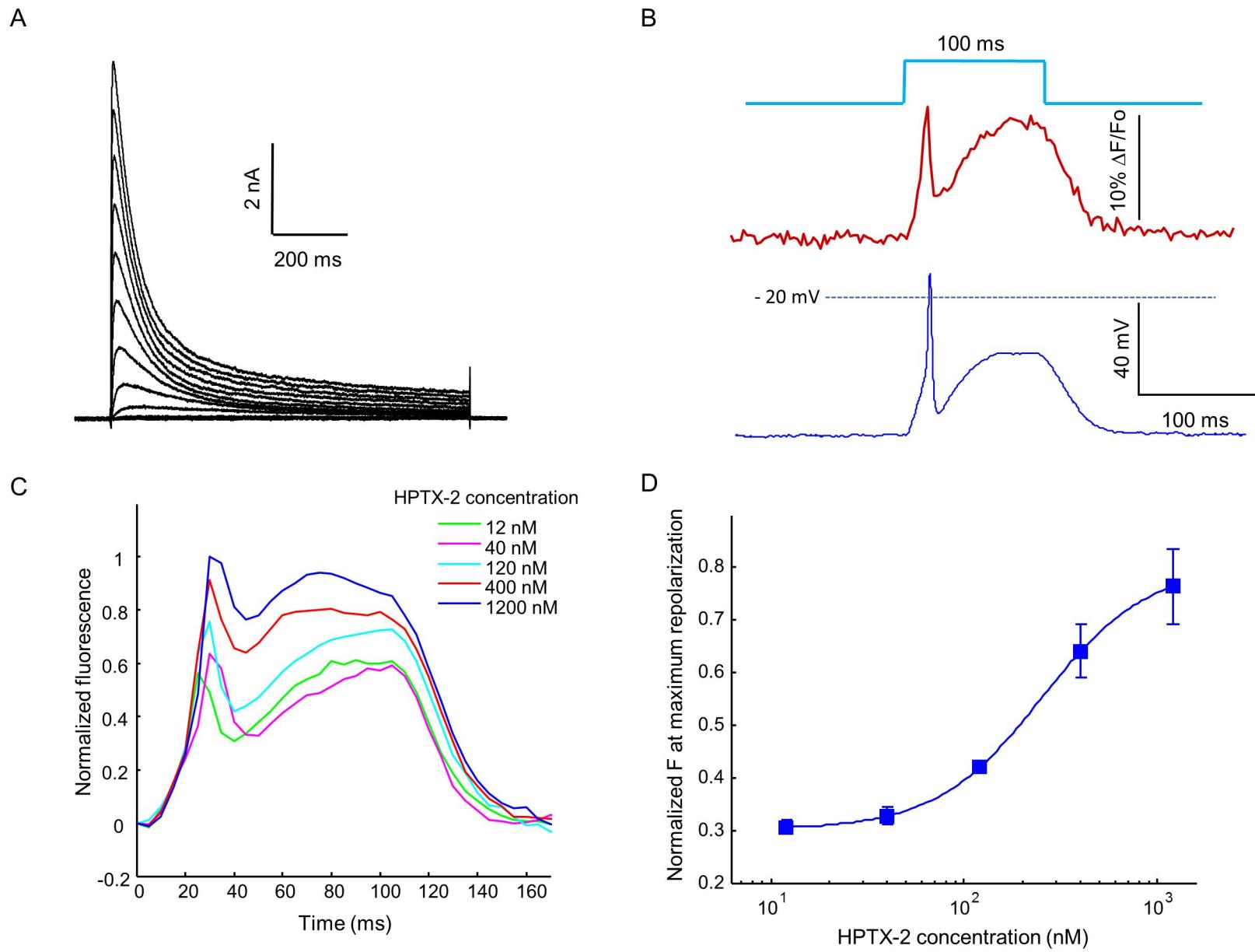


Figure 6