

Supplemental information

**Severe deficiency of the voltage-gated
sodium channel Na_v1.2 elevates
neuronal excitability in adult mice**

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Supplementary Figures:

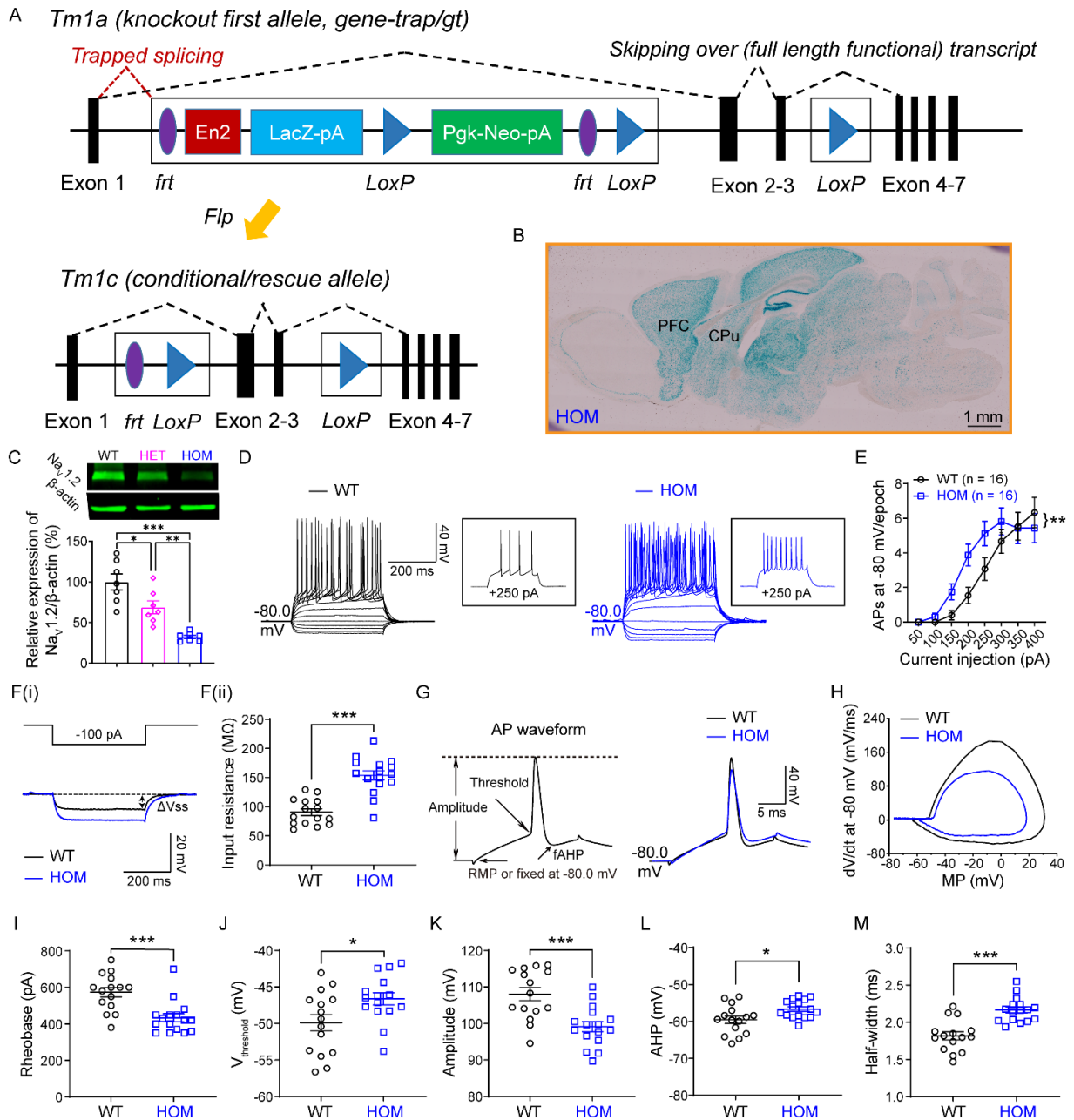


Figure S1. Elevated neuronal firings of striatal MSNs at a fixed membrane potential of -80 mV in adult *Nav1.2*-deficient mice. Related to Figure 1.

(A) Gene trap (gt) allele has an inserted *tm1a* trapping cassette between the Exon 1 and Exon 2 of *Scn2a* gene in the genome, which traps the transcription from Exon 1 to *tm1a* cassette, resulting in a deficiency of *Scn2a*. In the presence of Flp recombinase, *frt* sites flanked trapping cassette will be removed, producing conditional ("rescue") allele that allows the expression of

Scn2a like the WT. *flp*, *Flp* recognition target (purple); *En2*, engrailed-2 splice acceptor (red); *LacZ*, *lacZ* β -galactosidase (light blue); *LoxP*, locus of X-over P1 (dark blue); and *Neo*, neomycin (green).

(B) *gt* cassette contains a *LacZ* element and is driven by the native *Scn2a* promoter. Thus, the *LacZ* expression can be used as a surrogate of *Scn2a* expression. Representative *LacZ* staining of a sagittal slice from a *Scn2a^{gt/gt}* (HOM) mouse showing a strong blue signal across the brain including the prefrontal cortex (PFC) and dorsal striatum (CPu, caudate nucleus and the putamen). Scale bar, 1 mm.

(C) Upper: Representative Western blots of striatal tissues from WT (black circle), HET (magenta diamond), and HOM (blue square) mice. Lower: associated quantification of $\text{Na}_v1.2$ protein. One-way ANOVA followed by Tukey's multiple-comparison test: * $p < 0.05$; ** $p < 0.01$; *** $p < 0.001$.

(D) Representative current-clamp recordings of MSNs from WT (black) and HOM (blue) mice were obtained at a fixed membrane potential of -80 mV. A series of 400-ms hyperpolarizing and depolarizing steps in 50-pA increments were applied to produce the traces. Inset: representative trace in response to +250 pA current injection.

(E) The average number of APs generated in response to depolarizing current pulses at -80 mV. Unpaired two-tailed non-parametric Mann-Whitney *U*-test for each current pulse: ** $p < 0.01$.

(Fi) Representative traces in response to -100 pA current injection. $V_{\text{steady-state}}$ (V_{ss}) is the voltage recorded at 0-10 ms before the end of the stimulus.

(Fii) Individuals and average input resistance values at -80 mV. Unpaired two-tailed Student's *t*-test: *** $p < 0.001$.

(G) Left: plot of a typical AP showed its various phases. Right: typical spikes of MSNs from WT (black) and HOM (blue) mice were obtained at a fixed membrane potential of -80 mV.

(H) Associated phase-plane plots.

(I-M) Individuals and average spike rheobase, voltage threshold, amplitude, fast after-hyperpolarization (AHP), and half-width values. unpaired two-tailed Student's *t*-test: * $p < 0.05$; *** $p < 0.001$. Data were shown as mean \pm SEM.

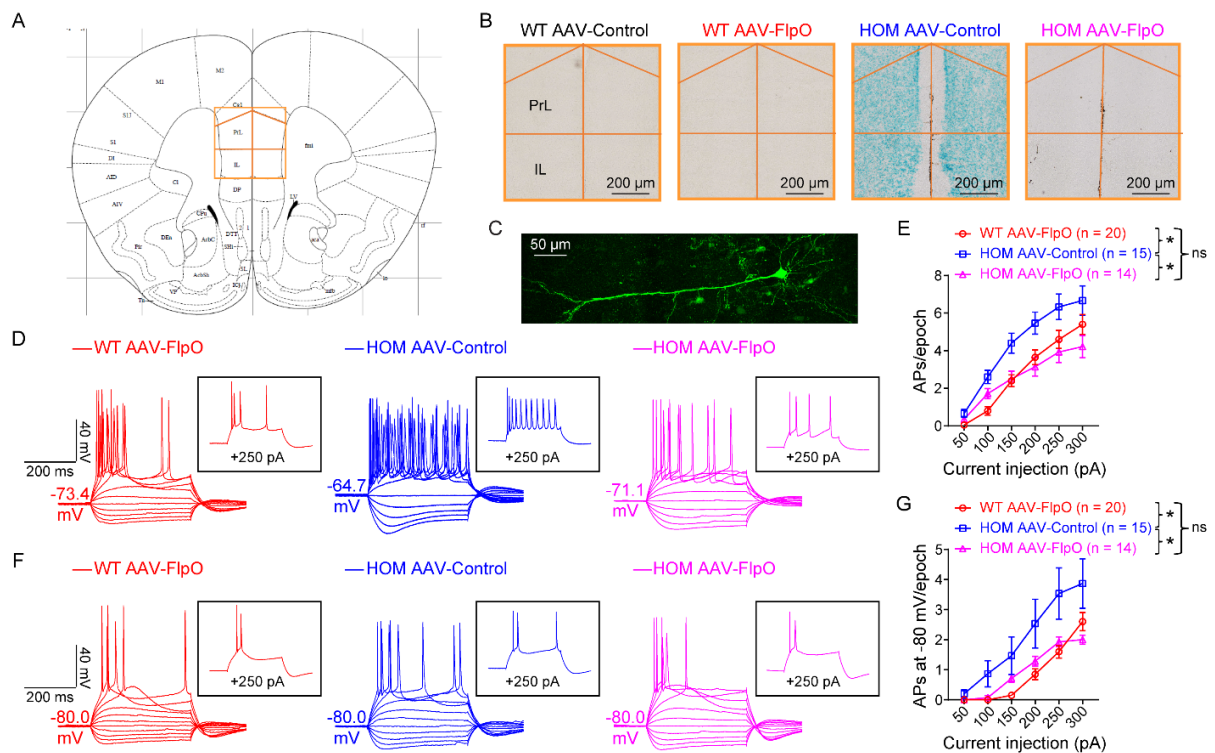


Figure S2. Elevated neuronal firings of layer V pyramidal cells in the mPFC are reversible by FlpO-mediated rescue in adult *Nav1.2*-deficient mice. Related to Figure 2.

(A-B) *LacZ* staining of coronal brain slices containing mPFC from WT and *Scn2a^{gt/gt}* (HOM) mice, which were systemically administered with AAV-Control or AAV-FlpO. PrL, prelimbic cortex; IL, infralimbic cortex.

(C) A typical layer V pyramidal neuron in the mPFC was labeled by neurobiotin. Scale bar, 50 μ m.

(D) Representative current-clamp recordings of pyramidal cells from WT mice transduced with AAV-FlpO (red), HOM mice transduced with AAV-Control (blue), and HOM mice transduced with AAV-Control (magenta) at the RMP. A series of 400-ms hyperpolarizing and depolarizing steps in 50-pA increments were applied to produce the traces. Inset: representative trace in response to +250 pA current injection.

(E) The average number of APs generated in response to depolarizing current pulses at the RMP. Unpaired two-tailed non-parametric Mann-Whitney *U*-test for each current pulse: ns, no significance, $p > 0.05$; * $p < 0.05$.

(F) Representative current-clamp recordings of layer V pyramidal cells in the mPFC from WT transduced with AAV-FlpO (red), HOM transduced with AAV-Control (blue) and HOM transduced with AAV-Control (magenta) at a fixed membrane potential of -80 mV. A series of 400-ms hyperpolarizing and depolarizing steps in 50-pA increments were applied to produce the traces. Inset: representative trace in response to +250 pA current injection.

(G) The average number of APs generated in response to depolarizing current pulses at -80 mV. Unpaired two-tailed non-parametric Mann-Whitney *U*-test for each current pulse: ns, no significance, $p > 0.05$; * $p < 0.05$. Data were shown as mean \pm SEM.

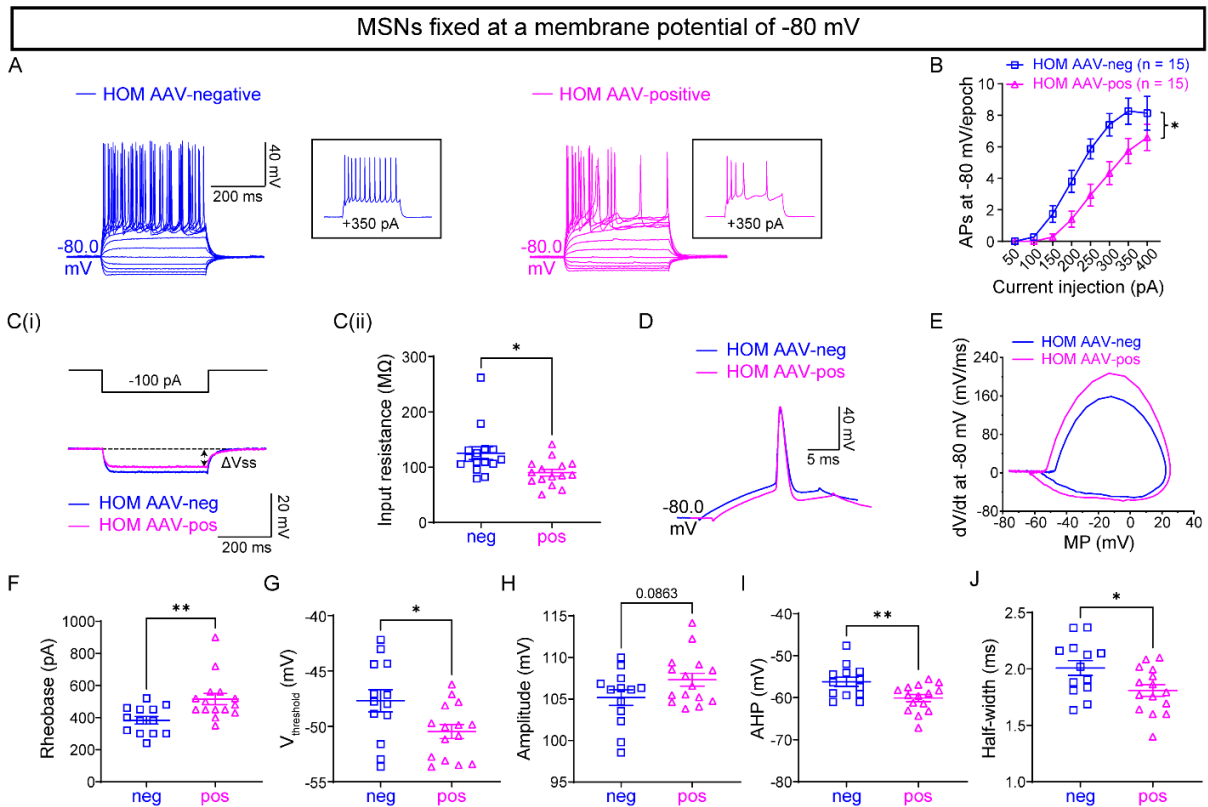


Figure S3. Ex vivo recordings of MSNs at a fixed membrane potential of -80 mV in adult *Nav1.2*-deficient mice with a dilute AAV-FlpO-mCherry injection. Related to Figure 3.

(A) Representative current-clamp recordings of MSNs with AAV-negative (blue) or with AAV-FlpO-positive (magenta) in *Scn2a^{gt/gt}* (HOM) mice were obtained at a fixed membrane potential of -80 mV. A series of 400-ms hyperpolarizing and depolarizing steps in 50-pA increments were applied to produce the traces. Inset: representative trace in response to +350 pA current injection.

(B) The average number of APs generated in response to depolarizing current pulses. Unpaired two-tailed non-parametric Mann-Whitney *U*-test for each current pulse: **p* < 0.05.

(C*i*) Representative traces in response to 100 pA negative current injection. $V_{\text{steady-state}}$ (V_{ss}) is the voltage recorded at 0-10 ms before the end of the stimulus.

(C*ii*) Individuals and average input resistance values at -80 mV. Unpaired two-tailed Student's *t*-test: **p* < 0.05.

(D) Typical spikes of MSNs with AAV-negative (blue) or AAV-FlpO-positive (magenta) in HOM mice were obtained at a fixed membrane potential of -80 mV.

(E) Associated phase-plane plots at -80 mV.

(F-J) Individuals and average spike rheobase, voltage threshold, amplitude, AHP, and half-width values. Unpaired two-tailed Student's *t*-test: ns, no significance, *p* > 0.05; **p* < 0.05; ***p* < 0.01. Data were shown as mean ± SEM.

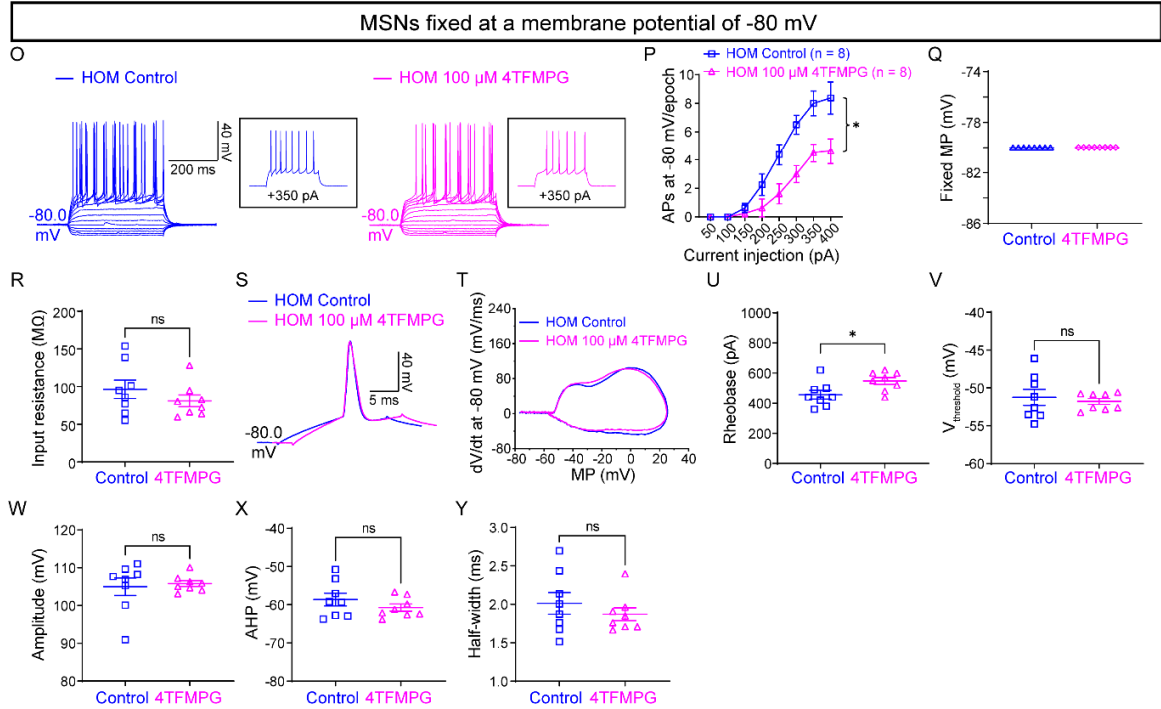
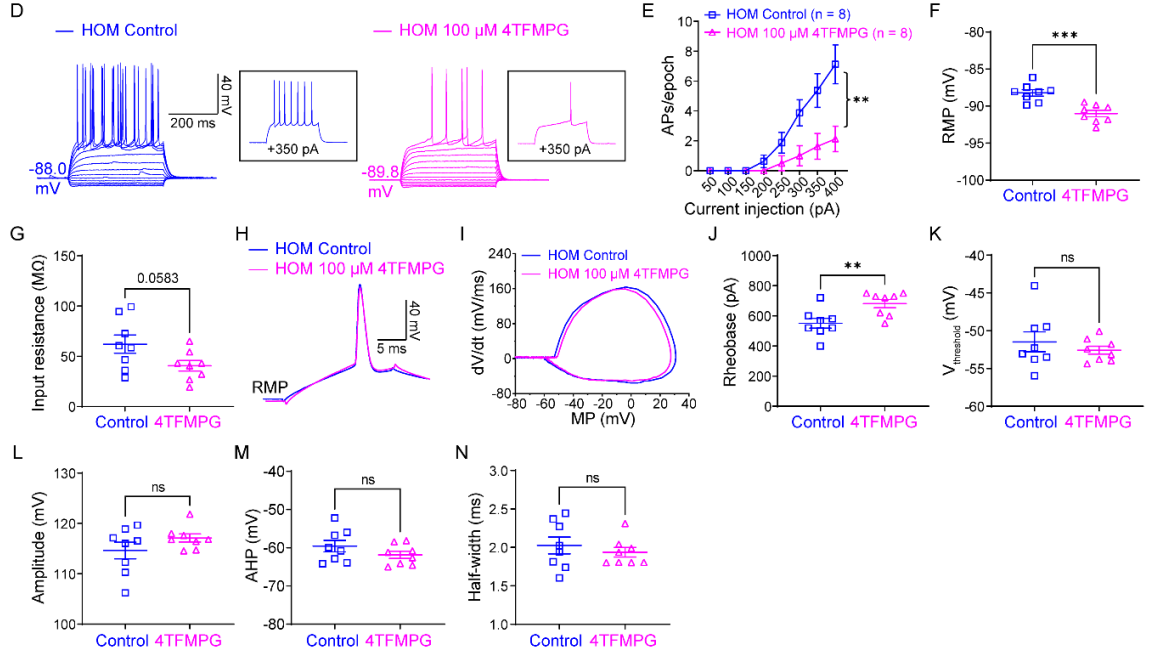
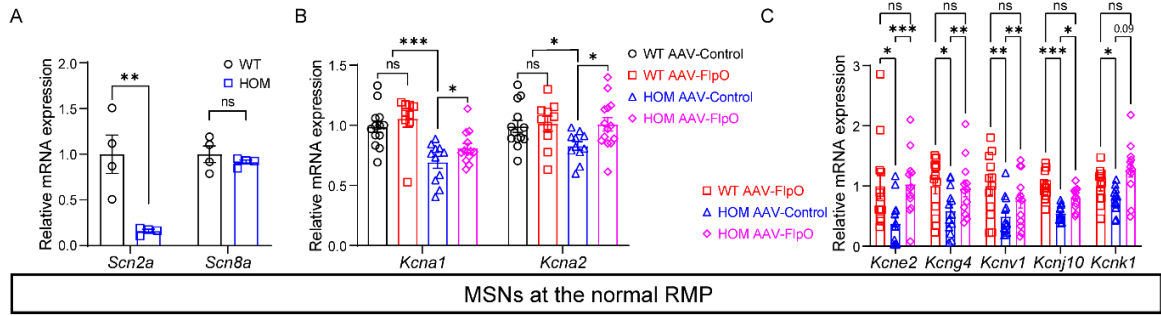


Figure S4. Specific activation of K_v1.1 channel by 4TFMPG reverses the elevated neuronal firings in adult Nav_{1.2}-deficient mice. Related to Figure 4.

(A) Quantitative (q)PCR analysis of *Scn2a* and *Scn8a* mRNA in the striatum samples from WT and *Scn2a^{gt/gt}* mice. Unpaired two-tailed Student's *t*-test for each group: ns, no significance, $p > 0.05$; ** $p < 0.01$.

(B) qPCR analysis of *Kcna1* and *Kcna2* mRNA in the striatum samples from WT and HOM mice transduced with AAV-Control or AAV-FlpO, showing that the downregulated mRNA levels of K_v1.1 and K_v1.2 were partially reversible by FlpO-mediated restoration of Nav_{1.2} expression in adult Nav_{1.2}-deficient mice. Unpaired two-tailed Student's *t*-test: ns, no significance, $p > 0.05$; * $p < 0.05$; *** $p < 0.001$.

(C) qPCR analysis of *Kcne2*, *Kcng4*, *Kcnv1*, *Kcnj10*, and *Kcnk1* mRNA in the striatum samples from WT and HOM mice transduced with AAV-Control or AAV-FlpO, showing that the downregulated mRNA levels of *Kcne2*, *Kcng4*, *Kcnv1*, *Kcnj10*, and *Kcnk1* were partially reversible by FlpO-mediated restoration of Nav_{1.2} expression in adult Nav_{1.2}-deficient mice. Unpaired two-tailed Student's *t*-test: ns, no significance, $p > 0.05$; * $p < 0.05$; ** $p < 0.01$; *** $p < 0.001$.

(D) Representative current-clamp recordings of MSNs from HOM slices perfused with aCSF (HOM Control, blue) and HOM slices perfused with aCSF containing 4TFMPG (HOM 100 μ M 4TFMPG, magenta) at the RMP. A series of 400-ms hyperpolarizing and depolarizing steps in 50-pA increments were applied to produce the traces. Inset: representative trace in response to +350 pA current injection.

(E) The average number of APs generated in response to depolarizing current pulses at the RMP. Unpaired two-tailed non-parametric Mann-Whitney *U*-test for each current pulse: ** $p < 0.01$.

(F) Individuals and average spike RMP values. Unpaired two-tailed Student's *t*-test: *** $p < 0.001$.

(G) Individuals and average input resistance values at the RMP. Unpaired *t*-test.

(H) Typical spikes of MSNs from HOM slices perfused with aCSF (HOM Control, blue) and HOM slices perfused with aCSF containing 4TFMPG (HOM 100 μ M 4TFMPG, magenta) were obtained at the RMP.

(I) Associated phase-plane plots.

(J-N) Individuals and average AP rheobase, voltage threshold, amplitude, AHP, and half-width values.

(O) Representative current-clamp recordings of MSNs from HOM slices perfused with aCSF (HOM Control, blue) and HOM slices perfused with aCSF containing 4TFMPG (HOM 100 μ M 4TFMPG, magenta) at a fixed membrane potential of -80 mV. A series of 400-ms hyperpolarizing and depolarizing steps in 50-pA increments were applied to produce the traces. Inset: representative trace in response to +350 pA current injection.

(P) The average number of APs generated in response to depolarizing current pulses at -80 mV. Unpaired two-tailed non-parametric Mann-Whitney *U*-test for each current pulse: * $p < 0.05$.

(Q) Fixed MP values for recording.

(R) Individuals and average input resistance values at -80 mV. Unpaired two-tailed Student's *t*-test: ns, no significance, $p > 0.05$.

(S) Typical spikes of MSNs from *Scn2a^{gt/gt}* slices perfused with aCSF (HOM Control, blue) and *Scn2a^{gt/gt}* slices perfused with aCSF containing 4TFMPG (HOM 100 μ M 4TFMPG, magenta) were obtained at a fixed membrane potential of -80 mV.

(T) Associated phase-plane plots.

(U-Y) Individuals and average AP rheobase, voltage threshold, amplitude, AHP, and half-width values. Unpaired two-tailed Student's *t*-test: ns, no significance, $p > 0.05$; * $p < 0.05$. Data were shown as mean \pm SEM.

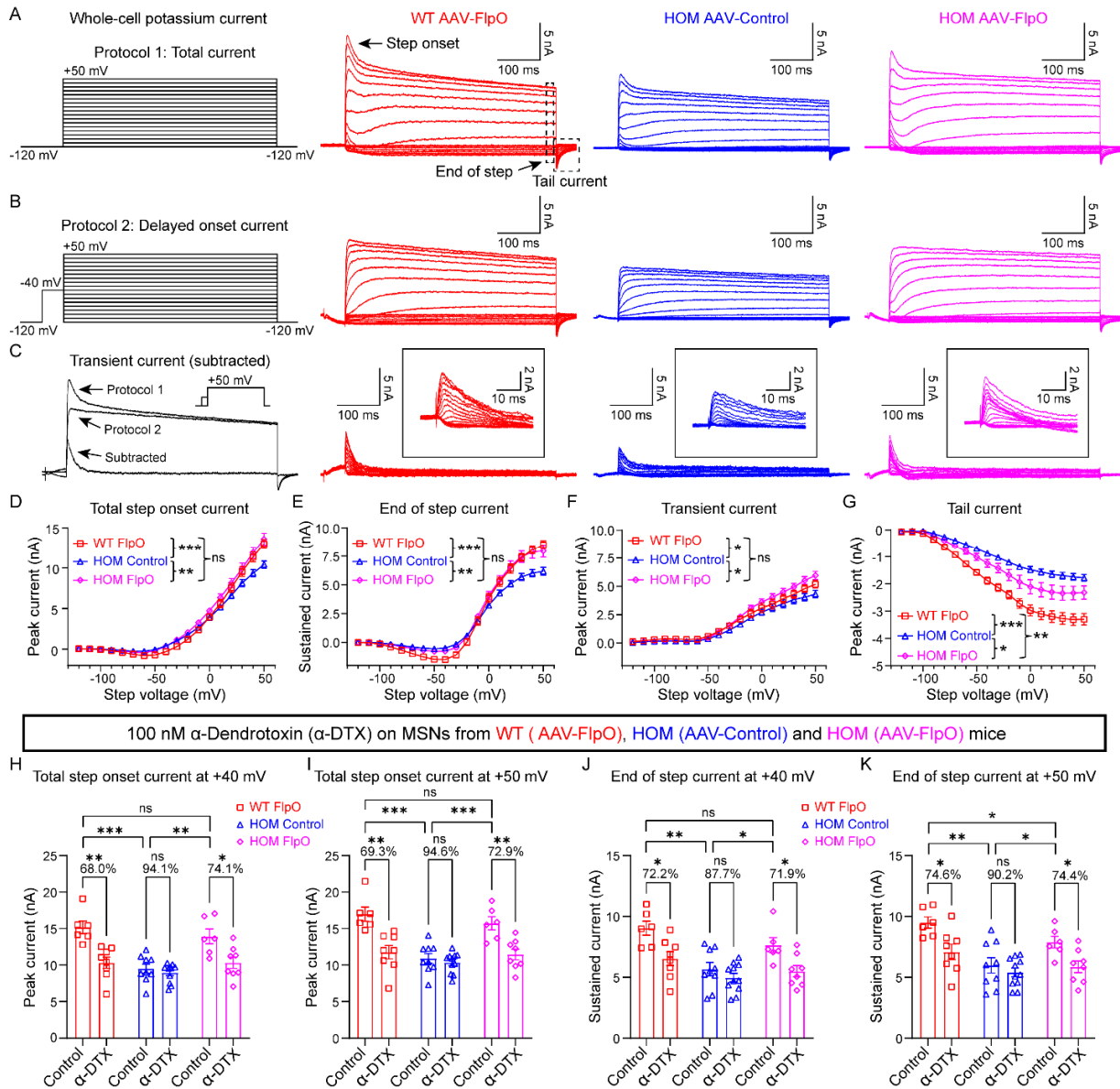


Figure S5. Reduced potassium currents in striatal MSNs are reversible by AAV-FlpO-mediated rescue in adult Nav1.2-deficient mice. Related to Figure 4.

(A-C) Representative whole-cell potassium currents of MSNs in slices from WT mice transduced with AAV-FlpO (red), HOM mice transduced with AAV-Control (blue), and HOM mice transduced with AAV-FlpO (magenta). Voltage steps were from -120 mV to +50 mV. Total currents were obtained from 500-ms steps followed by 100-ms pulse back to -120 mV. Step onset, end of step, and tail are shown. Delayed onset currents were measured following 50-ms prepulse to -40 mV. Transient currents were calculated by subtracting delayed onset currents from the total currents. Voltage-gated Ca^{2+} channels were not blocked to allow for activation of Ca^{2+} -dependent K^+ channels.

(D-G) Summary for total step onset current, end of step current, transient current, and tail current, respectively. $n = 17$ neurons in the group of WT mice transduced with AAV-FlpO (red), $n = 23$

neurons in the group of HOM mice transduced with AAV-Control (blue), and $n = 15$ neurons in the group of HOM mice transduced with AAV-FlpO (magenta). Two-way ANOVA with repeated measures: ns, no significance, $p > 0.05$; * $p < 0.05$; ** $p < 0.01$; *** $p < 0.001$.

(H-K) 100 nM α -Dendrotoxin (α -DTX, blocker of K_{V1}) reduces ~30% whole-cell total potassium currents of MSNs in slices from WT mice transduced with AAV-FlpO (red), ~10% total currents from HOM mice transduced with AAV-Control (blue), and ~30% total currents from HOM mice transduced with AAV-Control (magenta). Summaries for total step onset currents (**H**, at +40 mV; **I**, at +50 mV) and end of step current (**J**, at +40 mV; **K**, at +50 mV) were shown respectively. Values are the percentage for peak currents recorded in slices perfused 100 nM α -DTX compared to each corresponding control group. Two-way ANOVA with repeated measures: ns, no significance, $p > 0.05$; * $p < 0.05$; ** $p < 0.01$; *** $p < 0.001$. Data were shown as mean \pm SEM.

Table S1. Oligonucleotides used in this study. Related to STAR Methods.

REAGENT or RESOURCE	SOURCE	IDENTIFIER
Oligonucleotides		
<i>Scn2a</i> -Forward Sequence (5' -> 3') ATTTTCGGCTCATTCTTCACACT	https://pga.mgh.harvard.edu/primerbank/	PrimerBank ID: ID840654a1
<i>Scn2a</i> -Reverse Sequence (5' -> 3') GGGCGAGGTATCGGTTTTTGT	https://pga.mgh.harvard.edu/primerbank/	PrimerBank ID: 840654a1
<i>Scn8a</i> -Forward Sequence (5' -> 3') GCAAGCTCAAGAAACCACCC	https://pga.mgh.harvard.edu/primerbank/	PrimerBank ID: 951126a1
<i>Scn8a</i> -Reverse Sequence (5' -> 3') CCGTAGATGAAAGGCAAACCTCT	https://pga.mgh.harvard.edu/primerbank/	PrimerBank ID: 951126a1
<i>Kcna1</i> -Forward Sequence (5' -> 3') CCCTACCCGAGAAGGAGTACC	https://pga.mgh.harvard.edu/primerbank/	PrimerBank ID: 119395751c2
<i>Kcna1</i> -Reverse Sequence (5' -> 3') GGATGACCATGACCGACACAA	https://pga.mgh.harvard.edu/primerbank/	PrimerBank ID: 119395751c2
<i>Kcna2</i> -Forward Sequence (5' -> 3') GCACCCACAAGACACCTATGA	https://pga.mgh.harvard.edu/primerbank/	PrimerBank ID: 31543024a1
<i>Kcna2</i> -Reverse Sequence (5' -> 3') GTCTCTGGGAAGTGGGCTAAG	https://pga.mgh.harvard.edu/primerbank/	PrimerBank ID: 31543024a1
<i>Kcne2</i> -Forward Sequence (5' -> 3') CACATTAGCCAATTTGACCCAGA	https://pga.mgh.harvard.edu/primerbank/	PrimerBank ID: 19882205a1
<i>Kcne2</i> -Reverse Sequence (5' -> 3') GAACATGCCGATCATCACCAT	https://pga.mgh.harvard.edu/primerbank/	PrimerBank ID: 19882205a1
<i>Kcng4</i> -Forward Sequence (5' -> 3') CGCAGCCATGAGGAGATCAC	https://pga.mgh.harvard.edu/primerbank/	PrimerBank ID: 28076887a1
<i>Kcng4</i> -Reverse Sequence (5' -> 3') GCCAGGAAGCTCACGATCAC	https://pga.mgh.harvard.edu/primerbank/	PrimerBank ID: 28076887a1
<i>Kcnv1</i> -Forward Sequence (5' -> 3') ATTGCGCTCACTTGGGATGAC	https://pga.mgh.harvard.edu/primerbank/	PrimerBank ID: 28460685a1
<i>Kcnv1</i> -Reverse Sequence (5' -> 3') GTGAAGTTGTATCAGGAATGCT	https://pga.mgh.harvard.edu/primerbank/	PrimerBank ID: 28460685a1
<i>Kcnj10</i> -Forward Sequence (5' -> 3') ACCTCAAGGATCTATGGACGAC	https://pga.mgh.harvard.edu/primerbank/	PrimerBank ID: 87116685c2
<i>Kcnj10</i> -Reverse Sequence (5' -> 3') AGCTACCAGATACCACACCAC	https://pga.mgh.harvard.edu/primerbank/	PrimerBank ID: 87116685c2
<i>Kcnk1</i> -Forward Sequence (5' -> 3') GAGGAGCTGCCTTATGAGGAC	https://pga.mgh.harvard.edu/primerbank/	PrimerBank ID: 6680538a1

<i>Kcnk1</i> -Reverse Sequence (5' -> 3') TCCCAATTCCAATTTCCCGAG	https://pga.mgh.harvard.edu/primerbank/	PrimerBank ID: 6680538a1
<i>Actb</i> -Forward Sequence (5' -> 3') GGCTGTATTCCCCTCCATCG	https://pga.mgh.harvard.edu/primerbank/	PrimerBank ID: 6671509a1
<i>Actb</i> -Reverse Sequence (5' -> 3') GGCTGTATTCCCCTCCATCG	https://pga.mgh.harvard.edu/primerbank/	PrimerBank ID: 6671509a1
<i>Gapdh</i> -Forward Sequence (5' -> 3') AGGTCGGTGTGAACGGATTTG	https://pga.mgh.harvard.edu/primerbank/	PrimerBank ID: 6679937a1
<i>Gapdh</i> -Reverse Sequence (5' -> 3') TGTAGACCATGTAGTTGAGGTCA	https://pga.mgh.harvard.edu/primerbank/	PrimerBank ID: 6679937a1
<i>Scn2a1L</i> -Forward Sequence (5' -> 3') GAGGCAAAGAATCTGTACTGTGGGG	https://www.nlac.narl.org.tw/RMRC/upload/deposit_file/201704241012510.pdf	RMRC13300
<i>Scn2a1L</i> -Reverse Sequence (5' -> 3') GACGCCTGTGAATAAAACCAAGGAA	https://www.nlac.narl.org.tw/RMRC/upload/deposit_file/201704241012510.pdf	RMRC13300