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## Transient, Consequential Increases in Extracellular Potassium Ions Accompany Channelrhodopsin2 Excitation

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### SUMMARY

Channelrhodopsin2 (ChR2) optogenetic excitation is widely used to study neurons, astrocytes, and circuits. Using complementary approaches *in situ* and *in vivo*, we found that ChR2 stimulation leads to significant transient elevation of extracellular potassium ions by ~5 mM. Such elevations

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#### AUTHOR CONTRIBUTIONS

The initial observations for this project were made by S.H. and B.S.K. Subsequently, J.C.O., M.R.G., S.H., D.T., and B.S.K. worked together to finish the experiments. Modeling was performed by S.L.A., T.H.R., T.M.H.-T., and J.R.K. with feedback from B.S.K. P.G. gave input on *in vivo* experiments. B.S.K. directed the project and the experiments. J.C.O., D.T., and M.R.G. helped write the methods. B.S.K. wrote the paper, and all authors commented.

#### DECLARATION OF INTERESTS

The authors declare no competing interests.

#### SUPPLEMENTAL INFORMATION

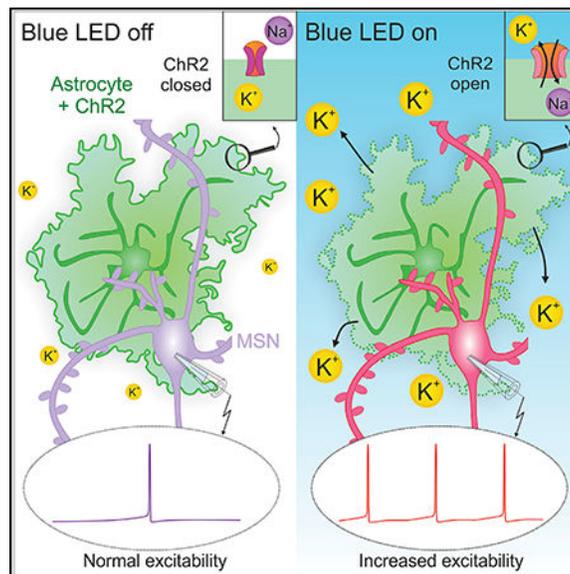
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were detected in ChR2-expressing mice, following local *in vivo* expression of ChR2(H134R) with adeno-associated viruses (AAVs), in different brain areas and when ChR2 was expressed in neurons or astrocytes. In particular, ChR2-mediated excitation of striatal astrocytes was sufficient to increase medium spiny neuron (MSN) excitability and immediate early gene expression. The effects on MSN excitability were recapitulated *in silico* with a computational MSN model and detected *in vivo* as increased action potential firing in awake, behaving mice. We show that transient, physiologically consequential increases in extracellular potassium ions accompany ChR2 optogenetic excitation. This coincidental effect may be important to consider during astrocyte studies employing ChR2 to interrogate neural circuits and animal behavior.

## In Brief

Using multiple approaches, Octeau et al. discover that optogenetic excitation of ChR2-expressing cells leads to significant transient extracellular potassium ion elevations that increase neuronal excitability and immediate early gene expression in neurons following *in vivo* stimulation.

## Graphical Abstract



## INTRODUCTION

The emergent field of optogenetics has revolutionized neuroscience since the landmark discovery that optical stimulation of channelrhodopsin2 (ChR2)-expressing cells elicits membrane potential depolarization (Nagel et al., 2003) and by the subsequent demonstration that such responses can be used to evoke action potential (AP) firing in neurons (Boyden et al., 2005). ChR2 and its variants are now widely used to explore neurons, glia, and multiple other cell types, permitting straightforward optical stimulation of genetically targeted cell populations and exploration of their contributions to neural circuit properties and animal behaviors (Bernstein and Boyden, 2011; Deisseroth, 2015).

The usual interpretation of neuronal studies that employ ChR2 is that selective, brief optical excitation of the targeted neurons causes sufficient membrane depolarization to engage endogenous voltage-gated ion channels and lead to AP firing in ChR2- expressing neurons, which then may cause ensuing changes in neural circuits controlling behavior. Thus, ChR2 provides an initial depolarizing inward current to bring the neurons to threshold. In this way, ChR2 has become a powerful tool for functional interrogation of neural circuits *in vivo* in a multitude of scenarios (Bernstein and Boyden, 2011; Deisseroth, 2015). Given its ease of use, ChR2-mediated excitation has also been used in a similar manner to explore the roles of astrocytes in regulating mouse behavior and circuits. For example, ChR2- driven astrocyte excitation has been used to implicate astrocytes in breathing (Gourine et al., 2010), response selectivity in the visual cortex (Perea et al., 2014), regulation of blood flow (Masamoto et al., 2015), sleep (Pelluru et al., 2016; Yamashita et al., 2014), and cerebellar motor function (Sasaki et al., 2012). It has also been used to regulate neuronal activity by engaging astrocyte mechanisms (Deemyad et al., 2018; Mederos et al., 2019; Shen et al., 2017) and to mimic the effects of neuronal activity on astrocyte glutamate transport (Gomez et al., 2019). However, astrocytes have few if any voltage-gated ion channels and display no propagated voltage signals. In the absence of a physiologically relevant endogenous voltage-dependent excitability mechanism to engage following brief ChR2 stimulation, several astrocyte studies have instead used quite long periods of ChR2 stimulation lasting from seconds to tens of seconds in order to observe subsequent effects on single neurons, circuits, and, in some cases, animal behavior. In some of these cases, ChR2-mediated excitation of astrocytes has suggested that astrocytes regulate circuits via gliotransmission, which was evoked by optical stimulation (Bang et al., 2016; Figueiredo et al., 2011).

We have been exploring astrocyte-neuron interactions in striatal microcircuits (Chai et al., 2017; Jiang et al., 2016; Octeau et al., 2018a; Srinivasan et al., 2016; Tong et al., 2014; Yu et al., 2018). As part of this, buoyed by the aforementioned studies, we used *Thy1-ChR2-YFP* mice (Arenkiel et al., 2007) to drive trains of APs in premotor cortical neurons and to study the consequences of this increased excitation upon striatal MSNs and astrocytes. However, during the experiments, we identified an unexpected slow inward current in both striatal medium spiny neurons (MSNs) and, surprisingly, in astrocytes. We describe the properties of the slow inward current and subsequent experiments that suggested that it was due to release of  $K^+$  into the extracellular space, which we carefully explored. The data reveal a potentially important consequence of ChR2-mediated optogenetic excitation, which is likely an important confounding feature for astrocyte studies employing prolonged ChR2 stimulation, especially for behavioral experiments where electrophysiological studies are not performed. Elevated  $K^+$  is expected to affect the physiology of not only neurons, but also diverse brain cell types (McDonough and Youn, 2017), as well as affect local blood flow and functional hyperemia (Girouard et al., 2010). These could influence the interpretation of, and possibly even cause, endpoint measures at different temporal scales, especially *in vivo*, in studies that employ ChR2 in astrocytes.

## RESULTS

### Transient Extracellular K<sup>+</sup> Elevations during Optical Stimulation in *Thy1-ChR2-YFP* Mice

ChR2 and its variants are the most commonly used tools for optogenetic excitation (Zhang et al., 2011). Immunohistochemistry (IHC) revealed strong expression of ChR2-YFP in *Thy1-ChR2-YFP* mice (Arenkiel et al., 2007) with strong expression in layer V cortical neurons (Figure 1A; n = 3 mice). As expected with the *Thy1* promoter, there was little expression in the striatum (Figure 1A), but at higher magnification, there was strong colocalization between ChR2-YFP and neurofilament (Nflm) in corticostriatal white matter tracts, but no significant colocalization between ChR2-YFP and neuronal nuclei (NeuN) or with the astrocyte marker S100 $\beta$  (Figures 1B and 1C; n = 12 images, n = 4 mice). Next, we recorded in whole-cell voltage-clamp mode from MSNs in sagittal brain slices and employed wide-field optical stimulation using parameters comparable to those used *in vivo* (Gibson et al., 2014). We used 473 nm light,  $\sim 0.02$  mW/mm<sup>2</sup> power, and 25 ms flashes at 20 Hz for 30 s and recorded optically evoked excitatory postsynaptic currents (EPSCs) in all MSNs from *Thy1-ChR2-YFP* mice, but not from wild-type (WT) mice (Figure 1D; n = 15 cells, n = 8 mice). The evoked EPSCs were accompanied by the development of a slow inward current (at  $-75.6$  mV after correcting for the tip potential) during the optical stimulation train (Figure 1D; see Figure S1 for expanded traces). This inward current peaked at  $-90 \pm 19$  pA, developed with a time constant ( $\tau$ ) of  $1.1 \pm 0.1$  s, was maintained during the stimulation train, and decayed with a  $\tau$  of  $3.7 \pm 0.2$  s (Figure 1D; n = 15 cells, n = 4 mice). Furthermore, recordings from striatal astrocytes revealed a similar inward current of  $-281 \pm 48$  pA and with rise and decay  $\tau$  values of  $1.0 \pm 0.1$  and  $4.3 \pm 0.2$  s, respectively (Figure 1E; n = 15 cells, n = 8 mice). The finding that optically evoked inward currents could be measured in striatal MSNs and astrocytes, even though neither cell type expressed ChR2-YFP (Figures 1A–1C), argues against photocurrents; the kinetics of the inward currents were also nearly three orders of magnitude slower than ChR2 activation and deactivation (Zhang et al., 2011). Additionally, the astrocyte responses argue against ionotropic glutamate receptor activation during corticostriatal axon stimulation (see later section) because such ion channels are unlikely to be expressed in striatal astrocytes (Chai et al., 2017). Moreover, the finding that the peak inward currents were larger in astrocytes than in MSNs suggests they may be due to elevation of K<sup>+</sup>. The astrocyte membrane has high resting K<sup>+</sup> conductance and is thus sensitive to extracellular K<sup>+</sup> increases.

We made K<sup>+</sup> selective microelectrodes to directly measure K<sup>+</sup> in the extracellular space of striatal slices before, during, and after optical stimulation in *Thy1-ChR2-YFP* mice. The calibrated electrodes displayed the expected  $\sim 54$  mV change for a 10-fold change in K<sup>+</sup> (Figure 1F; mean of 4 electrodes shown). Furthermore, single 25 ms flashes at a range of powers (0–0.09 mW/mm<sup>2</sup>) elicited clear, transient extracellular K<sup>+</sup> elevations in *Thy1-ChR2-YFP* mice, but not in WT mice (Figure 1G; n = 14, n = 4 mice). Such transients peaked and decayed over seconds, as expected for release and subsequent washout of K<sup>+</sup> in the extracellular space (Figure 1G). At power of 0.02 mW/mm<sup>2</sup>, 30 s trains of light flashes at 20 Hz elicited significant increases in extracellular K<sup>+</sup> that were maintained for the duration of the stimulus, with rise and decay  $\tau$  values of  $2.1 \pm 0.6$  and  $6.1 \pm 0.5$  s (Figure 1H; Table 2). The kinetics of the K<sup>+</sup> elevations were thus similar to the steady-state inward

currents recorded from MSNs and astrocytes (Figures 1D and 1E): the slightly slower time course measured with the electrodes may reflect sampling from the tip of the K<sup>+</sup> electrode, whereas the MSN and astrocyte recordings sampled within the tissue. At peak, the extracellular K<sup>+</sup> levels were elevated significantly to  $7.2 \pm 0.3$  mM from a bath concentration of 4.5 mM (Figure 1H). Furthermore, the change in K<sup>+</sup> evoked by a train of stimuli was not blocked by tetrodotoxin (TTX; 500 nM) or a combination of TTX, cyanquinoxaline (6-cyano-7-nitroquinoxaline-2,3-dione (CNQX; 10  $\mu$ M), and D 2-Amino-5-phosphonopentanoic acid (APV; 10  $\mu$ M) to block APs and ionotropic glutamate receptors (Figure 1I; n = 14–23 slices, n = 4 mice). Similar, albeit as expected smaller, changes were measured when the flash duration was reduced to 2 ms (Figures S2A–S2D; n = 14–23 slices, n = 4 mice, 20 Hz). We also performed a set of experiments in the cortex to ascertain if the K<sup>+</sup> elevations were a striatal specific effect. However, during optical stimulation, we measured slow-seconds time scale cortical pyramidal neuron depolarization (rise and decay  $0.3 \pm 0.1$  and  $6.8 \pm 0.7$  s, respectively) and significant-seconds time scale elevations in extracellular K<sup>+</sup> ( $8.7 \pm 0.5$  mM, with rise and decay  $2.8 \pm 0.2$  and  $9.1 \pm 1.1$  s, respectively) that were not blocked by TTX (Figures 1J and 1K; n = 16 slices, n = 4 mice). Responses were also observed with 2 ms flashes (Figures S2E and S2F; n = 12 slices, n = 4 mice).

### Transient Extracellular K<sup>+</sup> Elevations after AAV-Driven Expression of ChR2

Channelrhodopsin is frequently deployed using adeno-associated viruses (AAVs) for localized expression. We explored extracellular K<sup>+</sup> changes following local expression of the commonly used ChR2 variant, ChR2(H134R), in the striatum. Intra-striatal hSynapsin1-driven expression of ChR2(H134R)-YFP was robustly localized to NeuN positive neurons (Figures 2A and 2B), and optical stimulation with trains of increasing duration resulted in significant elevations of extracellular K<sup>+</sup> to  $8.8 \pm 0.8$  mM (Figure 2C; n = 16 slices, n = 4 mice). The rise and decay  $\tau$  values of these elevations were  $3.2 \pm 0.3$  and  $9.5 \pm 1.4$  s, respectively (Figures 2A–2C; n = 16 slices, n = 4 mice). Furthermore, the K<sup>+</sup> elevations were not significantly affected by TTX (Figure 2D; p > 0.05, n = 11–12 slices, n = 4 mice), showing they were not due to AP firing. Next, we expressed ChR2(H134R)-mCherry in striatal astrocytes using AAV2/5 and the *GfaABC1D* promoter (Haustein et al., 2014; Shigetomi et al., 2013): as expected, we detected no colocalization with NeuN positive neurons but detected significant colocalization with S100 $\beta$  positive astrocytes (Figures 2E and 2F). Optical stimulation with trains of increasing duration resulted in significant elevation of extracellular K<sup>+</sup> to  $7.4 \pm 0.2$  mM (Figure 2G; n = 16 slices, n = 4 mice). The rise and decay  $\tau$  values of such elevations were  $3.0 \pm 0.2$  and  $11.0 \pm 1.4$  s, respectively (Figures 2E–2G; n = 16 slices, n = 4 mice). Responses were also observed with 2 ms flashes (Figures S2G and S2H; n = 16 slices, n = 4 mice). Astrocytes are proposed to regulate extracellular K<sup>+</sup> dynamics via Ba<sup>2+</sup> sensitive Kir4.1 ion channels (Nwaobi et al., 2016). Ba<sup>2+</sup> block (100  $\mu$ M) of Kir4.1 did not significantly alter the peak change in K<sup>+</sup> evoked by ChR2(H134R) activation; however, suggesting a possible role for Kir4.1 in K<sup>+</sup> clearance, Ba<sup>2+</sup> significantly increased the rise and decay times of the extracellular K<sup>+</sup> elevations (Figure 2H; n = 16 slices, n = 4 mice). To explore the possibility that the ChR2-mediated extracellular K<sup>+</sup> elevations may be due to brain slicing, we repeated the experiments following expression of ChR2(H134R) in cortical astrocytes *in vivo* (Figure 2I). We measured significant, albeit smaller, slow K<sup>+</sup> elevations in the cortex *in vivo* of awake head-fixed mice (rise and decay

times of  $0.5 \pm 0.4$  and  $1.4 \pm 0.7$  s, respectively), but not in mice that received a control AAV (Figure 2J;  $n = 4$  mice each). Finally, we compared the  $K^+$  elevations measured at room temperature and at more physiological temperatures ( $34^\circ\text{C}$ ): we found significant elevations at both temperatures and only subtle effects of temperature on the kinetics of extracellular  $K^+$  (Table 1; Figure S4).

### ChR2-Mediated Extracellular $K^+$ Elevations Increased MSN Excitability in Brain Slices

The finding that ChR2(H134R) activation in astrocytes caused an increase in extracellular  $K^+$  ( $\sim 7.4$  mM) comparable to that caused by neuronal ChR2 activation ( $\sim 8.8$  mM; Figures 2C and 2G) provided an opportunity to evaluate if such  $K^+$  fluxes were consequential for neuronal function. To this end, we expressed ChR2(H134R)-mCherry in adult striatal astrocytes using AAV2/5 and the *GfaABC1D* promoter (Haustein et al., 2014; Shigetomi et al., 2013). We confirmed in these specific experiments that ChR2(H134R)-mCherry was expressed in  $\sim 80\%$  of striatal S100 $\beta$  positive astrocytes, but not in striatal NeuN positive neurons (Figures 3A and 3B;  $n = 12$  sections,  $n = 4$  mice). In this experimental approach, we identified and recorded from MSNs but optically stimulated ChR2(H134R)-mCherry expressing astrocytes nearby in the dorsolateral striatum. A single brief train of ten 2 ms flashes significantly and slowly depolarized MSNs by  $\sim 2$  mV (Figure 3D;  $n = 14$  MSNs,  $n = 4$  mice). A longer 30 s train of flashes depolarized MSNs by  $\sim 7$  mV for the duration of the stimulation train (Figure 3D;  $n = 14$  MSNs,  $n = 4$  mice). The depolarization also peaked and decayed with seconds time scale kinetics similar to those observed for extracellular  $K^+$  elevations (Figures 2G and 3D). Some past studies have suggested that ChR2 stimulation of astrocytes evokes astrocyte glutamate release, which is proposed to be a gliotransmitter acting on neurons. However, we detected no change in excitatory postsynaptic potentials (EPSPs) arriving onto MSNs during optical stimulation of astrocytes, and of most relevance here, the change in MSN membrane potential was not affected by the bath application of CNQX and APV ( $10 \mu\text{M}$ ), which completely blocked all EPSPs (Figure 3D). These data suggest that ChR2-mediated stimulation of striatal astrocytes depolarized MSNs significantly, likely by  $K^+$  elevations and not via glutamate release (Chai et al., 2017; see also Nedergaard and Verkhratsky, 2012 and Sloan and Barres, 2014). To explore neuronal consequences more carefully in the striatum, we applied slow ramps to determine the amount of current required to trigger APs in MSNs (rheobase) before, during, and after optical stimulation of ChR2-expressing astrocytes (Figure 3E). In all cases, we measured significant MSN membrane potential depolarization and a significant drop in the rheobase (Figure 3E;  $n = 14$  MSNs,  $n = 4$  mice). Furthermore, following *in vivo* stimulation of astrocytes expressing ChR2(H134R), we detected a significant increase in the expression of the immediate early gene, cFos, in 20% of DAPI-positive cells in the striatum (Figures 3F–3H;  $n = 7$  slices,  $n = 3$  mice), which was significantly greater than that observed in mice expressing tdTomato in astrocytes (Figures 3G and 3H). Of the cells with elevated cFos expression following *in vivo* astrocyte optogenetic stimulation,  $\sim 35\%$  were S100 $\beta$  positive astrocytes (Figures 3F and 3H), whereas  $\sim 18\%$  were NeuN positive neurons (Figures 3G and 3H). These data are consistent with the interpretation that optogenetic excitation of ChR2(H134R)-expressing astrocytes *in vivo* was sufficient to alter astrocyte and neuronal excitability, resulting in significantly elevated cFos expression in a proportion of the cells. During our experiments, we observed no overt change in open-field activity or general

behavior during optical stimulation of ChR2(H134R)-expressing astrocytes relative to control mice (Figure S3).

### Computational Modeling: MSN Excitability Increase Recapitulated by K<sup>+</sup> Elevation Alone

We adapted a MSN model (Mahon et al., 2000) (Figure 4A; Table 2), which comprised a single compartment representing a MSN soma and proximal dendrites, and eight active Hodgkin-Huxley (HH)-based models of variable membrane conductances. The modeled currents are persistent sodium ( $I_{NaP}$ ); slowly inactivating sodium ( $I_{NaS}$ ); transient sodium ( $I_{NaT}$ ); delayed rectifier potassium ( $I_{KDR}$ ); inwardly rectifying potassium ( $I_{KIR}$ ); persistent potassium ( $I_{KRP}$ ); fast inactivating A-type potassium ( $I_{KAf}$ ); slow inactivating A-type potassium ( $I_{KAs}$ ); and a lumped, non-specific ion leak current. We replaced the non-specific leak current with three Goldman-Hodgkin-Katz (GHK)-based specific ion leaks ( $Cl^-$ ,  $Na^+$ , and  $K^+$ ). We used the model to explore how  $K^+$  concentration changes measured in the experiments reported in Figure 3 affect MSN excitability *in silico*. We adjusted the model's parameters to replicate basal MSN spiking features (Mahon et al., 2000) (see STAR Methods). We then varied conductance and permeability parameters to constrain a population of MSN models to reproduce the observed ranges of resting membrane potential values and resting levels of excitability during light off and light on conditions. We required that each model reproduce these measures at both 4.5 and 7.5 mM extracellular  $K^+$  to be included in the population. By asking if models exist sensitive to  $K^+$  changes within experimentally observed ranges from Figures 1 and 2, we aimed to test if the modeled range of extracellular  $K^+$  was sufficient to explain the data. Current injection ramp protocols were replicated for each model instance using two extracellular  $K^+$  concentration parameters related to Figure 3 (i.e., 4.5 and 7.5 mM). A model instance comprised a unique set of ionic conductances and specific leak permeability parameters. Using a modified version of the parameter optimization via evolutionary algorithms approach (Rumbell et al., 2016) (see STAR Methods), we identified a population of model instances that reproduced the range of excitability features measured before and after optical stimulation of astrocytes expressing ChR2 (Figure 3) when the extracellular  $K^+$  increased from 4.5 to 7.5 mM  $K^+$  (Figure 4B), respectively. The resulting population of model instances recapitulates a range of excitability features measured empirically, which are modulated consistently in the same direction by extracellular  $K^+$  across all model instances, thus establishing extracellular  $K^+$  as an independent parametric change in the model sufficient to explain changes in MSN excitability observed during ChR2 excitation (Figure 4). We further optimized these models, constraining with specific spiking features from one of the experimental traces. One such outcome generated the waveforms depicted in Figure 4C, which closely recalls experimental data of Figure 3. To explore the full range of extracellular  $K^+$  concentrations, we incrementally varied  $K^+$  in the model used to generate Figure 4C. Increases in extracellular  $K^+$  depolarized MSNs and decreased rheobase in the model (Figure 4D).  $K^+$  above 9 mM caused seizure-like bursting activity in the MSN model; to generate plots in Figure 4D, we thus did not exceed 7.5 mM  $K^+$ . Overall, this MSN model supports the hypothesis that extracellular  $K^+$  elevations are sufficient per se to reproduce the increased excitability changes measured during experiments in Figure 3 without the need to invoke additional factors.

## ChR2-Mediated Extracellular K<sup>+</sup> Elevations Increased MSN Excitability *In Vivo*

Experiments reported in the preceding sections show that optical stimulation of ChR2-expressing astrocytes causes extracellular K<sup>+</sup> increases (Figure 2) and MSN depolarization (Figure 3D), as well as increased MSN excitability in brain slices (Figure 3E) and *in silico* (Figure 4). In order to determine whether such responses translated into increased MSN AP firing *in vivo*, we recorded MSN activity in awake behaving mice (Figure 5) using multisite silicon probes (Bakhurin et al., 2016). ChR2 was selectively expressed in striatal astrocytes using approaches described in the preceding sections. Silicon probes with an inbuilt light-emitting diode (LED) were used for optical excitation and electrophysiological recording (Figure 5A) in awake behaving mice on a spherical treadmill. MSN APs were analyzed offline, sorted using established criteria (Bakhurin et al., 2016) and AP frequency assessed before, during, and after optical stimulation (Figure 5A). *Post hoc*, we verified in each mouse that AAV-driven ChR2 expression was overlapping anatomically with the site at which the silicon probes were placed for electrophysiological assessments (Figure 5B). As a control group, we used mice in which tdTomato was identically expressed in astrocytes: MSN APs from tdTomato and ChR2 groups were identified and recorded (Figure 5C). Figure 5D shows AP firing of 60–70 MSNs before, during, and after optical stimulation for the control tdTomato and the ChR2 groups. From the raster plots, it was clear that optical stimulation did not evoke alterations in MSN AP firing in the control group, but optical stimulation evoked clear increases in AP firing in the ChR2 group (Figure 5D; n = 5 mice in each group). These data are summarized as average firing over time (Figure 5E) or as histograms before, during, and after optical stimulation (Figure 5F). Together with the studies of extracellular K<sup>+</sup> increases, MSN excitability, and computational modeling, these *in vivo* recordings strongly indicate that optical stimulation of ChR2-expressing astrocytes can cause significantly elevated MSN firing. Our brain slice data and modeling suggest that astrocyte ChR2-mediated K<sup>+</sup> elevations are the underlying cause.

## DISCUSSION

We show that transient extracellular K<sup>+</sup> elevations accompany optogenetic excitation with two widely used variants of ChR2 (wild-type and H134R) in a common transgenic mouse model, following AAV-mediated delivery in brain slices and *in vivo*, regardless of whether the ChR2 is expressed in neurons or astrocytes. The simplest interpretation is that K<sup>+</sup> flows down its electrochemical gradient through the ChR2 pore from the inside to the outside of cells. The ChR2 pore displays a K<sup>+</sup> to Na<sup>+</sup> relative permeability of ~0.4 (Lin et al., 2009; Nagel et al., 2003) and a minimal open pore diameter of ~6.2 Å (Richards and Dempksi, 2012). Thus, opening of the ChR2 pore carries inward Na<sup>+</sup> current and outward K<sup>+</sup> current under physiological conditions as determined by the respective ion equilibrium potentials relative to the cell's membrane potential. Our data show that this outward K<sup>+</sup> flux is significant and elevates K<sup>+</sup> concentrations in the extracellular space to consequential levels during seconds-long optical stimulations usually employed in astrocyte studies.

Interestingly, our observations recall work at the neuromuscular junction where a slow endplate potential due to K<sup>+</sup> release directly by nerve terminal APs was detected (Katz and Miledi, 1982). Several other papers have similarly documented K<sup>+</sup> elevations during

neuronal activity (Heinemann et al., 1990; McCreery and Agnew, 1983; Walton and Chesler, 1988). However, the  $K^+$  elevations we report are not due to  $K^+$  release during the AP because they occurred when APs were blocked and when ChR2 was expressed in astrocytes devoid of APs. Furthermore, although  $K^+$  is expected to flow through open endogenously expressed voltage-dependent  $K^+$  channels that open during ChR2-mediated depolarization, this cannot account for our observations: astrocytes have little if any rectification indicative of voltage-gated  $K^+$  conductance, and owing to their low membrane resistance and high resting ohmic  $K^+$  conductance, they rarely deviate from near the  $K^+$  equilibrium potential. In light of these considerations, we discuss the relevance of ChR2-mediated  $K^+$  fluxes in relation to the diverse uses of optogenetic excitation strategies, but with a focus on astrocyte studies.

A logical interpretation of our astrocyte studies is that optogenetic stimulation increased neuronal excitability via extracellular  $K^+$  elevations, which was recapitulated in a computational model. Our studies should not be overinterpreted, however, as indicating that we dismiss gliotransmission (Araque et al., 2014), but our studies highlight an important consideration with the use of ChR2 in these regards. It will be important to candidly consider if cellular and behavioral responses ascribed to ChR2-evoked gliotransmission onto neurons (reviewed by Bang et al., 2016) could instead reflect elevations of extracellular  $K^+$ . Even subtle changes in  $K^+$  will change neuronal excitability directly (Hille, 2001) and indirectly by reducing glutamate clearance by electrogenic transporters (Barbour et al., 1988; Szatkowski et al., 1990). This will lead to elevated ambient glutamate levels, which could conceivably be interpreted as glutamate gliotransmission. If so, the relevance of this phenomenon to circuits and behaviors may have been misinterpreted. Furthermore, strong excitation of astrocytes with ChR2 may change extracellular electric fields, which may affect neuronal ephaptic signaling (Jefferys, 1995). The use of optogenetic excitation approaches to study astrocytes, therefore, requires careful consideration not least because these cells are largely electrically silent (Kuffler, 1967). It is not immediately clear why artificially changing membrane potential is appropriate to explore astrocyte physiology, although such approaches are obviously meaningful for neurons.

Our goals in this study were not to identify a past behavioral study based on ChR2 and highlight it as being better explained by transient  $K^+$  elevations. Instead, we followed empirical observations that show  $K^+$  elevations occur, and we suggest this should be considered in the interpretation of future studies using ChR2, especially in astrocytes. We also did not perform detailed behavior analysis, but during our experiments, we observed no overt change in open-field activity or general behavior during optical stimulation of ChR2(H134R)-expressing striatal astrocytes relative to control mice. This is unsurprising because astrocytes do not have axonal projections and do not discriminate between D1 and D2 MSNs. Thus, we suggest that the elevation of  $K^+$  would not discriminate between D1 and D2 MSNs, both of which exist within single astrocyte territories (Chai et al., 2017; Octeau et al., 2018a) and may drive compensatory signals in relation to simple behaviors. Instead, we interpret the *in vivo* data to show that elevated cFos responses most likely reflect neuronal firing caused by transient  $K^+$  elevations, a finding confirmed by recording neuronal activity *in vivo*.

It is problematic to search for a specific striatal microcircuit-related behavior that may be caused by astrocyte-mediated  $K^+$  increases because unlike a neurotransmitter,  $K^+$  will affect the functions of all cells, not just neurons. Instead, we explored the effect of astrocyte ChR2-mediated  $K^+$  elevations on the behavior of single neurons using modeling and experiments. These data strongly suggest that ChR2-mediated optical excitation of astrocytes can affect the excitability of neurons via  $K^+$  elevations. Since  $K^+$  is tightly regulated, even small changes are expected to affect neurons, astrocytes, microglia, and blood vessels (McDonough and Youn, 2017). For example,  $K^+$  is potently ictogenic at around 9 mM (de Curtis et al., 2018), has vasoconstrictor and vasodilator actions on brain vasculature (Girouard et al., 2010), contributes to neurological disease phenotypes (Nwaobi et al., 2016; Tong et al., 2014), and will synchronize neuronal activity (Jefferys, 1995). Subtle changes in  $K^+$  will depolarize neurons and alter the open and closed probabilities of the activation and inactivation gates of voltage-gated  $K^+$  and  $Na^+$  channels, which of course will alter AP properties through Hodgkin-Huxley outcomes (Hodgkin and Huxley, 1952). Such effects may explain why ChR2-mediated stimulation of astrocytes silenced neurons in the subthalamic nucleus in pioneering early studies (Gradinaru et al., 2009). Hence, long-term optogenetic stimulation *in vivo* may cause complex effects on cells in parallel with the effects desired by expression of ChR2 in specific cell types.  $K^+$  changes may also engage local circuits that modulate behavior through ionic or ephaptic mechanisms (Jefferys, 1995) in addition to the pathways targeted by ChR2 expression. These are relevant considerations with the use of optogenetic strategies *in vivo* and may predict the emergence of unrelated, possibly pathogenic, effects during use. Additional consequences of the larger and more compartmentalized intracellular  $Na^+$  fluxes also merit consideration. Perhaps, infrared induced electro-mechanical effects (Plaksin et al., 2018) may provide opportunities for remote neuronal excitation (Shapiro et al., 2012) without reliance on substantial ionic currents. In the case of astrocytes, new tools are clearly needed to remotely manipulate physiological levels of activity.

It would be egregiously misleading of us to suggest that transient  $K^+$  elevations markedly contribute to most optogenetic excitation studies. This is not our intention. More parsimoniously, we suggest that the impact of extracellular  $K^+$  dynamics will depend on the membrane properties of the neurons in question (e.g., their input resistance, levels of open  $K^+$  channels, and rectification), as well as the activation and inactivation thresholds for voltage-gated ion channels. Local neuroanatomy will also be relevant, as this will determine how  $K^+$  elevations are dissipated in the extracellular milieu and will affect local electric fields. Overall, our insights echo caution by others emphasizing effects of light itself (Rungta et al., 2017) and the importance of depolarization block (Herman et al., 2014) and of circuit-related off-target effects in the case of silencing approaches (Otchy et al., 2015; Südhof, 2015). Admittedly, our work may have little relevance to the use of ChR2 in neuronal studies where optical stimuli are usually brief, but we suggest transient  $K^+$  increases are likely to be markedly consequential in astrocyte and behavioral studies when optical stimuli are typically prolonged. The impact of such changes may have been missed when independent electrophysiological evaluations were not employed to measure cellular level electrical changes and extracellular  $K^+$  levels. We suggest that cellular neurophysiology is necessary in parallel with *in vivo* studies in order to make adequate

interpretation of neural circuit and behavioral outcomes in studies employing Chr2 and other related optogenetic strategies.

## STAR★METHODS

### CONTACT FOR REAGENT AND RESOURCE SHARING

Requests for resources and reagents should be directed to and will be fulfilled by the Lead Contact, Baljit S. Khakh (bkhakh@mednet.ucla.edu). Adeno-associated viruses and plasmids are, or will, be made available on Addgene or through the University of Pennsylvania Vector Core.

### EXPERIMENTAL MODEL AND SUBJECT DETAILS

All animal experiments were conducted in accordance with the National Institute of Health Guide for the Care and Use of Laboratory Animals and were approved by the Chancellor's Animal Research Committee at the University of California Los Angeles. Mice were housed with food and water available *ad libitum* in a 12-hour light/dark environment. Mice were healthy with no abnormal behavioral phenotypes and were not involved in prior studies. For all experiments, mice were sacrificed during the light cycle. Both male and female mice were used for experiments (between 8–16 weeks of age).

### METHODS DETAILS

**Mouse lines**—Most experiments were conducted on C57BL/6N mice (from Taconic) from an in-house colony. Heterozygous *Thy1-ChR2-YFP* mice were obtained from Jackson Laboratories (strain B6.Cg-Tg(Thy1-COP4/EYFP)9Gfng/J; stock #007615) and have been published by others (Arenkiel et al., 2007). *Thy1-ChR2* mice were maintained as heterozygous  $\times$  wt breeding pairs in an in-house colony. For all experiments, wt mice were C57BL/6N.

**Plasmid and adeno-associated viruses (AAV)**—All plasmid constructs were generated using standard molecular biology techniques and the In-Fusion HD Cloning Kit (Clontech). All constructs were sequenced before AAV generation. For all constructs, we modified plasmid pZac2.1 *GfaABC<sub>1</sub>D*Lck-GCaMP6f (Addgene plasmid # 52924). Briefly, we generated adeno-associated virus plasmids (AAV2/5) capable of expressing the selected cDNA using the following strategy. For the plasmid, *GfaABC<sub>1</sub>D*-ChR2(H134R)-mCherry, we removed Lck-GCaMP6f using *Xho1* and *Xba1* restriction enzymes and infused the ChR2-(H134R)-mCherry cDNA. The fully sequenced “pZac2.1” plasmids were sent to the Penn Vector Core, which used them to generate AAV serotype 2/5 for this construct yielding a concentration of  $\sim 1 \times 10^{13}$  genome copies/ml (gc/ml). This construct has been deposited at Addgene in the Khakh lab repository, and the AAVs are available from the UPenn Vector Core Catalog.

**In vivo microinjections of AAVs**—Male and female P42-P56 C57BL/6N mice were used in all AAV experiments. Briefly, animals were deeply anesthetized using isoflurane (induction at 5%, maintenance at 1%–2.5% v/v). After induction of anesthesia, the mouse's head was carefully placed into a stereotaxic frame and gently secured by blunt ear bars and

their nose was placed into an anesthesia system (David Kopf Instruments). Before the surgery, mice were administered 0.05 mL of buprenorphine (Buprenex, 0.1 mg/ml) subcutaneously. 10% povidone iodine and 70% ethanol were used to sterilize the incision site. Short (5–8 mm) skin incisions were made, followed by craniotomies ~2 mm in diameter using a small steel burr (Fine Science Tools) powered by a highspeed drill (K.1070; Foredom) directly above the site of the injection. Sterile saline was applied onto the skull to lessen heating caused by drilling. Viral injections were performed using a stereotaxic apparatus (David Kopf Instruments) which guided the placement of angled bevel glass pipettes (1B100 – 4; World Precision Instruments) into the appropriate brain region. Our stereotaxic coordinates (Axes: Anterior-Posterior / Medial-Lateral / Dorsal-Central; relative to Bregma) for AAV microinjection were as follows: dorsolateral striatum: +0.8 / +2.0 / –2.4 and superficial motor cortex: –0.1 / +0.7 / –0.1. In this study, the following viruses and their titers were used: AAV2/5 *GfaABC1D*-ChR2(H134R)-mCherry ( $8.1 \times 10^{12}$  gc/ml), AAV2/5 *hSynapsin1*-ChR2(H134R)-YFP ( $2.9 \times 10^{12}$  gc/ml, Lot# AV4319J, UNC GTC Vector Core). AAVs were injected by using a syringe pump at 200 nl/min (Pump11 PicoPlus Elite; Harvard Apparatus). The glass needles were left in place for at least 6 min and slowly removed over the course of 1 minute. The incisions were closed with single external 6 – 0 nylon sutures. After surgery, animals recovered overnight in cages which were placed halfway on a low-voltage heating pad. As an analgesic, subcutaneous buprenorphine was given twice daily for 48 hours following surgery. In addition, trimethoprim/sulfadiazine diet (275 & 1,365 ppm, TD.06596, Teklad) was fed *ad lib* prophylactically for one week after surgery. Following AAV injection and recovery, the mice were sacrificed 22–28 days later for all experiments.

**Acute brain slice preparation and single cell recordings**—Acute brain slices were prepared from P42–70 wt, *Thy1*-ChR2 mice or AAV microinjected mice. Briefly, animals were deeply anesthetized and decapitated. The brains were placed in ice-cold modified artificial CSF (aCSF, containing in mM: 194 sucrose, 30 NaCl, 4.5 KCl, 1 MgCl<sub>2</sub>, 26 NaHCO<sub>3</sub>, 1.2 NaH<sub>2</sub>PO<sub>4</sub>, and 10 D-glucose) and cut into 300  $\mu$ m-thick coronal or parasagittal slices containing areas of the striatum and cortex. Brain slices equilibrated for ~30 min at 32–34°C in normal aCSF (containing in mM; 124 NaCl, 4.5 KCl, 2 CaCl<sub>2</sub>, 1 MgCl<sub>2</sub>, 26 NaHCO<sub>3</sub>, 1.2 NaH<sub>2</sub>PO<sub>4</sub>, and 10 D-glucose) which was continuously bubbled with a mixture of 95% O<sub>2</sub>/5% CO<sub>2</sub>, stored at room temperature in the same buffer. All slices were used for experiments within ~6 hr of animal death. Cells were visualized using IR-DIC optics on an upright microscope (BX61WI, Olympus). pCLAMP10 software and a MultiClamp 700B amplifier and Digidata 1322A was used for electrophysiology (Axon Instruments). For all recordings we used normal aCSF at 20–25°C. For patch recordings, the intracellular solution contained the following (in mM): 135 potassium gluconate, 3 KCl, 0.1 CaCl<sub>2</sub>, 10 HEPES, 1 EGTA, 8 Na<sub>2</sub>-phosphocreatine, 4 Mg-ATP, 0.3 Na<sub>2</sub>-GTP, pH 7.3 adjusted with KOH. The initial access resistances were < 20 M $\Omega$  for all cells; if this changed by > 20% the cell was discarded.

**Recording K<sup>+</sup> ion dynamics in slices and *in vivo***—To record slice K<sup>+</sup> dynamics, we made K<sup>+</sup> selective microelectrodes (Octeau et al., 2018b). Briefly, borosilicate glass capillaries were pulled to a fine tip of ~2–5  $\mu$ m and then baked at ~200°C in the presence of

5% dichlorodimethylsilane for 1 hour up to overnight to enhance the hydrophobicity of the glass by silanization. The silanized capillaries were then allowed to cool to room temperature and then broken to a diameter of approximately 15  $\mu\text{m}$  using a blunt instrument. The capillaries were then backfilled with a solution containing 300 mM NaCl and 10mM HEPES, pH 7.4. The microelectrode tip was then immersed in a small volume ( $\sim 1 \mu\text{l}$ ) of potassium ionophore cocktail (5% w/v valinomycin, 93% v/v 1,2-dimethyl-3-nitrobenzene, 2% w/v potassium tetrakis(4-chlorophenyl)borate) and this solution was drawn approximately 1 mm into the tip by capillary action. The primed microelectrode was then fixed into an electrode holder on the connected to an amplifier and digitizer which transmitted voltage responses to a PC recording the responses in pCLAMP10. The voltage responsiveness of all microelectrodes were calibrated with solutions of aCSF which had various concentrations (in mM 0.1, 1.0, 10 and 100) of KCl, for these solutions iso-osmolar quantities of NaCl were subtracted to maintain total solution osmolarity at  $\sim 310$  mOsm. The  $\text{K}^+$ -selective microelectrodes were then slowly inserted into prepared brain slices which were superfused with 4.5 mM  $\text{K}^+$  aCSF and the voltage responses to blue LED flashes were measured. For *in vivo* measurement of  $\text{K}^+$  ion dynamics mice were prepared with a cranial bar fixed to the skull to maintain brain position. Mice were anesthetized with isoflurane (3%–5% induction, 1.5% maintenance) ten minutes after injection of a systemic analgesic (carprofen, 5 mg per kg of body weight) and placed in a stereotaxic frame. During surgery, mice were always kept at 37°C using a feedback-controlled heating pad. Pressure points and incision sites were injected with lidocaine (2%), and eyes were protected from desiccation using artificial tear ointment. The skin above the skull was cut and a custom-made lightweight metal head holder was implanted on the skull using Vetbond (3M) and a recording chamber was built using dental cement (Ortho-Jet, Lang). Access to the brain was closed using a thin layer ( $\sim 2$  mm) fast drying silicone elastomer (Kwik-sil, WPI) and a top layer of dental cement. Mice had a recovery period from surgery of five days, during which they were administered amoxicillin (0.25 mg per ml in drinking water through the water supply). After the recovery period, mice were habituated to head fixation on the spherical treadmill. 20–25 days following viral injection mice were placed onto the spherical treadmill and maintained in a head fixed position using a metal arm to grip the head holder attached to the mouse's skull. Following head restriction, the dental cement and silicone covering was removed and ACSF was applied to the surface of the brain to prevent tissue drying. To assess potassium ion changes a  $\text{K}^+$  selective microelectrode was prepared, calibrated and then lowered  $\sim 50$  microns perpendicularly into the cortical surface. Electrode potentials were allowed to stabilize over the course of 8–10 minutes before recording. Light flashes from a LED-Blue optogenetics light source were delivered via a fiber which was controlled remotely using a TTL pulse from Pulser software version 2.3.1 (Prizmatix, Israel). Electrode responses were captured and recorded using an Axopatch 700B amplifier connected via a digitizer to a computer with pCLAMP9 software. After completion of the experiments, the mice were euthanized by pentobarbital injection and immunostaining was performed on the brains to confirm viral expression.

***In vivo* Chr2 activation and behavioral assessment**—To deliver blue light to the striatum we constructed a fiber optic cannula using a DIY-cannula kit (Prizmatix). The fiber optic was cleaved into the desired length, which was approximately 10 mm. A small,  $\sim 1 \mu\text{L}$

droplet of epoxy resin was applied to the flat opening of a 6.5 mm long cannula. The fiber was then inserted through the epoxy bead into the cannula until it protruded approximately 1 mm from the opposite end of the cannula. Using a heat gun, we then cured the epoxy resin and secured the fiber in place. When the epoxy cooled, the fiber on the convex end of cannula was made transparent using polishing paper of increasing grits; from 4500 to 60000 final grit strength. Cannulated fibers were tested for their integrity and maximum light output (8 mW) before being used for the implantation. To implant the cannulated fiber optic, mice were first anesthetized with isoflurane (induction at 5%, maintenance at 1%–3%). The scalp was shaved and animals were gently fitted into a stereotaxic frame with their heads secured by ear bars. Ophthalmic ointment was applied to eyes to prevent dryness and postoperative discomfort. The surgical area was sterilized thrice with 10% povidone iodine and 70% isopropanol. A skin incision was made on top of the head and the periosteum was removed by swiping with a cotton swap wetted with 3% hydrogen peroxide. A mark was made on the skull at coordinates above the dorsolateral striatum using the stereotaxic apparatus. At this mark, a small opening (~1 mm in diameter) was drilled using small steel burr powered by high speed drill; during which saline was applied to the skull to diffuse heating and remove bone debris. Afterward, 1  $\mu$ L of AAV was injected at 200 nl/min (dorsolateral striatum coordinates relative to bregma: in mm +0.8/+1.8/–2.4). The needle was kept in place for 10 minutes and slowly removed over a 5 minute period. Subsequently, the cannulated fiber optic was slowly lowered into the striatum and secured in place using Vetbond tissue adhesive (3M Maplewood, MN) and dental cement. Three weeks after the surgery the mouse was connected to optical stimulation system (Prizmatix). The optical cannula was connected to the patch cord through the mating sleeve (Prizmatix). Blue light pulses were applied in a stimulation paradigm consisting of 10 s light on and 10 s light off for a period of 60 min. During this period the mouse behavior in an opaque open chamber (L  $\times$  W  $\times$  H in cm: 29  $\times$  29  $\times$  15) was recorded using a camera placed above the field. Locomotion was tracked by ANY-maze software (Stoelting Co.) and the tracking accuracy was verified by the investigator.

**Immunohistochemistry (IHC)**—Prior to transcardial perfusion, mice were euthanized with 200 mg/kg pentobarbital (i.p.). Once reflexes were lost, the abdominal cavity was opened and heparin (0.1 ml, 100 USP) was injected into the heart to avert blood clotting. Mice were perfused with ~50 mL of 0.1 M phosphate buffered saline (PBS) followed by ~50 mL 10% buffered formalin (Fisher Scientific). After careful removal from the skull, the brain was post fixed in 10% buffered formalin overnight. The tissue was cryoprotected (0.1 M phosphate buffered saline with 30% sucrose) the following day for at least 2 days at 4°C until use. 40  $\mu$ m sections were sectioned using a cryostat (Leica) and processed for immunohistochemistry. Sections were washed 4 times in 0.1 M PBS for 5 min each, and then incubated in a blocking solution comprising 5% NGS in 0.1 M PBS with 0.2% Triton X-100 for 1 hr at room temperature with gentle rocking. Sections were then incubated in primary antibodies diluted in 'blocking solution overnight at 4°C. The following primary antibodies were used: chicken anti-GFP (1:1000; Abcam Cat# ab13970 RRID:AB\_300798), rabbit anti-NeuN (1:1000; Cell Signaling; RRID: AB\_2630395), rabbit anti S100 $\beta$  (1:2000, Abcam Cat# ab41548 RRID:AB\_956280), guinea pig anti-cFOS (1:500, Synaptic systems #226004 RRID:AB\_2619946), rabbit anti-dsRed (1:1000, Clontech Laboratories Cat#

632496 RRID:AB\_10013483), mouse anti-mCherry (1:1000, St. John's Cat# STJ97087), guinea pig anti-NeuN (1:500; Synaptic Systems Cat # 266 004; RRID:AB\_2619988), mouse anti-neurofilament (1:250, BioLegend Cat# SMI-312R, RRID:AB\_2314906). The next day the sections were washed 4 times in 0.1 M PBS for 5 min each before incubation at room temperature for 1–2 hr with secondary antibodies diluted in 0.1 M PBS with 5% NGS and 0.2% Triton X-100. The following Alexa conjugated secondary antibodies were used: goat anti-guinea pig 488 (1:1000, Molecular Probes Cat# A-11073 RRID:AB\_142018), goat anti-rabbit 546 (1:1000, Molecular Probes Cat# A-11035 RRID:AB\_143051), goat anti-chicken 488 (1:1000, Molecular Probes Cat# A-11039 RRID:AB\_142924), goat anti-guinea pig 546 (1:1000, Thermo Fisher Scientific Cat# A-11074 RRID:AB\_2534118) and goat anti-mouse 546 (1:1000, Molecular Probes Cat# A-11030 RRID:AB\_144695). The sections were then rinsed 4 times in 0.1 M PBS for 5 min each, and briefly dipped in distilled water before being dried, mounted and coverslipped on microscope slides. Fluorescence images were taken using UPlanSApo 20X 0.85 NA and UPlanFL 40X 1.30 NA oil immersion objective lens and FV1000 Fluoview confocal laser-scanning microscope. We used the 488 nm emission wavelength of an Argon laser to excite Alexa488, with the intensity adjusted to 0.5%–10% of the maximum output, which was 10 mW. The sample emission light path consisted of a band-pass emission filter (505–525 nm) before the photomultiplier tube. Alexa 546 was excited by the 543 nm wavelength of a Helium-neon laser at 2%–30% of the maximum output (1 mW). The emitted light pathway consisted of a dichroic mirror (SDM560) and a 560–600 nm band-pass filter. For imaging Alexa 647, we used a 635 nm light emitting diode (LD635, Olympus) at 0.1%–15% of its power output which was 23 mW. The emitted light pathway consisted of a dichroic mirror (SDM640) and a 655–755 band-pass filter. For fluorescence images of brain hemisections, individual images were acquired using identical settings and stitched together to show a single composite image of the tissue section.

**Computational model of MSNs**—The computational model of the medium spiny neuron (MSN) was refined from a past study (Mahon et al., 2000), but is reported in full here. The model comprises a single compartment, representing a MSN soma and proximal dendrites, and eight active Hodgkin-Huxley (HH) based models of variable membrane conductances associated with MSN ion channel currents. The eight modeled currents are the persistent sodium current ( $I_{NaP}$ ), the slowly inactivating sodium current ( $I_{NaS}$ ), the transient sodium current ( $I_{NaT}$ ), the delayed rectifier potassium current ( $I_{KDR}$ ), the inward rectifying potassium current ( $I_{KIR}$ ), the persistent potassium current ( $I_{KRP}$ ), the fast inactivating A-current ( $I_{KAf}$ ), the slow inactivating A-current ( $I_{KAs}$ ), and a lumped, non-specific ion leak current.

The model is computationally inexpensive compared to multicompartment models (Wolf et al., 2005), and in its original form captured MSN spike timing dynamics such as characteristic latency to first spike and short-term facilitation of intrinsic excitability. A major drawback of the original model is its assumption of different reversal potentials ( $E_K$  or  $E_{Na}$ ) among ion channels selective for the same ion species, rendering it biophysically unrealistic, because these channels likely share the same intracellular and extracellular ion concentrations. To resolve this discrepancy, we replaced the model's non-specific leak

current with specific leak currents for potassium, sodium and chloride, each based on the Goldman-Hodgkin-Katz (GHK) equation:

$$\Phi_s = P_s z_s^2 \frac{V_m F^2}{RT} \frac{[S]_i - [S]_o \exp(-z_s V_m F/RT)}{1 - \exp(-z_s V_m F/RT)},$$

where  $\Phi_s$  is the current density (flux) (amperes per unit area) of ion S,  $P_s$  is the permeability of ion S ( $\mu\text{m}/\text{ms}$ ),  $[S]_i$  is the intracellular concentration of ion S (mM),  $[S]_o$  is the extracellular concentration of ion S (mM), and  $V_m$  is the membrane potential (mV). Reversal potentials for each ion species are then calculated once for all currents using the Nernst equation:

$$E_s = \frac{RT}{z_s F} \ln \left( \frac{[S]_o}{[S]_i} \right),$$

where R is the universal gas constant  $8.314 \text{ J}\cdot\text{K}^{-1}\cdot\text{mol}^{-1}$ , T is the temperature 295.15 K, z is the valence of the ionic species, and F is the Faraday's constant  $96,485 \text{ C}\cdot\text{mol}^{-1}$ . We set external and internal ion concentrations to those used for the experiments reported in preceding sections. The calculation of MSN membrane potential in our revised model then follows:

$$C \frac{dV}{dt} = I_{NaT} + I_{KDR} + I_{KIR} + I_{KRP} + I_{KAf} + I_{KAs} + I_{NaS} + I_{NaP} + I_{Cl, leak} + I_{Na, leak} + I_{K, leak} + I_{Inj}$$

where V is the membrane potential. Ion channel currents followed  $I = \bar{g} m^k h (V - E)$ , where each  $\bar{g}$  is the conductance of an ion channel current (noted by subscript).  $I_{inj}$  is an injected current.  $I_X$  is either the specific leak current for each ion species or the ion channel current. Specific leak current is derived by multiplying current density ( $\Phi_s$ ) from Equation 1 by the compartment surface area (assumed to be  $1 \mu\text{m}^2$ ). Ion channel currents are derived by multiplying channel conductance ( $\bar{g}_X$ ) by membrane potential minus the ion specific reversal potential ( $V - E_X$ ) by compartment surface area. The activation m and optional inactivation h gating variables were as reported in Mahon et al. (2000) followed:

$$\frac{dp}{dt} = \alpha_p (1 - p) - \beta_p p,$$

where p represents (m, h, n). For channel Nat and KDR, the gating parameters were as follows:

$$\alpha_m = -0.1(V + 28) / \exp((-0.1(V + 28)) - 1)$$

$$\beta_m = 4.0 \exp(-(V + 53)/18)$$

$$\alpha_h = 0.07 \exp(-(V + 51)/20)$$

$$\beta_h = 1/(\exp(-0.1(V + 21)) + 1)$$

$$\alpha_n = -0.01(V + 27)/\exp((-0.1(V + 27)) - 1)$$

$$\beta_n = 0.125 \exp(-(V + 37)/80)$$

All other currents obeyed the Boltzmann equation for steady state activation

$$s_\infty(V) = 1/(1 + \exp(-(V - V_s)/k_s))$$

and the voltage dependent time constants followed

$$\tau(V) = \frac{\tau_0}{\left( \exp^{-\frac{V - V_\tau}{k_\tau}} + \exp^{\frac{V - V_\tau}{k_\tau}} \right)}$$

except for the time constant of  $I_{As}$  and  $I_{KRP}$ . The inactivation for these currents, followed

$$\tau_{hAs}(V) = 1790 + 2930 \exp\left(\frac{-(V + 38.2)}{28^2}\right) \left(\frac{V + 38.2}{28}\right)$$

and

$$t_{KRP}(V) = 3 * t_{hAs}(V)$$

For numerical integration, we chose a time step of 0.01 ms. The datasets used for the model were archival at the time they were shared with researchers from IBM, and no new experiments were suggested, designed, or performed based on these models and analyses. All simulations were performed using the IBM Neural Tissue Simulator (NTS) (Kozloski and Wagner, 2011) on IBM Cloud. NTS executes simulations based on model descriptions (written in the Model Description Language) and resource allocation scripts (written in the Graph Specification Language), both of which are available in Supplemental Information.

The software is experimental, and readers are therefore encouraged to contact the IBM authors if interested in using the tool.

We employed an optimization technique to search parameter space of this model. Each optimization targeted features extracted from measures taken during experiments under differing conditions as reported in Figure 3. Specifically, we optimized the single neuron model to reproduce both the Light off and Light on experimental conditions. By first setting the extracellular  $K^+$  concentrations to 4.5 mM and 7.5 mM, respectively for each condition, we guided automatic changes to model conductance parameters toward models that replicated both experimental measurements and conditions simultaneously. The optimizations accessed and varied 11 parameters, the maximal conductances  $\bar{g}_{NaT}$ ,  $\bar{g}_{NaP}$ ,  $\bar{g}_{NaS}$ ,  $\bar{g}_{KDR}$ ,  $\bar{g}_{KRP}$ ,  $\bar{g}_{KIR}$ ,  $\bar{g}_{KAf}$ ,  $\bar{g}_{KAs}$  and GHK based permeability coefficients  $P_{Cl, leak}$ ,  $P_{Na, leak}$  and  $P_{K, leak}$ .

The optimization employed the non-dominated sorting (NS) differential evolution (DE) algorithm (NSDE) (Deb et al., 2002; Price et al., 2005), previously used to search parameter space of compartmental neuron models in Rumbell et al. (2016). To run the algorithm, we used a modified version of the BluePyOpt (Van Geit et al., 2016) python framework for single neuron optimization. Each parameter set was simulated for 5 s. Trial-and-error was used to assess optimization metaparameters such as population size and number of generations. A single optimization of ~500 generations of a population of ~100 models took approximately ~12 hours of computing time on an X86\_64 Intel architecture (2 Ghz, 64 bit, 56 cores, 128 Gb of RAM). The neuron model error score for each target feature was calculated by extracting feature measures and subtracting them from the exact target values based on empirical measures and dividing the absolute value of this quantity by a deviation variable based on variability of the experimental measures. Dominance ranking according to the NS algorithm was used as the first criterion for model selection, and total error was used to sort models within dominance ranks. We ran three optimizations, first to validate the modified version of the model with specific ion leaks to reproduce MSN model spiking features as in the original implementation (Mahon et al., 2000). All of the relevant features targeted and parameters thus obtained are reported in Tables S1-S5. A second optimization to find a population of parameter sets that reproduced the range of empirically observed excitability measures with Light off and on conditions by the two extracellular  $K^+$  conditions, and third to target several spiking features from one neuron recording to produce a more detailed characterization of MSN firing, sufficiently accurate to allow interpolation of excitability measures when tested with various extracellular  $K^+$  values. We targeted 4 features for the second optimization: empirical measurements of resting membrane potential (RMP) and rheobase (Rh) in both light on and light off conditions, modeled here as 4.5 mM  $K^+$  and 7.5 mM  $K^+$  respectively. Target feature values were chosen as (target value  $\pm$  deviation)  $70.0 \pm 4.0$  (RMP, 4.5 mM  $K^+$ ),  $60.4 \pm 4.0$  (RMP, 7.5 mM  $K^+$ ),  $310.23 \pm 10.0$  pA (Rh, 4.5 mM  $K^+$ ), and  $200.89 \pm 8.10$  pA, to approximate the variability around the empirical median of these features. These models were also constrained by change in initial RMP voltage variability  $\sim 0$ . For this optimization, the NSDE algorithm (Deb et al., 2002; Price et al., 2005) used in Rumbell et al. (2016) was modified by subtracting 2.0 from the calculated error score for each feature, rounding up to 0.0. This ensured that any feature values

calculated from models that lie within two standard deviations of the mean would be considered equally acceptable as a potential model of an empirical observation. As this meant that any models with all 4 feature values within 2 standard deviations of the target mean would have a total error score of 0.0, we used a crowdedness function to sort models with equal total error score, calculated as Euclidean distance to the nearest neighbor in feature space. This additional selection criterion encouraged the algorithm to find models filling the range of acceptable features within 2 standard deviations of the target mean. We ran this optimization for 1000 generations with population size 110, simulating 110,000 unique parameter sets, among which 1119 acceptable models with total error score of 0.0 were identified. From among these, 20 were selected randomly and their resulting feature values displayed in Figure 4B. For the third optimization, we chose 14 features to optimize in the models, which characterized measurements of neuron properties (Figure 4) such as Rheobase, resting membrane potential, inter-spike interval distributions, action potential shape. Here the target feature values were precise and extracted from one empirical recording, and the ‘standard deviations’ used here was selected as 5% of the target feature value for each feature, except RMP where smaller ‘standard deviation’ values ~0.1 were used, to prevent the optimization from entering local minima (Tables S1-S5). As this optimization targeted one particular voltage trace, the above method for reducing errors to 0.0 within 2 standard deviations was not used. We ran this optimization for 5000 generations with a population size of 220, testing 1,100,000 unique parameter sets, at which point we terminated the optimization and selected the best fitting member of the population to demonstrate the voltage traces shown in Figure 4C. This model was then simulated in conditions with extracellular  $K^+$  between 4.5 and 7.5 mM in increments of 0.5mM, demonstrating reliability of the change in excitability produced by this modulation (Figure 4D).

***In vivo* multi-unit recordings of neuronal activity**—Three weeks following AAV microinjection in the dorsolateral striatum, adult 9 week old male C57Bl6N mice were anesthetized with isoflurane (3%–5% induction, 1.5% maintenance). Ten minutes after injection of a systemic analgesic (carprofen, 5 mg per kg of body weight) mice were placed in a stereotaxic frame. Mice were maintained at 37°C at all times using a feedback-controlled heating pad. Pressure points and incision sites were injected with lidocaine (2%), and eyes were protected from desiccation using artificial tear ointment. The skin above the skull was incised, a custom-made lightweight metal head holder was implanted on the skull using Vetbond (3M) and a recording chamber was built using dental cement (Ortho-Jet, Lang). Mice had a recovery period from surgery of five days, during which they were administered amoxicillin (0.25 mg per ml in drinking water through the water supply). After the recovery period, mice were habituated to head fixation on the spherical treadmill. On the day of the recording, mice were anesthetized with isoflurane. To fix the ground wire, a small craniotomy (0.5 mm diameter) was made above the right cerebellum and a silver wire was implanted at the surface of the craniotomy and fixed with dental cement. A circular craniotomy 1 mm in diameter was made at a location above the dorsolateral striatum and the mice was allowed to recover from anesthesia. The recording chamber was filled with cortical buffer containing (in mM) 135 NaCl, 5 KCl, 1.8 CaCl<sub>2</sub> and 1 MgCl<sub>2</sub>. The head-bar was

fixed to a post and the mouse was placed on the spherical treadmill consisting of a ball floating on small cushion of air allowing for 2D movement.

All recordings were performed at least two hours after the end of anesthesia, when the mouse was alert and could actively behave. A 128-channel silicon nanoprobe was lowered into the brain (Shobe et al., 2015). Each nanoprobe had 2 shanks with 64 electrodes on each shank. Each shank was 1.05 mm long and 86  $\mu\text{m}$  at its widest point and tapered to a tip. Each prong was separated by 400  $\mu\text{m}$ . A multiplexed analog output signal was sent from the head stage (Intan Technologies, RHA-2164B) of each probe via thin flexible cables to a 16-bit analog to digital conversion cards (USB-6356, National Instruments), which sampled each signal at a rate of 25 kHz per channel. Signals were then filtered offline and a background signal subtraction was performed. Unit clustering was performed using JRClust (Jun et al., 2017), a fully-automated method for clustering units based on unit spike properties across multiple channels. We only included high-quality units (Butterworth filter 300–6000 Hz, spikes count > 1000, SNR > 7.0). We performed firing rate analysis in MATLAB; we averaged each unit firing rates across 5 trials with 5 s activity bins. For each experimental condition a raster graph was plotted by combining all spike times from 5 trials for each unit and down sampling in 5:1 ratio.

**General reagents**—All general chemicals used were from Cayman Chemical, Tocris, Thermo-Fisher or Sigma. Other specific reagents and kits are listed in the methods.

## QUANTIFICATION AND STATISTICAL ANALYSIS

Data from every experiment represents a minimum of  $n$  animals with a balanced number of male and female mice. In all the experiments,  $n$  was 4 mice; sample sizes were not calculated *a priori*. For AAV injections, mice were randomly assigned to each experimental group. No experimental data points were excluded. Statistical tests were run in OriginPro 2015. Summary data are presented as mean  $\pm$  SEM. Note that in some of the graphs, the bars representing the s.e.m. are smaller than the symbols used to represent the mean. For every dataset, we determined within Origin whether the data were normally distributed or not using the Shapiro-Wilk test. For tests of statistical significance, we used parametric tests for normally distributed data and non-parametric tests for data that was not normally distributed. Paired and unpaired Student's two-tailed t tests (as appropriate) and two tailed Mann-Whitney or Wilcoxon paired sign tests were used for most statistical analyses with significance declared at  $P < 0.05$ . When a statistical test was employed that was not a Student's t test or a Mann-Whitney test for a specific case, then it is stated as such in the text. Specific P values are stated in the figures, however, when the P value was less than 0.001, it is stated as  $p < 10^{-3}$  to save space on the figure panels and text. However, where appropriate key statistics are also reported in the text. If the P value was greater than 0.05, then it is stated as  $p > 0.05$ . The sample size,  $n$ , is defined as the number of cells, numbers of slices, fields-of-view or numbers of mice depending on the experiment on a case-by-case basis.

## DATA AND SOFTWARE AVAILABILITY

The AAV expression vector for ChR2(H134R)-mCherry using the *GfaABC1D* promoter will be available from Addgene (#112496). Data analyses and software guidance is available upon request. The MSN model is available from James R. Kozlowski at IBM.

## Supplementary Material

Refer to Web version on PubMed Central for supplementary material.

## ACKNOWLEDGMENTS

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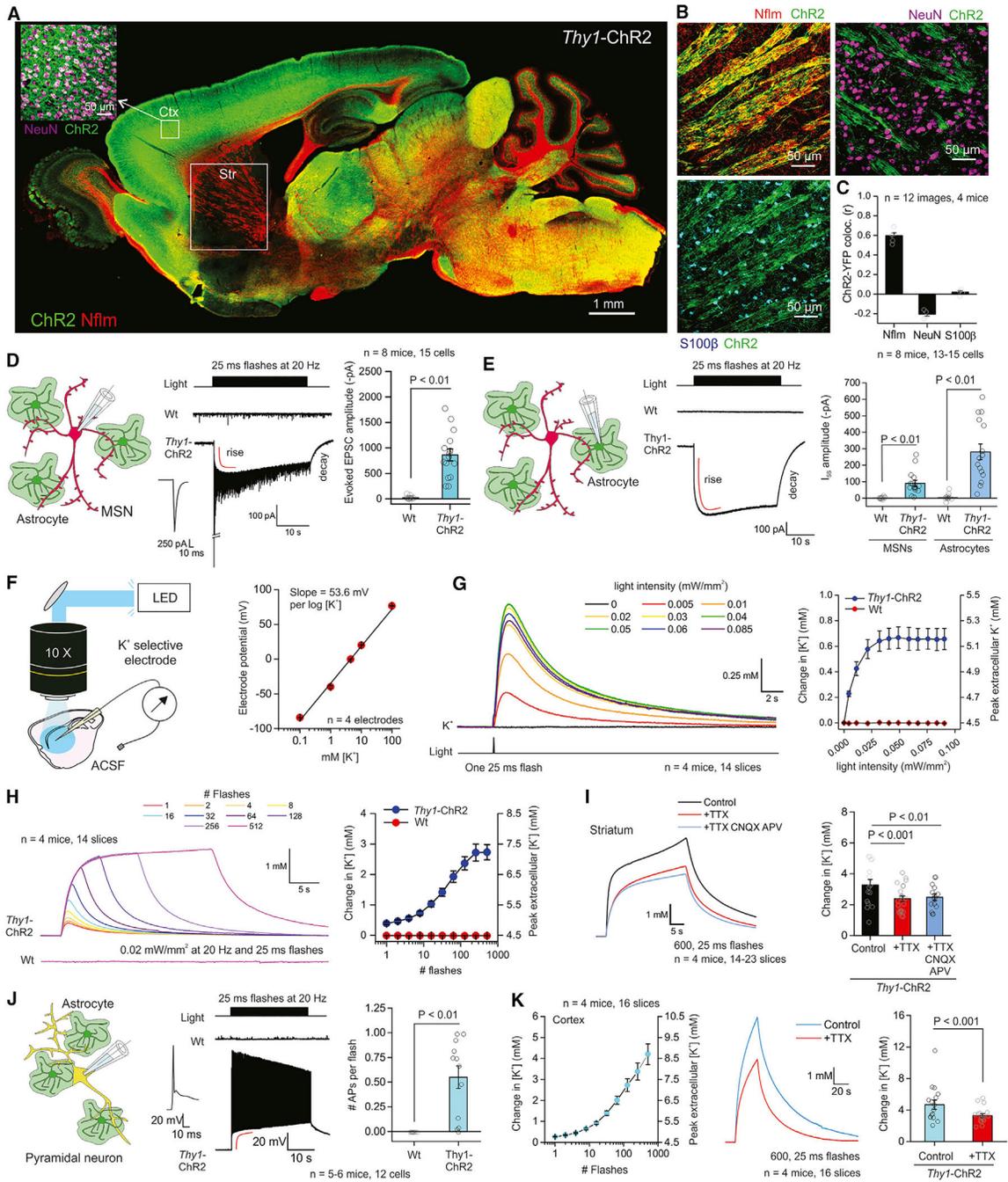
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### Highlights

- Multiple approaches reveal transient  $K^+$  elevations during ChR2 excitation
- ChR2-mediated  $K^+$  elevations increase neuronal excitability and cFos expression
- Neuronal effects of  $K^+$  are recapitulated with a model and *in vivo*
- Increased  $K^+$  may contribute to astrocyte experiments employing ChR2 *in vivo*



**Figure 1. Optical Stimulation of ChR2-Expressing Cells Evokes Transient Elevation of Extracellular K<sup>+</sup>**

(A) Representative image showing brain distribution of channelrhodopsin2 (ChR2-YFP; green) in a *Thy1-ChR2-YFP* mouse brain section together with neurofilament (Nflm, red). The inset shows strong colocalization of ChR2 and NeuN positive neurons in cortex. (B) The images show colocalization of ChR2 with Nflm positive axonal tracts in the striatum, but not with NeuN or S100β positive neurons and astrocytes. (C) Pearson's colocalization correlation coefficient (*r*) for images such as those shown in (B). (D) Evoked EPSC amplitude in MSNs for Wt and *Thy1-ChR2*. (E) *I<sub>h</sub>* amplitude in astrocytes for Wt and *Thy1-ChR2*. (F) K<sup>+</sup> selective electrode setup and its calibration. (G) K<sup>+</sup> transients in response to light flashes of varying intensities. (H) Extracellular K<sup>+</sup> transients in response to light flashes in *Thy1-ChR2* and Wt slices. (I) TTX and APV block the K<sup>+</sup> transients in the striatum. (J) Evoked APs in pyramidal neurons for Wt and *Thy1-ChR2*. (K) Extracellular K<sup>+</sup> transients in the cortex for Wt and *Thy1-ChR2*.

(D) Left: schematic depicting patch-clamp of medium spiny neurons (MSNs) in the striatum. Middle: representative current waveforms from MSNs in *Thy1-ChR2-YFP* or WT mice in response to a train of flashes to stimulate corticostriatal axons. Right: peak amplitude of the first EPSC upon blue light flashes.

(E) Left: schematic depicting patch-clamp of astrocytes in the striatum. Middle: representative current waveforms from striatal astrocytes in *Thy1-ChR2-YFP* or WT mice in response to a train of flashes. Right: quantification of the steady-state inward current ( $I_{ss}$ ) from MSNs and astrocytes in the striatum from WT or *Thy1-ChR2-YFP* mice.

(F) Left: schematic depicting the experimental setup for recording of  $K^+$  dynamics in brain slices with blue light activation of ChR2. Right: graph plots the response of the  $K^+$  electrode to various  $K^+$  concentrations in the extracellular fluid.

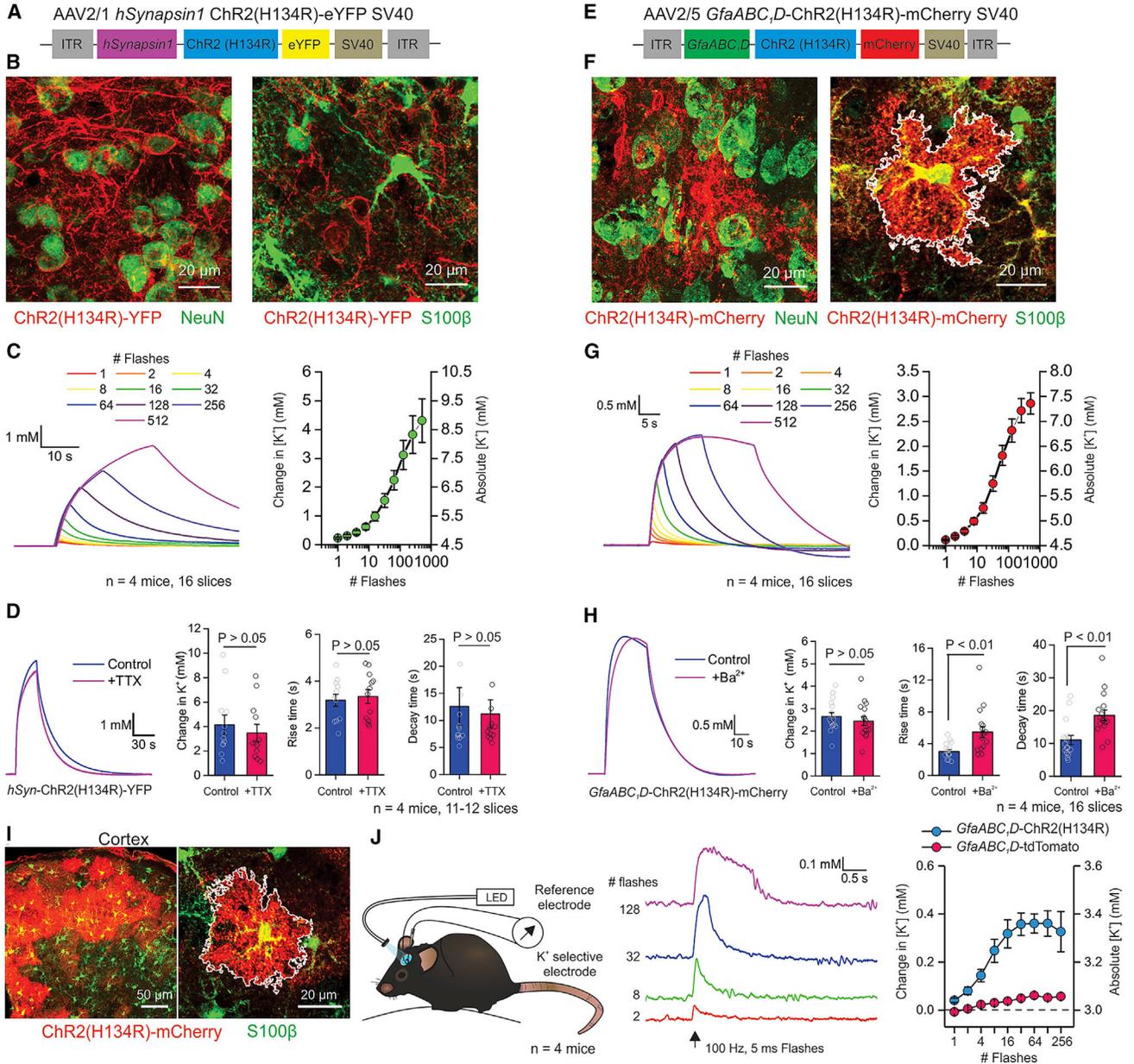
(G) Left: waveforms of changes in  $K^+$  concentration in response to a single flash of various light intensities. Right: quantification of the peak amplitude of  $K^+$  in WT and *Thy1-ChR2-YFP* mice in response to single flashes of various light intensities.

(H) Left: waveform depicting changes in  $K^+$  in response to different numbers of flashes in *Thy1-ChR2* and WT mice. Right: quantification of the peak amplitude of the  $K^+$  elevation in WT and *Thy1-ChR2-YFP* mice in response to different numbers of flashes.

(I) Left: waveform depicting  $K^+$  responses in the striatum with TTX or TTX, CNQX, and APV. Right: average data for experiments in the left panel.

(J) Left: schematic depicting patch-clamp of pyramidal neurons in the cortex. Middle: representative voltage waveform in cortical pyramidal cells from WT or *Thy1-ChR2* mice with light flashes. Right: quantification of the number of action potentials.

(K) Left: quantification of the peak  $K^+$  concentration in the cortex. Middle: representative waveform of  $K^+$  concentrations in the cortex. Right: average data from the experiment depicted in the left panel. Data shown as mean  $\pm$  SEM. In some cases, the error bars are smaller than the symbol used to represent the mean. See also Figures S1—S3.



**Figure 2. Transient Extracellular K<sup>+</sup> Increases Occur Following AAV-Driven Expression of ChR2**

(A) AAV vector for neuron-specific expression of ChR2(H134R).  
 (B) Left: representative image of the dorsolateral striatum showing the colocalization of *hSyn*-ChR2(H134R)-YFP (red) and NeuN (green). Right: representative image of *hSyn*-ChR2(H134R)-YFP (red) and S100β (green).  
 (C) Left: waveforms depicting striatal K<sup>+</sup> elevations in mice expressing the *hSyn*-ChR2. Right: average data for *hSyn*-ChR2(H134R)-YFP K<sup>+</sup> elevations.  
 (D) Left: waveforms of K<sup>+</sup> elevations with and without TTX. Right: quantification of various properties of the responses shown in the left panel.  
 (E) AAV vector for astrocyte-specific expression of ChR2(H134R).

(F) Left: representative image showing the distribution of *GfaABC<sub>1</sub>D-ChR2* showing overlap with S100 $\beta$  (green). Right: representative image of the dorsolateral striatum showing colocalization of *GfaABC<sub>1</sub>D-ChR2* (red) and NeuN (green).

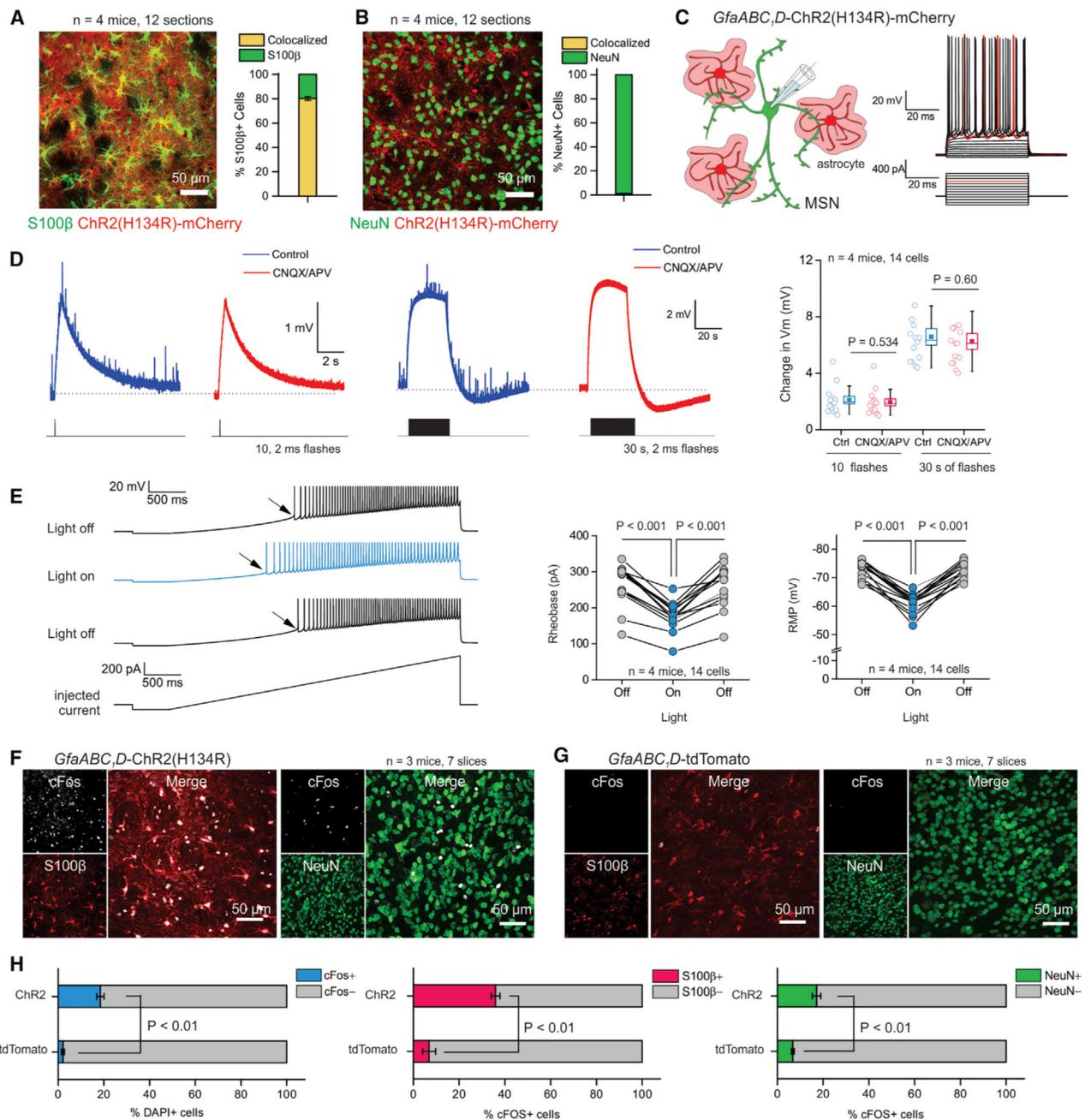
(G) Left: waveforms of K<sup>+</sup> elevations in mice expressing the *GfaABC<sub>1</sub>D-ChR2*. Right: average data for the experiment in the left panel.

(H) Left: waveform of K<sup>+</sup> elevations with and without Ba<sup>2+</sup> applied to the slice. Right: quantification of various properties of the responses in the left panel.

(I) Representative images from animals expressing ChR2(H134R) in cortical astrocytes, showing overlap with S100 $\beta$  in green.

(J) Left: schematic depicting the experimental setup for *in vivo* monitoring of K<sup>+</sup> concentrations. Middle: waveforms of K<sup>+</sup> concentration changes in the cortex from *in vivo* recordings. Right: average peak K<sup>+</sup> concentration in the *in vivo* recordings. The LED power was 0.33 mW/mm<sup>2</sup>. Data shown as mean  $\pm$  SEM. In some cases, the error bars are smaller than the symbol used to represent the mean.

See also Figures S2 and S4.



**Figure 3. Chr2-Mediated Extracellular K<sup>+</sup> Increased Neuronal Excitability**

(A) Left: representative image of a striatal section from mice expressing *GfaABC<sub>1</sub>D*-Chr2(H134R) (red), showing overlap with S100β (green). Right: fraction of S100β positive cells expressing *GfaABC<sub>1</sub>D*-Chr2(H134R) in the dorsolateral striatum.

(B) Left: representative image from mice expressing *GfaABC<sub>1</sub>D*-Chr2(H134R) (red), showing no overlap with NeuN (green). Right: fraction of NeuN positive cells expressing *GfaABC<sub>1</sub>D*-Chr2(H134R) in the dorsolateral striatum.

(C) Left: schematic depicting patch-clamp of medium spiny neurons (MSNs) in the striatum near astrocytes expressing ChR2(H134R). Right: waveform of membrane potential changes of a MSN to a series of stepwise current injections.

(D) Left: traces of membrane potential changes from MSNs in response to a train of 10 flashes with and without glutamate antagonists in mice expressing *GfaABC1D*-ChR2. Traces for 30 s of flashes with and without glutamate antagonists are also shown. Right: average data from experiments shown with the traces.

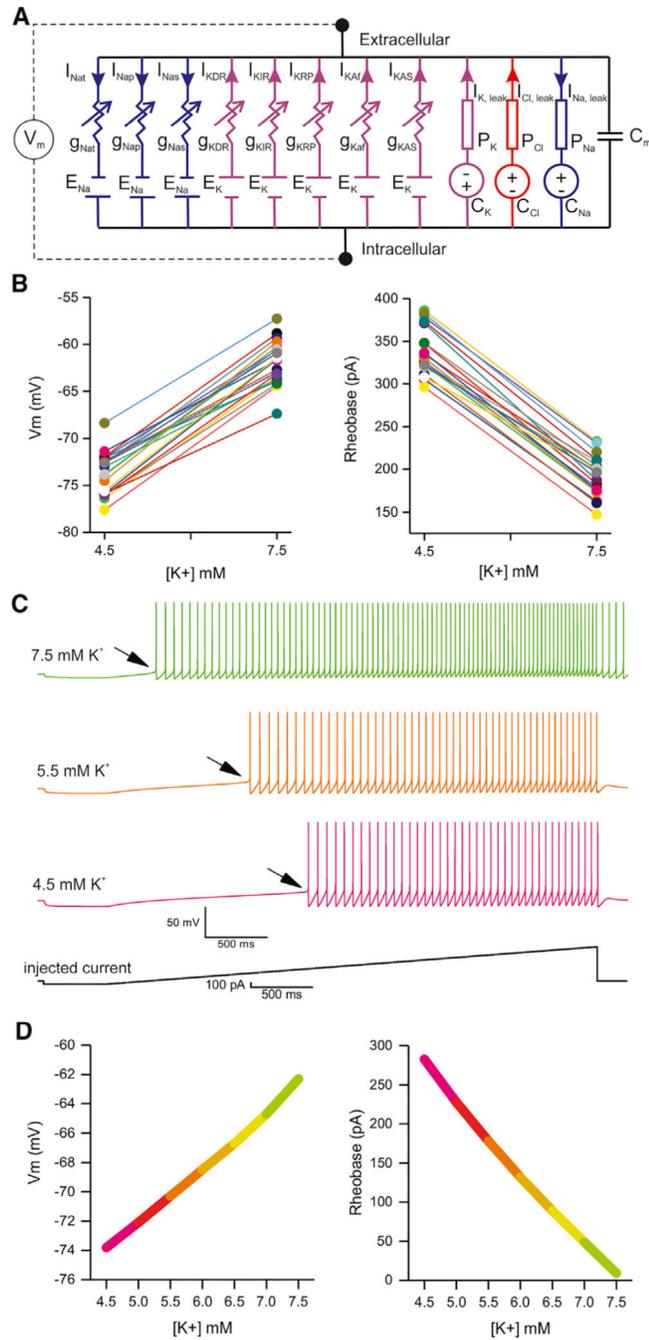
(E) Left: waveforms of membrane potential responses of a MSN to a ramp current injection with and without blue light. Inset: highlighted region showing rheobase shift. The scatter graphs show average data for rheobase and membrane potential.

(F) Left: representative images showing overlap between cFos (upper left inset, white) and S100 $\beta$  (lower left inset, red) from animals in which ChR2 was expressed in astrocytes and activated with blue light. Right: as in left panels, but for NeuN (green) instead of S100 $\beta$ .

(G) As in (F), but for control experiments expressing tdTomato instead of ChR2.

(H) Summary plots of cFos expression following optical stimulation of astrocytes expressing ChR2 in relation to control experiments where astrocytes expressed tdTomato. For the experiments in (F)-(H), the stimulation conditions were 0.05 Hz (10 s on and 10 s off for 1 hr). The power was ~3.5 mW from the tip of the cannula before implantation. Data shown as mean  $\pm$  SEM. In some cases, the error bars are smaller than the symbol used to represent the mean.

See also Figures S3 and S4.



**Figure 4. Computational Modeling of  $K^+$  Effects In Silico**

(A) Diagrammatic representation of the circuit model of a MSN with the indicated active conductances and ion-specific leak currents. Key values used in the model are listed in Table 1: kinetic parameters were unchanged from those reported in published work (Mahon et al., 2000).

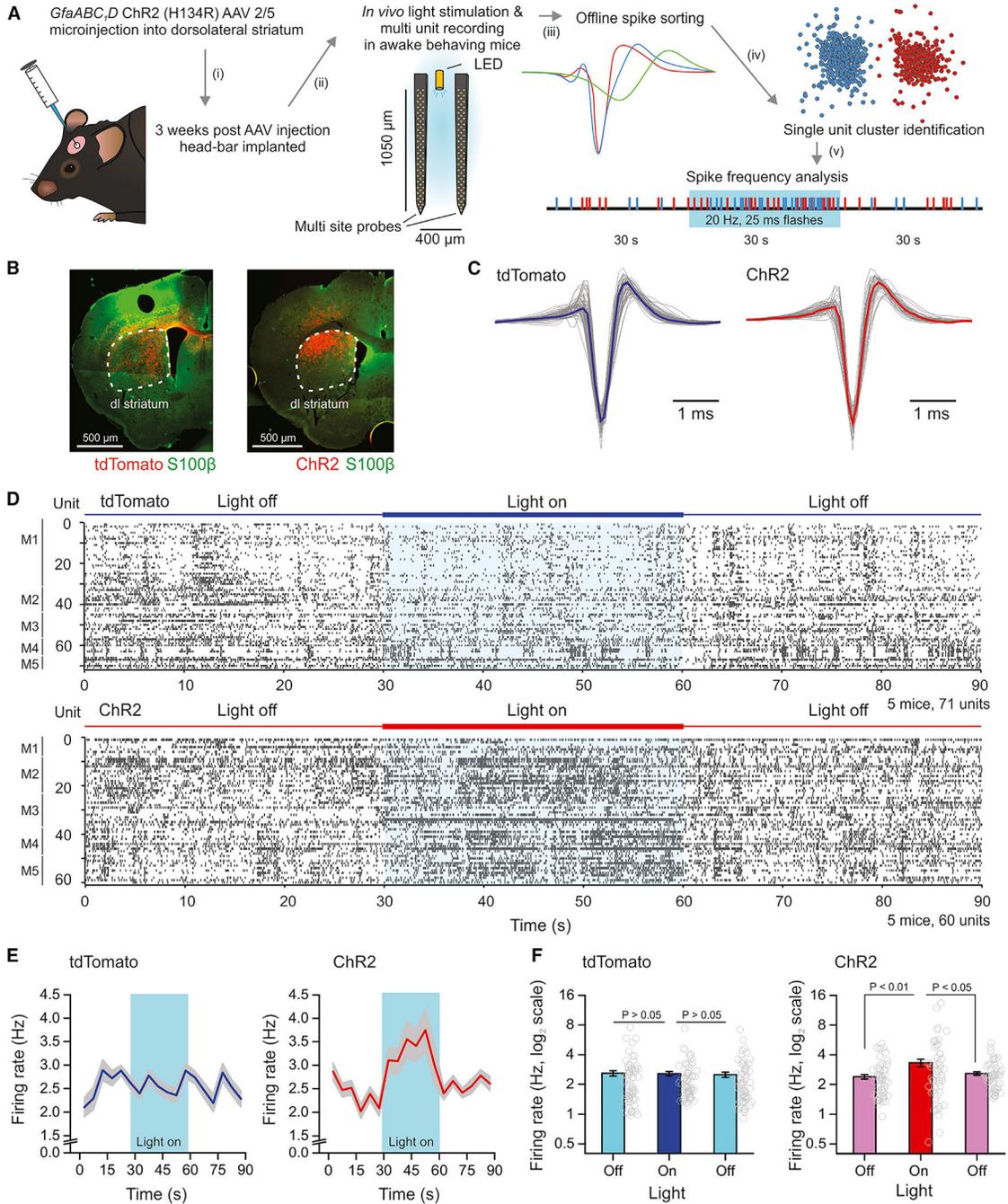
(B) Scatter graphs plot the effect of  $K^+$  elevations from 4.5 to 7.5 mM on the MSN model population's resting membrane potentials and rheobases. Each conductance parameter set

(model instance) is represented as a pair of points joined by a line. Across scatterplots, colors represent the same model instance.

(C) Traces of MSN excitability from the computational model at the indicated levels of  $K^+$ . The arrows emphasize the shift in rheobase with increasing  $K^+$ .

(D) Summary plots of how resting membrane potential and rheobase change as a function of extracellular  $K^+$  in the MSN model depicted in (C). Full details of the MSN model are provided in the STAR Methods section.

See also Tables S1 — S5.



**Figure 5. *In Vivo* Activation of ChR2 in Astrocytes Enhances MSN AP Firing Measured with Silicon Probes in Awake Behaving Mice**

(A) Schematic depicting experimental workflow using multi-site silicon probes *in vivo*. Mice were injected with AAVs to express tdTomato or ChR2 within astrocytes. Light flashes were produced by 30 s of 25 ms flashes at 20 Hz and at 470 nm using a LED light source. (B) IHC staining of coronal sections from mice injected with AAVs expressing tdTomato (left) or ChR2 (right) in astrocytes of the dorsolateral striatum (white dashed outline) and used for *in vivo* recordings.

- (C) Representative waveforms from narrow spiking single units in either tdTomato (left) or ChR2 (right) expressing mice. The average trace is the thick colored line.
- (D) Raster plots of single unit recordings from 5 mice expressing tdTomato (top graph) or ChR2 (bottom graph) during 30 s periods with the LED off or on.
- (E) Average firing rate of all units over the recording period (in 5 s bins) in tdTomato or ChR2 groups of mice. The period optical stimulation is highlighted in blue.
- (F) Bar graphs of average activity of all units in the 30 s periods prior to, during, or following optical stimulation in either condition. Average data are represented as mean  $\pm$  SEM from n = 5 mice.

Comparison of Chr2-Mediated Extracellular K<sup>+</sup> Elevations at Room Temperature and at More Physiological Temperatures

Table 1.

	# Slices [# Mice]		Peak Change in K <sup>+</sup> (mM)		Rise Time (τ; s)		Decay Time (τ; s)		p		
	20°C	34°C	20°C	34°C	20°C	34°C	20°C	34°C			
Transgenic mouse	15 [4]	18 [4]	2.7 ± 0.2	2.7 ± 0.3	0.55	2.1 ± 0.6	5.3 ± 0.7	< 0.001	7.2 ± 0.5	14.4 ± 3.8	0.07
<i>Thy1</i> Chr2											
AAV <i>hSynapsin1</i> Chr2	12 [4]	16 [4]	4.3 ± 0.8	3.5 ± 0.3	0.29	3.2 ± 0.3	4.7 ± 0.5	0.02	9.5 ± 1.4	18.9 ± 1.5	< 0.001
AAV <i>GliaABC/D</i> Chr2	16 [4]	20 [4]	2.9 ± 0.2	2.7 ± 0.1	0.52	3.0 ± 0.2	2.9 ± 0.3	0.74	11.0 ± 1.4	12.8 ± 1.0	0.12

The peak change in K<sup>+</sup> concentrations were not significantly different at room (20°C) versus a more physiological temperature (34°C) in acute slice experiments in the striatum. However, some aspects of the kinetics (τ) were. All values are reported as mean ± SEM from the number of slices indicated (n) from 4 mice in each condition.

**Table 2.**

Model Parameters Used in MSN Computations Shown in Figure 4

Current	Gate	$\bar{g}$ (nS· $\mu\text{m}^{-2}$ )	$S_{\infty}$			$\tau$ (V)		
			$V_s$ (mV)	$k_s$ (mV)	$\tau_0$ (ms)	$V_c$ (mV)	$k_c$ (mV)	
$I_{NaT}$	$m_{NaT}^3 h_{NaT}$	1.61	See STAR Methods					
$I_{Nap}$	$m_{Nap}$	$3.1 \times 10^{-3}$	-47.8	3.1	1.0	-	-	
$I_{Nas}$	$m_{Nas}$	0.02	-16.0	9.4	637.8	-33.5	26.3	
$I_{KDR}$	$n_{KDR}^4$	0.11	See STAR Methods					
$I_{KIR}$	$m_{KIR}$	$0.33 \times 10^{-3}$	-100	-10	-	-	-	
$I_{KRP}$	$m_{KRP}$	0.155	-13.4	12.1	206.2	-53.9	26.5	
	$h_{GRP}$		See STAR Methods					
$I_{KAF}$	$m_{KAF}$	0.0017	-33.1	7.5	1.0	-	-	
	$h_{CAF}$		-70.4	-7.6	25.0	-	-	
$I_{KAS}$	$m_{KAS}$	$6.7 \times 10^{-6}$	-25.6	13.3	131.4	-37.4	27.3	
	$h_{CAS}$		See STAR Methods					

Full details are in the STAR Methods. Here,  $P_{Na}$ , leak =  $1.96 \times 10^{-7}$ ,  $P_{Cl}$ , leak =  $1.5 \times 10^{-7}$ , and  $P_K$ , leak =  $0.6 \times 10^{-7}$  ( $\mu\text{m}/\text{ms}$ ). The equilibrium potentials for  $\text{Na}^+$ ,  $\text{K}^+$  and  $\text{Cl}^-$  were 56.64 mV, -87.06 mV, and -95.07 mV at 295.15 K, respectively, and were calculated using the Nernst equation for each ion with concentrations identical to those used in the experiments.

## KEY RESOURCES TABLE

REAGENT or RESOURCE	SOURCE	IDENTIFIER
Antibodies		
Chicken anti-GFP	Abcam	Cat# ab13970; RRID: AB_300798
Rabbit anti-NeuN	Cell Signaling	Cat# 12943; RRID: AB_2630395
Mouse anti-mCherry	Saint John's	Cat#STJ97087; RRID:AB_2687534
Rabbit anti-DsRed	Clontech Laboratories	Cat# 632496; RRID:AB_10013483
Rabbit anti-S100b	Abcam	Cat# ab41548; RRID: AB_956280
Guinea pig anti-Neun	Synaptic Systems	Cat# 266 004; RRID:AB_2619988
Guinea pig anti-cFOS	Synaptic Systems	Cat# 226 004; RRID:AB_2619946
Rabbit anti-cFOS	Abcam	Cat# ab190289; RRID:AB_2737414
Mouse anti-neurofilament	Biologend	Cat# SMI-312R; RRID:AB_2314906
Alexa goat anti-rabbit 488	Molecular Probes	Cat# A11008; RRID: AB_143165
Alexa goat anti-chicken 488	Molecular Probes	Cat# A11039; RRID: AB_142924
Alexa goat anti-rabbit 546	Molecular Probes	Cat#A11010; RRID: AB_143156
Alexa goat anti-mouse 546	Molecular Probes	Cat#A11003; RRID: AB_141370
Alexa goat anti-guinea pig 488	Molecular Probes	Cat# A-11073; RRID:AB_142018
Alexa goat anti-guinea pig 546	Molecular Probes	Cat# A-11074 RRID:AB_2534118
Bacterial and Virus Strains		
AAV2/5 <i>GfaABC<sub>1</sub>D</i> ChR2(H134R)-mCherry SV40	This manuscript, UPenn Vector Core	Available upon request from UPenn
AAV2/1 <i>hSynapsin1</i> ChR2(H134R)-eYFP SV40	UNC GTC vector core	Lot# AV4319J
Chemicals, Peptides, and Recombinant Proteins		
TTX	Cayman Chemical Company	Cat# 14964
CNQX	Tocris	Cat# 0190
D-AP5	Tocris	Cat# 0106
Barium chloride	Sigma	Cat# 202738
Normal goat serum	Vector	Cat# S-1000
Critical Commercial Assays		
QIAquick PCR Purification Kit	QIAGEN	Cat# 28104
QIAGEN Mini kit	QIAGEN	Cat# 27104
QIAGEN Endofree MAXI kit	QIAGEN	Cat# 12362
QIAquick Gel extraction Kit	QIAGEN	Cat# 28704
Experimental Models: Organisms/Strains		
Mouse: C57BL/6N inbred mice	Taconic	JAX Stock # 005304
Mouse: <i>Thy1</i> -ChR2-YFP	Arenkiel et al., 2007	JAX Stock # 007615
Recombinant DNA		
pZac2.1 <i>GfaABC<sub>1</sub>D</i> ChR2(H134R) SV40	This manuscript; Addgene	Plasmid # 112496
Software and Algorithms		
OriginPro 8.5/9/2015	Origin Lab	N/A

<b>REAGENT or RESOURCE</b>	<b>SOURCE</b>	<b>IDENTIFIER</b>
GraphPad InStat 3	GraphPad Software	N/A
pCLAMP10	Molecular Devices	N/A
ClampFit 10.5	Molecular Devices	N/A
Fluoview FV3000	Olympus	N/A
ImageJ v1.30	ImageJ	N/A
ANY-maze behavioral tracking software	Stoelting Co.	N/A
Prizmatix Pulser	Prizmatix Ltd.	N/A
MSN model parameter file	GSL file submitted here	N/A

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