

Supplemental information

Improvement of sensory deficits in Fragile X mice by increasing cortical interneuron activity after the critical period.

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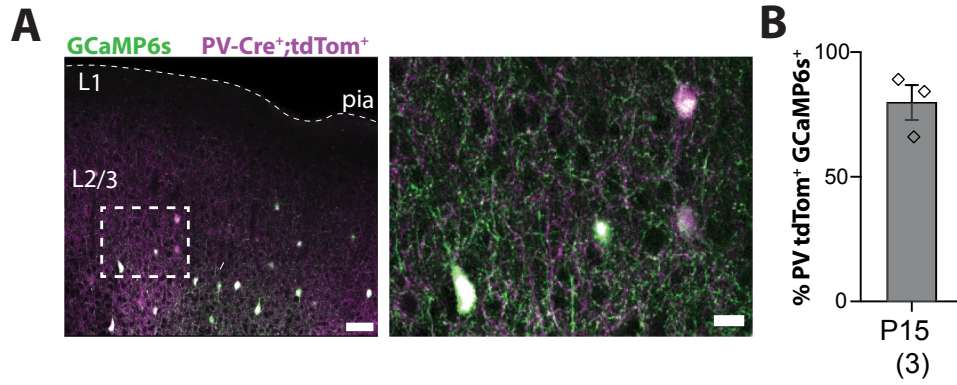
Supplementary figures: 13

Supplementary table: 1

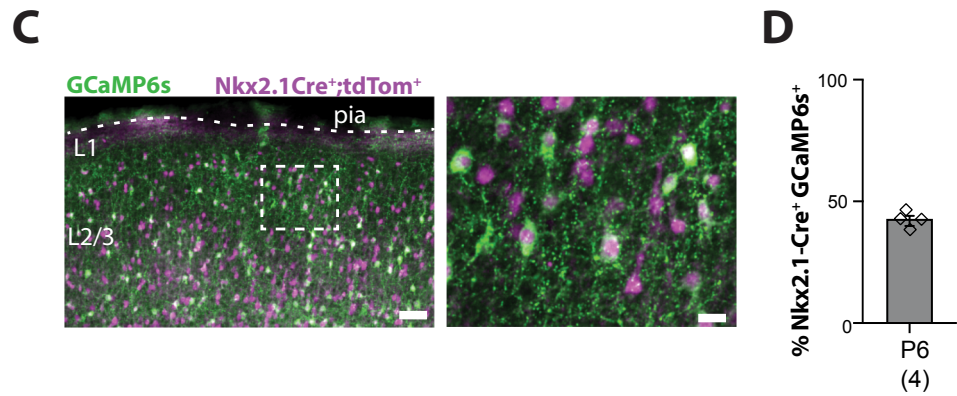
Supplementary videos: 2

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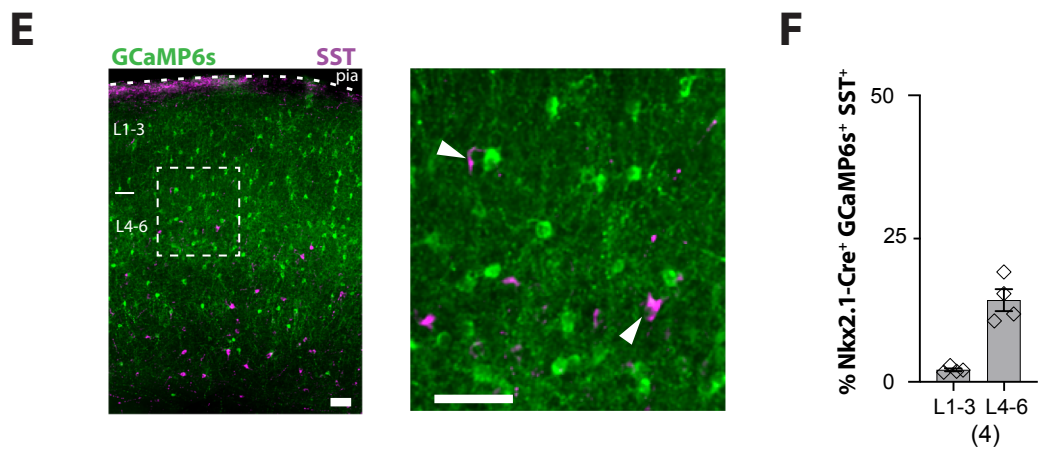
Percentage of PV-tdTom⁺ cells that express GCaMP6S at P15



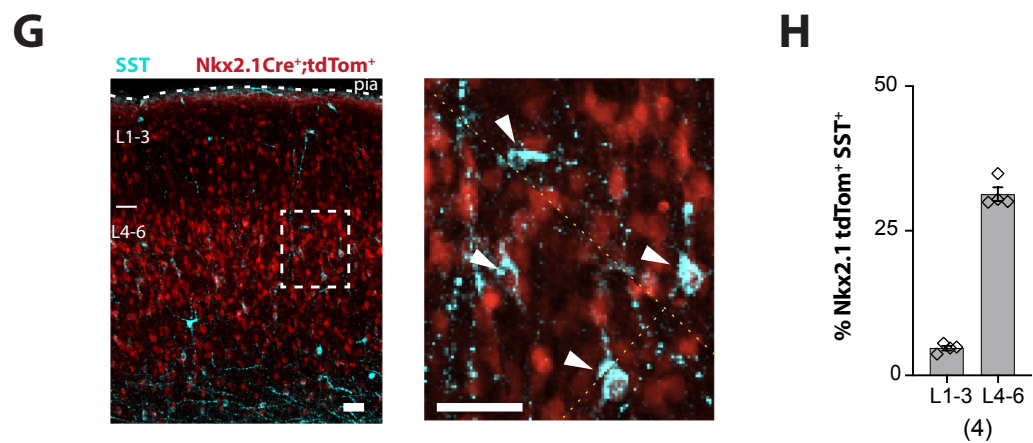
Percentage of Nkx2.1-tdTom⁺ cells that express GCaMP6S at P6



Percentage of Nkx2.1⁺-GCaMP6s⁺ cells that express SST at P6



Percentage of Nkx2.1-tdTom⁺ cells that express SST at P6

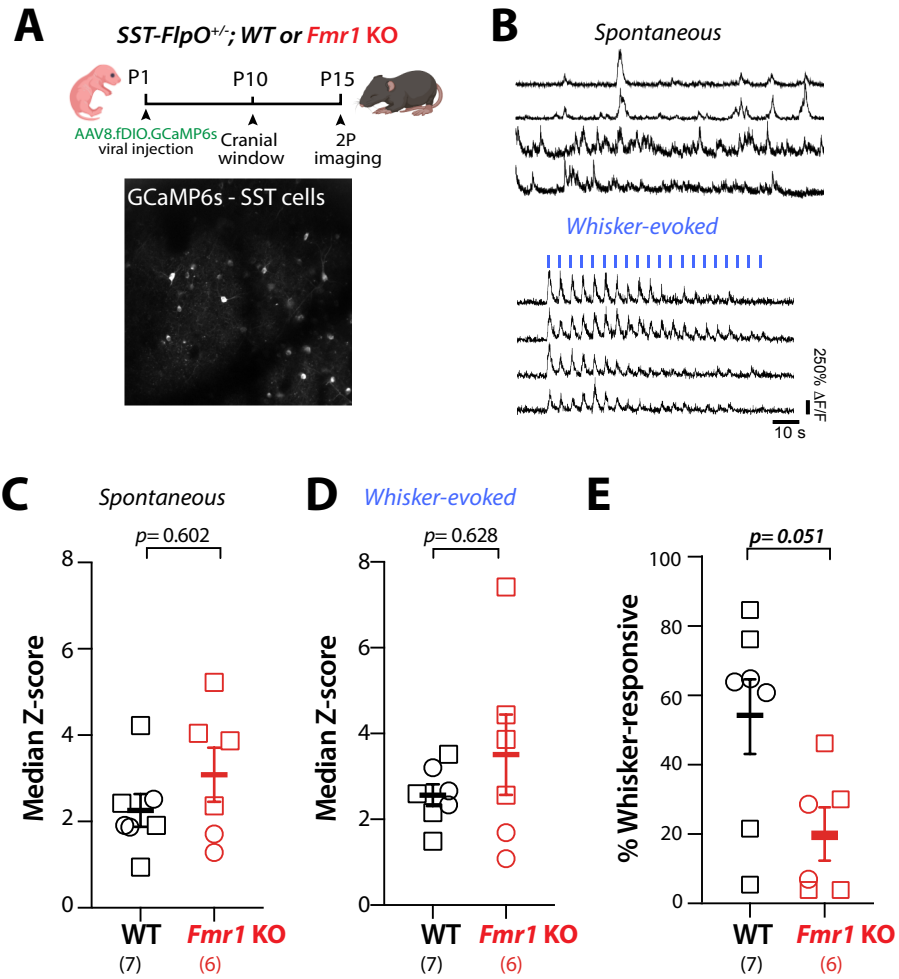


Supplementary Fig. S1

Supplementary Fig. S1: Efficacy of viral strategy to express GCaMP6s in different IN populations in P15 mice. (Related to Fig. 1)

- A. Example field of view of PV-Cre⁺-tdTom⁺ (magenta) expressing GCaMP6s (green) in PV-Cre;Ai14^{+/-}; *Fmr1* KO mice at P15.
- B. An injection of rAAV-CAG-DIO-GCaMP6s virus in S1 at P10 resulted in a high proportion of PV-INs also expressing GCaMP6s by P15 (79.8± 7%, n=3 mice). This pertains to Fig. 1A-F. Scale= 20 μm.
- C. Example field of view of Nkx2.1-Cre⁺-tdTom⁺ (magenta) expressing GCaMP6s (green) in Nkx2.1-Cre;Ai14^{+/-} *Fmr1* KO mice at P15.
- D. An injection of rAAV-CAG-DIO-GCaMP6s virus in S1 at P1 resulted in 43.4± 1.6% of Nkx2.1-INs also expressing GCaMP6s (n=4 mice). Note that the density of Nkx2.1-Cre⁺-GCaMP6s⁺ cells is comparable to other reports¹. This pertains to Fig. 1G-L. Scale= 20 μm.
- E. Representative image of Nkx2.1-INs expressing GCaMP6s (arrowheads, following injection of rAAV-CAG-DIO-GCaMP6s virus in S1 at P1) and immunostained for SST at P6. Scale= 50 μm.
- F. Percentage of Nkx2.1-Cre⁺;GCaMP6s⁺ INs that co-express SST at P6 (2.1± 0.2% in L1-L3 and 14.2± 1.9% in L4-6).
- G. Representative image of Nkx2.1-Cre⁺;tdTom⁺ INs immunostained for SST at P6. Arrowheads depicts the co-expressing cells. Yellow dotted line indicates uneven stitching by Zeiss Apotome. Scale= 50 μm.
- H. Percentage of Nkx2.1-Cre⁺;tdTom⁺ INs that co-express SST at P6 (4.7± 0.3% in L1-L3 and 31.3± 1.2% in L4-6).

SST-IN activity in P15 WT and *Fmr1* KO mice

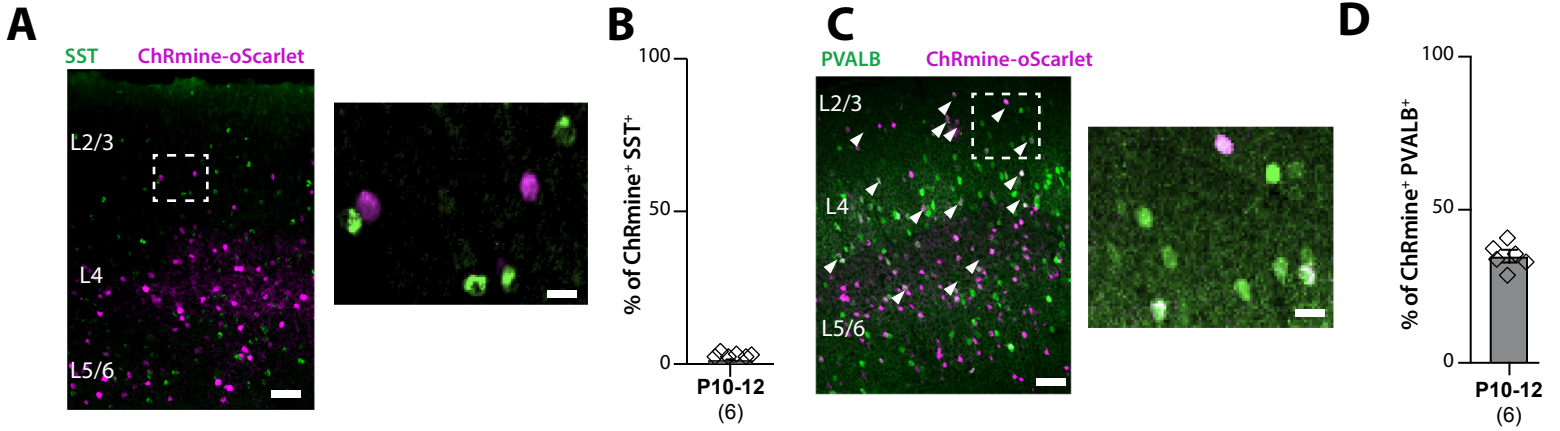


Supplementary Fig. S2

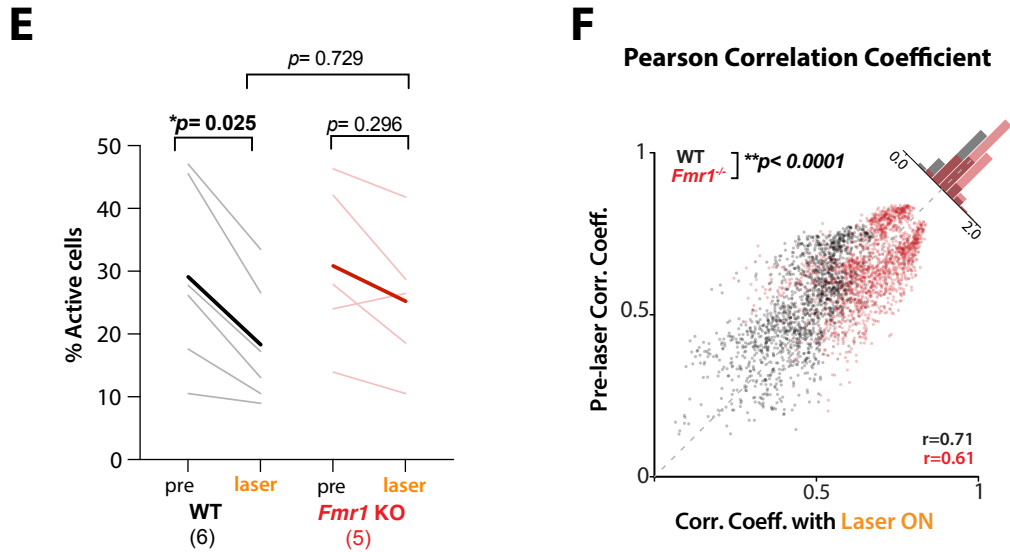
Supplementary Fig. S2: Activity of SST INs in S1 of P15 *Fmr1* KO mice is not different from WT controls. (Related to Fig. 1)

- A. Top: Cartoon of experimental design for calcium imaging recordings in SST-INs. Bottom: Example field of view of SST-INs expressing GCaMP6s in SST-FlpO mice at P15.
- B. Example traces of calcium transients for spontaneous and whisker-evoked activity in both SST-FlpO ; WT and SST-FlpO; *Fmr1* KO mice (we show traces from 2 different SST-INs of 2 different animals). The vertical blue bars represent the 20 whisker stimulations.
- C. Mean Z-scores for spontaneous activity of SST-INs in *Fmr1* KO and WT mice at P15 (2.56 ± 0.38 for WT vs. 3.08 ± 0.62 for *Fmr1* KO, n=6 and 7 mice, respectively; $p=0.602$, MW t-test).
- D. Mean Z-scores for whisker-evoked activity of SST-INs in *Fmr1* KO and WT mice at P15 (2.56 ± 0.25 for WT vs. 3.51 ± 0.94 for *Fmr1* KO, $p=0.628$, MW t-test).
- E. Percentage of whisker-responsive SST-INs in *Fmr1* KO and WT mice at P15 ($53.8 \pm 11.0\%$ for WT vs. $19.6 \pm 7.6\%$ for *Fmr1* KO, $p= 0.051$, M-W t-test).

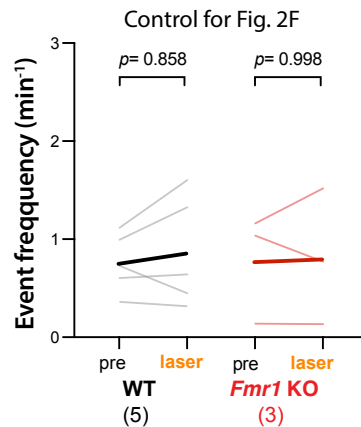
Percentage of ChRmine⁺ cells that express PVALB or SST in *Nkx2.1-Cre^{+/+};SST-FlpO^{+/-}* mice at P10-12



Effect of optogenetic activation of *Nkx2.1*⁺ INs on network activity at P10 (excitatory cells)



G Pyramidal cell activity in mCherry control mice

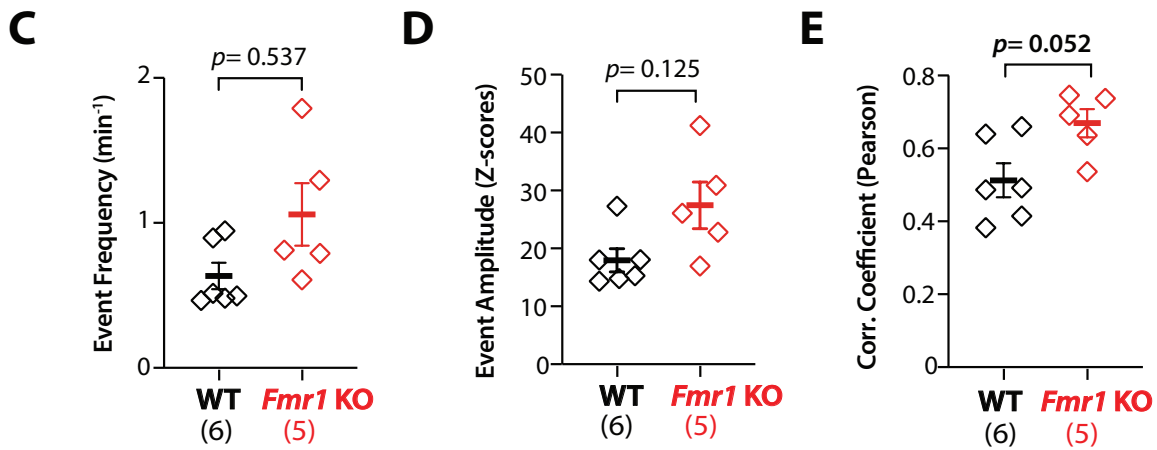
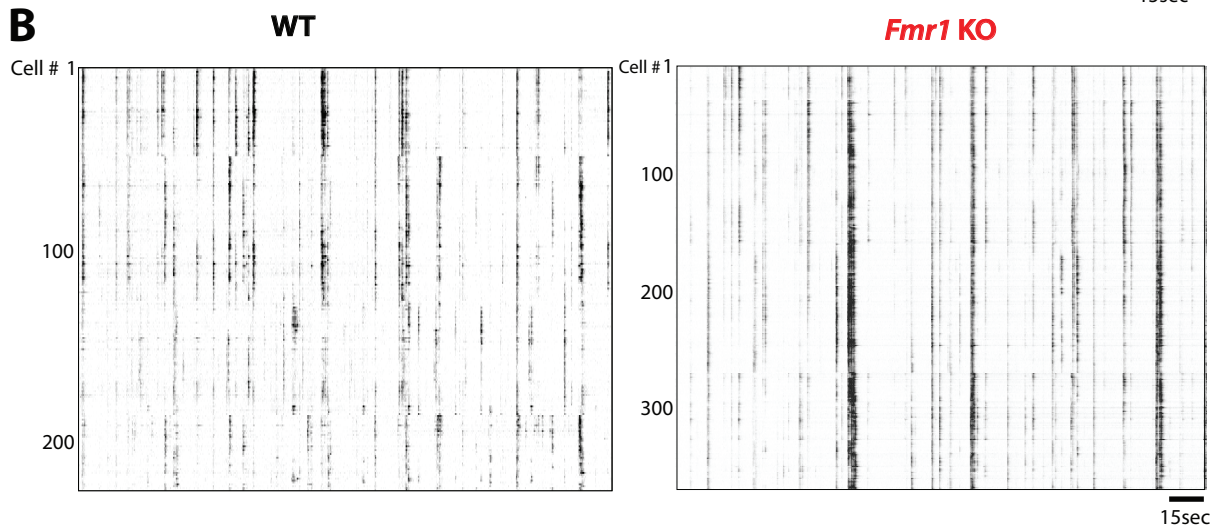
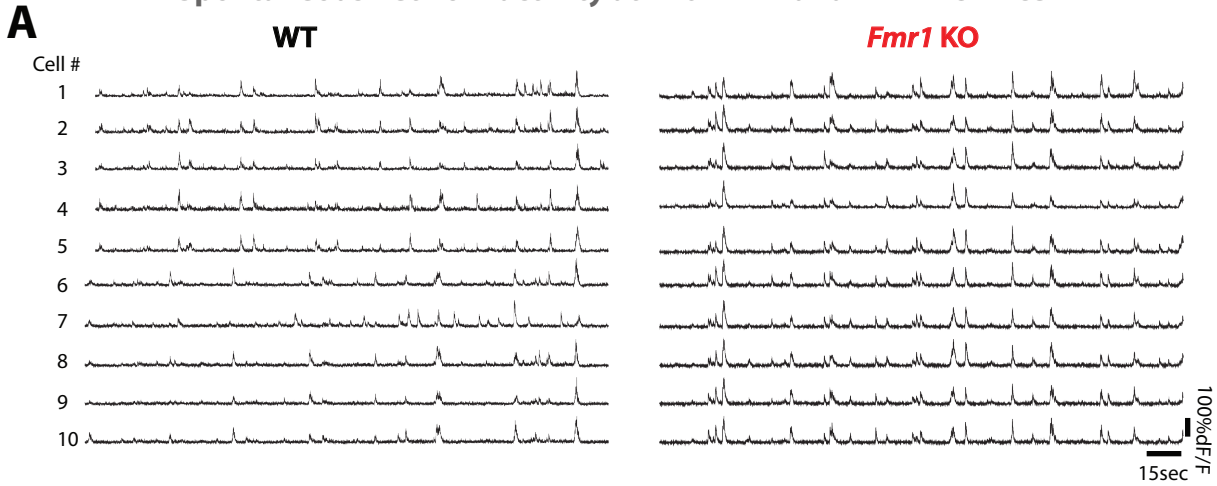


Supplementary Fig. S3

Supplementary Fig. S3: Controls for optogenetic experiments to stimulate Nkx2.1-INs in an attempt to modulate Pyr neurons in neonatal *Fmr1* KO mice at P10. (Related to Fig. 2)

- A. Example field of view in S1 of Nkx2.1-Cre;Sst-FlpO; *Fmr1* KO expressing ChRmine (magenta), using AAV-EF1-CreON/FlpOFF-ChRmine-oScarlet- virus) and immunostained for SST (green).
- B. Quantification of ChRmine⁺ cells that also express either PVALB or SST (by immunohistochemistry). As expected almost none of the ChRmine-labeled cells were SST immunoreactive ($1.2 \pm 0.4\%$, n= 6 mice), confirming the intersectional strategy worked as intended to target Nkx2.1⁺ but SST-FlpO⁺ cells at P10.
- C. Example field of view in S1 of Nkx2.1-Cre;Sst-FlpO; *Fmr1* KO expressing ChRmine (same as in b) and immunostained for SST (green). White arrowheads indicate double labeled cells.
- D. Quantification of ChRmine⁺ cells that are also immunoreactive for PVALB. Overall, $34.9 \pm 2.1\%$ the ChRmine-labeled also expressed PVALB (n= 6 mice). This is consistent with the fact that at this developmental stage (P10-P12) not all future PV-IN have started expressing PVALB.
- E. The percentage of active Pyr cells is significantly reduced upon laser stimulation of ChRmine-expressing future PV-INs in WT mice but is unchanged in *Fmr1* KO mice (29.1 ± 6.0 pre laser vs. 18.3 ± 4.0 laser On, p=0.025 in WT mice vs. 30.9 ± 5.9 pre laser vs. 25.2 ± 5.2 laser On in *Fmr1* KO mice, p=0.296, two-way ANOVA post-hoc Tukey).
- F. Change in pairwise correlation coefficients after optogenetic Nkx2.1-IN activation in Nkx2.1-Cre;Sst-FlpO mice for individual pairs of neurons (Spearman r=0.71, n=1,449 neuron pairs from n=6 WT mice; Spearman r=0.61, n=1,842 neuron pairs from n=5 *Fmr1*^{KO} mice). Note that correlation coefficients are significantly reduced by laser stimulation in WT mice, but not in *Fmr1* KO mice, as shown by the frequency distribution (p<0.001, Kolmogorov-Smirnov test).
- G. Mean frequency of Pyr cell calcium transients remained unchanged upon optogenetic stimulation in Nkx2.1-Cre;Sst-FlpO WT and *Fmr1* KO mice that do not express the opsin ChRmine (mCherry controls; 0.67 ± 0.13 pre laser vs. 0.86 ± 0.25 laser On, p=0.312 in WT mice; 0.78 ± 0.32 pre laser vs. 0.81 ± 0.4 laser On, p=0.500, n=5 WT and n=3 *Fmr1* KO , two-way ANOVA, post-hoc Tukey).

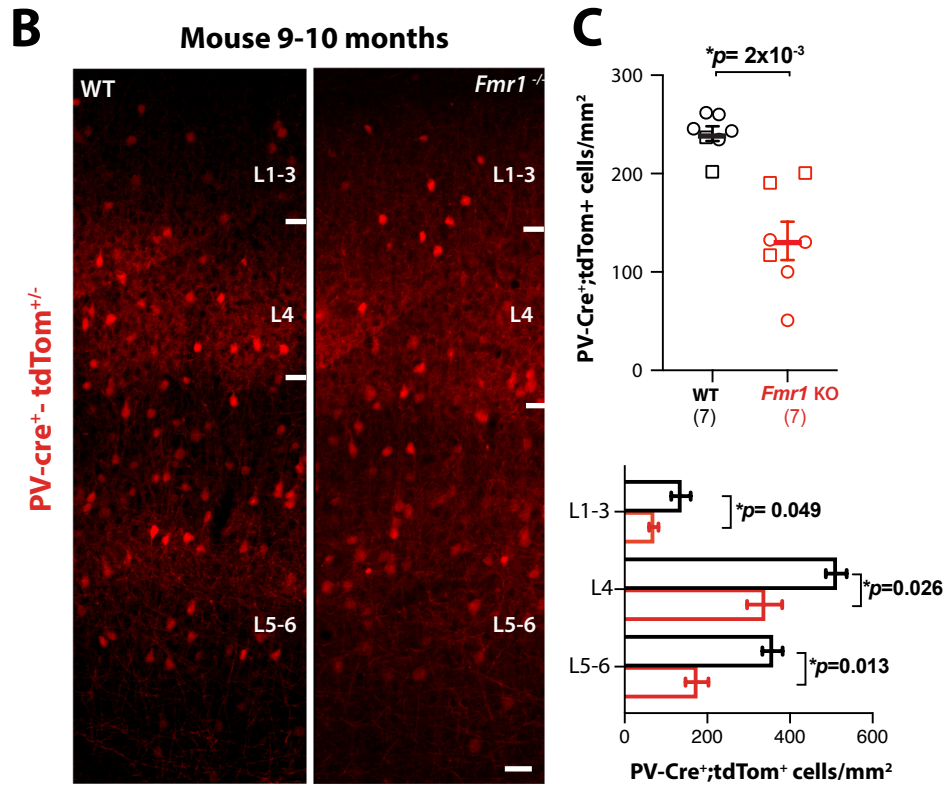
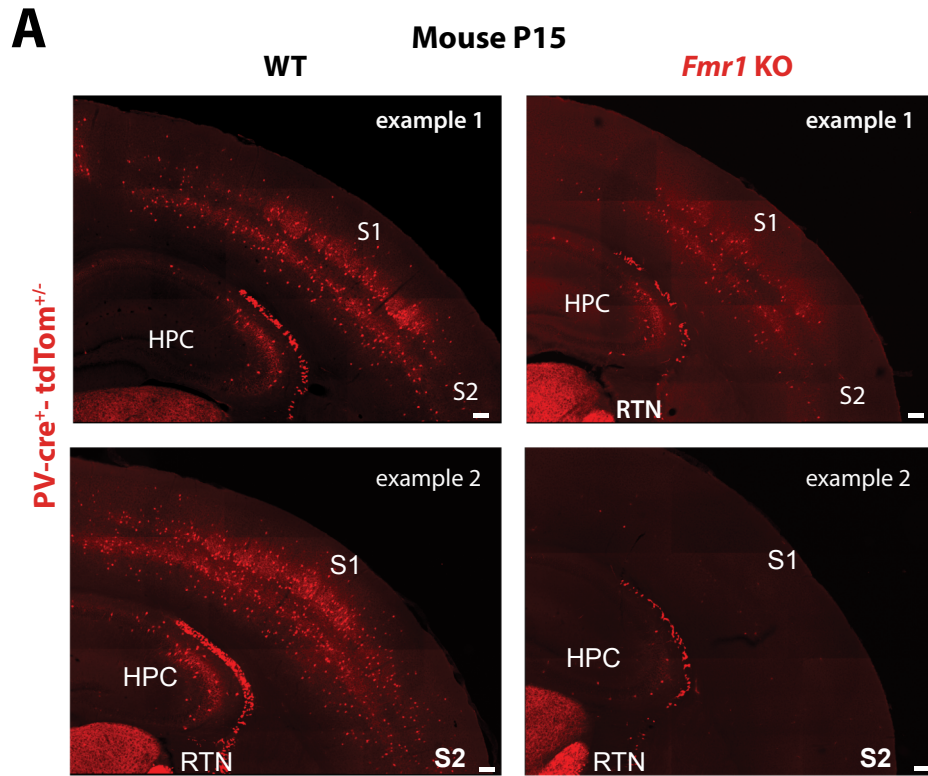
Spontaneous network activity at P10 in WT and *Fmr1* KO mice



Supplementary Fig. S4

Supplementary Fig. S4: Pyr cell activity at P10 in *Fmr1* KO is hypersynchronous. (Related to Fig. 2)

- A. Representative calcium traces of 10 example Pyr neurons in S1 of P10 WT and *Fmr1*^{KO}.
- B. Raster plots of Pyr cell activity in a representative WT and *Fmr1* KO mouse. Note the higher synchrony of network *Fmr1* KO.
- C. Mean frequency of Pyr cell calcium transients is not significantly different in WT and *Fmr1* KO mice (0.63 ± 0.09 in WT mice; 1.60 ± 0.21 , $n=6$ WT and $n=5$ *Fmr1* KO mice, $p=0.537$ MW test).
- D. Mean amplitude of Pyr cell calcium transients is not significantly different in WT and *Fmr1* KO mice (23.6 ± 2.00 in WT mice; 27.5 ± 4.0 , $n=5$ WT and $n=6$ *Fmr1* KO mice, $p=0.125$, MW test).
- E. Mean pairwise correlation coefficients is higher in *Fmr1* KO mice as compared to WT (0.51 ± 0.05 in $n=6$ WT mice; 0.67 ± 0.04 , and $n=5$ *Fmr1* KO mice, $p=0.052$, MW test).



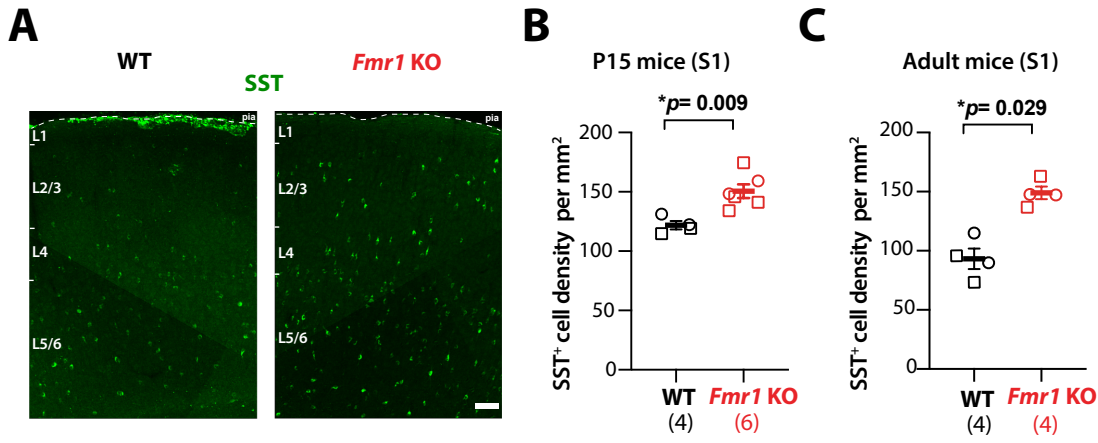
Supplementary Fig. S5

Supplementary Fig. S5: Reduced PV-IN density in *Fmr1* KO mice at P15 and at 9-10 months.

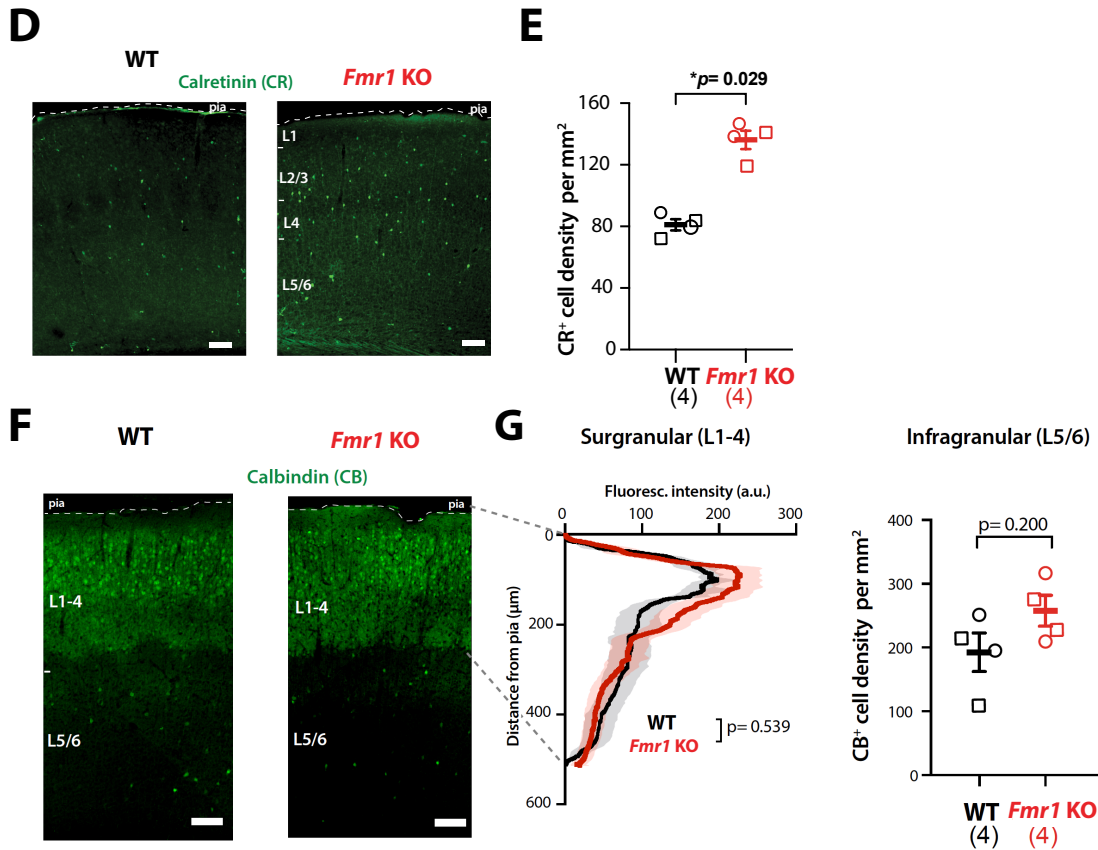
(Related to Fig. 3)

- A. Example coronal sections through S1 from *PV-Cre;tdTom^{+/-}* mice (WT and *Fmr1^{KO}*) at P15 showing the range of PV-IN density in *Fmr1* KO mice across the dorsal brain. Notably some *Fmr1* KO mice (example 2) exhibit a dramatic loss of PV-INs in neocortex and hippocampus (HPC), while other brain regions, such as the reticular thalamic nucleus (RTN) are much less affected. S2: secondary somatosensory cortex. Scale=100 μ m.
- B. Coronal sections through the barrel field of S1 from *PV-Cre;tdTom^{+/-}* mice (WT and *Fmr1* KO) at 9-10 months (corresponding approximately to age of human tissue in Fig. 3C-D). Scale= 50 μ m.
- C. Mean density of PV-tdTom⁺ INs in S1 is significantly lower in adult *Fmr1* KO mice (top), even across individual cortical layers (bottom). (all layers: 241 ± 8 cells/mm² for WT vs. 132 ± 2 for *Fmr1* KO; $p=0.002$, MW *t*-test; L2/3: 136 ± 23 vs. 69 ± 10 , $p=0.049$, L4: 512 ± 26 vs. 339 ± 43 , $p=0.026$; L5/6: 358 ± 24 , two-way ANOVA, post-hoc Holm-Sidak test, $p=0.013$, $n=7$ per genotype).

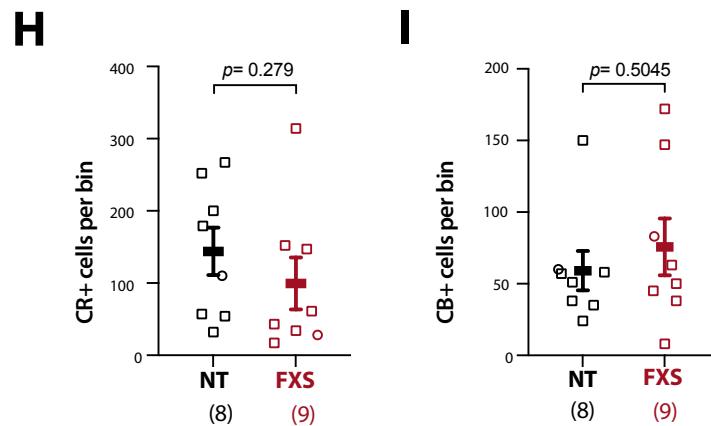
SST-IN density in 15 day-old and adult WT and *Fmr1* KO mice



Density of Calretinin- and Calbindin-expressing interneurons subtypes in adult adult WT and *Fmr1*^{-/-} mice

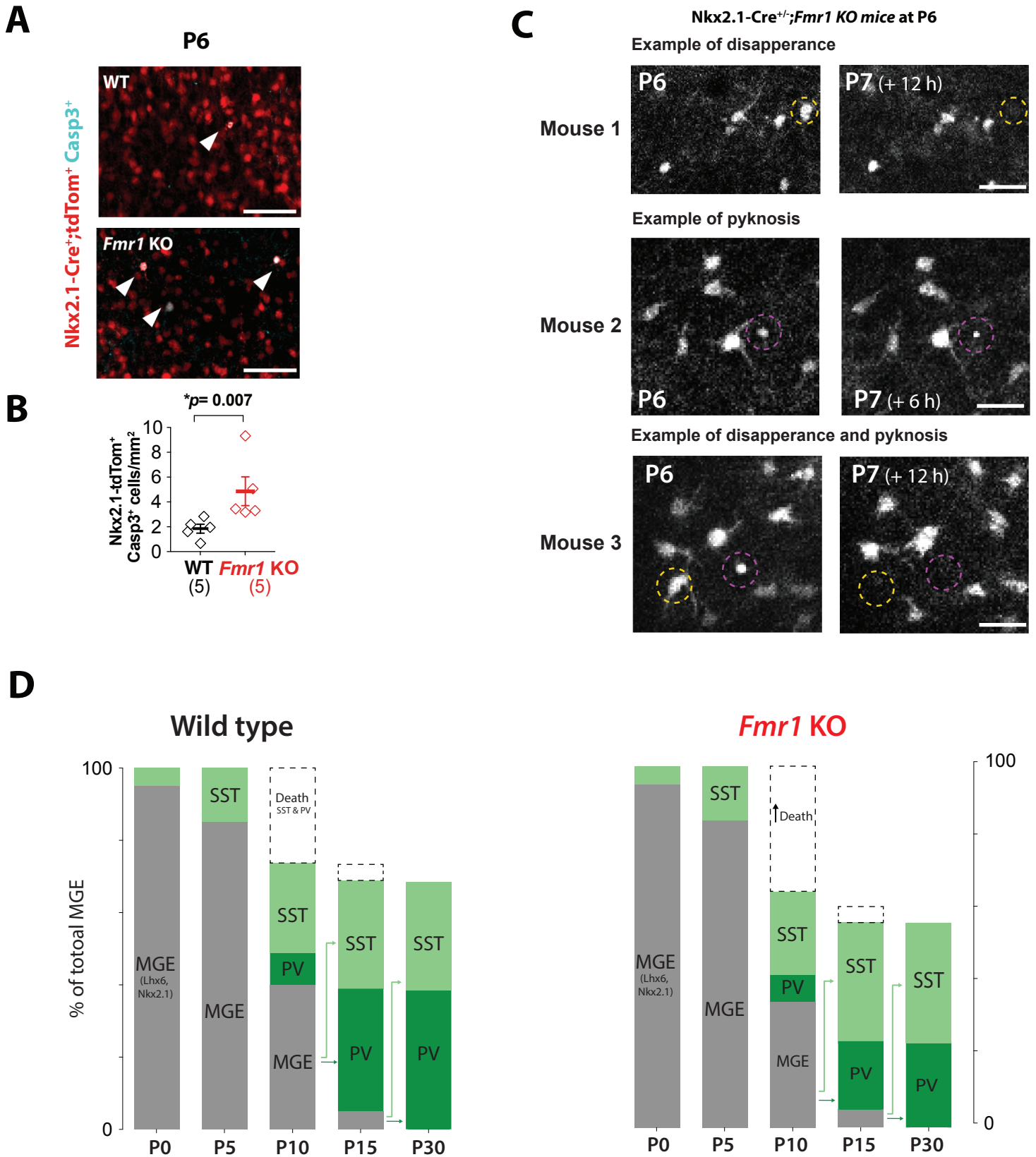


Density of Calretinin- and Calbindin-expressing interneurons in human samples



Supplementary Fig. S6: Density of SST, Calbindin, and Calretinin INs in FXS human cases and *Fmr1* KO mice. (Related to Fig. 3)

- A. Representative images of SST immunostaining in S1 in P15 WT and *Fmr1* KO mice.
- B. Quantification of total SST-immunoreactive cell density in the barrel field of S1 in WT and *Fmr1* KO mice in P15 (WT: 121.9 ± 3.7 cells/mm², *Fmr1* KO: 150.5 ± 5.9 , n= 4 and 6, respectively; p=0.009, MW test).
- C. Quantification of total SST-immunoreactive cell density in adult (4-5 months old) WT and *Fmr1* KO mice in and adult mice (WT: 93.14 ± 8.6 cells/mm², *Fmr1* KO: 149.1 ± 5.3 , n= 4 per group; p=0.029, MW test)
- D. Representative images of Calretinin immunostaining in S1 in adult WT and *Fmr1* KO mice.
- E. Quantification of Calretinin-immunoreactive INs from adult WT and *Fmr1* KO mice (81 ± 3.6 for WT vs. 136 ± 5.9 for *Fmr1* KO, n=4 mice per group; p=0.029 MW *t* test).
- F. Representative images of Calbindin immunostaining in S1 in adult WT and *Fmr1* KO mice. Note that the CB immunoreactivity is similar to previous reports.^{2,3}
- G. Quantification of Calbindin-immunoreactive INs in WT and *Fmr1* KO mice. The distribution and intensity of fluorescence is comparable in between WT and *Fmr1* KO mice in the supragranular layers (AUC= $48,036 \pm 704$ for WT vs. $42,093 \pm 578$ for *Fmr1* KO, n=4 mice per group; p=0.539, MW *t* test). The density of CB⁺ INS in infragranular layers is similar between WT and *Fmr1* KO mice (192.6 ± 30.2 for WT vs. 257.7 ± 24.2 for *Fmr1* KO; p=0.200, MW *t* test).
- H. Quantification of Calbindin-immunoreactive INs in human FXS and neurotypical control (NT) cases (59 ± 14 for NT vs. 76 ± 20 for FXS, n=8 and 9, respectively; p=0.279 MW *t* test).
- I. Quantification of Calretinin-immunoreactive INs in FXS human and control (NT) cases (144 ± 33 for NT vs. 100 ± 36 for FXS; p=0.279 MW *t* test).



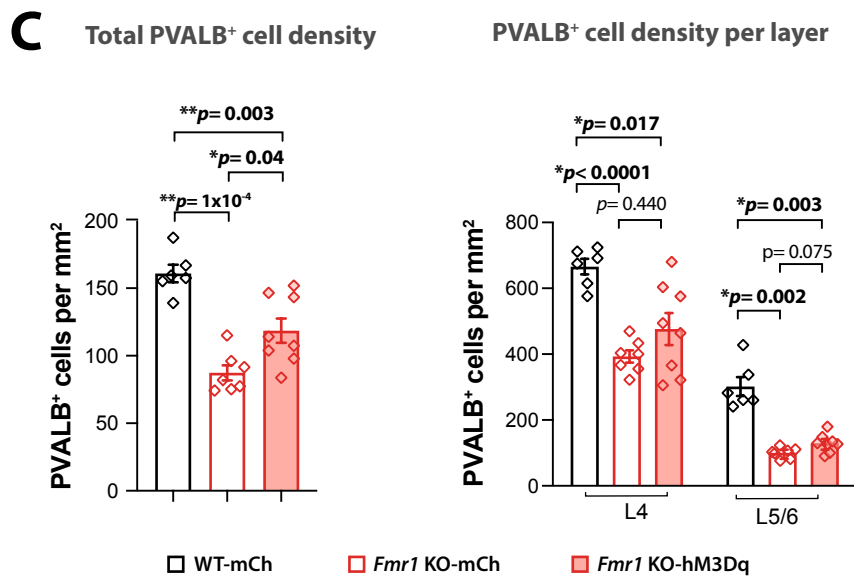
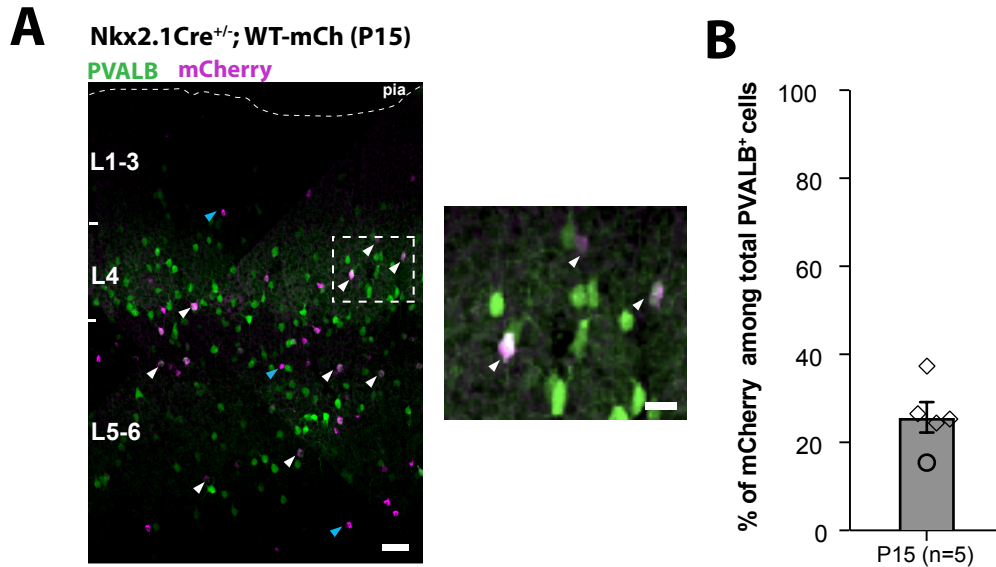
Supplementary Fig. S7

Supplementary Fig. S7: Evidence of apoptotic cell death of *Nkx2.1*-tdTom⁺ INs in neonatal *Fmr1* KO mice. (Related to Fig. 3)

- A. Coronal sections through S1 from *Nkx2.1-Cre* mice (WT and *Fmr1* KO) at P6 immunostained for cleaved Caspase-3 (cyan) showing rare double-labeled cells (arrowheads). Scale= 100 μ m.
- B. Mean density of *Nkx2.1*⁺-tdTom⁺-INs that co-express Caspase-3 was significantly higher in *Fmr1* KO mice (1.85 ± 0.35 cells/mm² for WT vs. 4.87 ± 1.16 for *Fmr1* KO; $p=0.007$, M-W test).
- C. Representative maximum intensity projection (~ 20 μ m) of *Nkx2.1*-tdTom⁺ IN image stacks acquired by in vivo 2P microscopy of the same FOV at P6 and 6-12h later in 3 example *Nkx2.1-Cre;Ai14*^{+/-}; *Fmr1* KO mice. Yellow dotted contours indicate *Nkx2.1*-tdTom⁺ INs present at P6 but absent at P7. Purple dotted contours indicate pyknotic *Nkx2.1*⁺-tdTom⁺-INs, indicative of apoptosis. Scale= 50 μ m.
- D. Cartoon representing developmental sequence of events related to differentiation of MGE-derived INs into SST and PV populations in WT (left) and *Fmr1* KO mice (right). Note that in WT mice, SST-INs differentiate before PV-INs, that naturally occurring cell death of SST-INs and PV-INs occurs roughly between P5 and P10, and that additional differentiation of PV-INs from MGE-derived precursors goes on after P15. In contrast, in *Fmr1* KO mice, there is 1. A lower density of PV-INs (due to an excess cell death), and 2. a higher density of SST-INs (presumably due to less cell death and/or greater differentiation from MGE-INs) compared to WT mice.

Chronic DREADD manipulation (P5 to P9)
in *Nkx2.1-Cre^{+/+}*; WT or *Fmr1* KO

Percentage of mCherry⁺ cells among PVALB⁺-IN

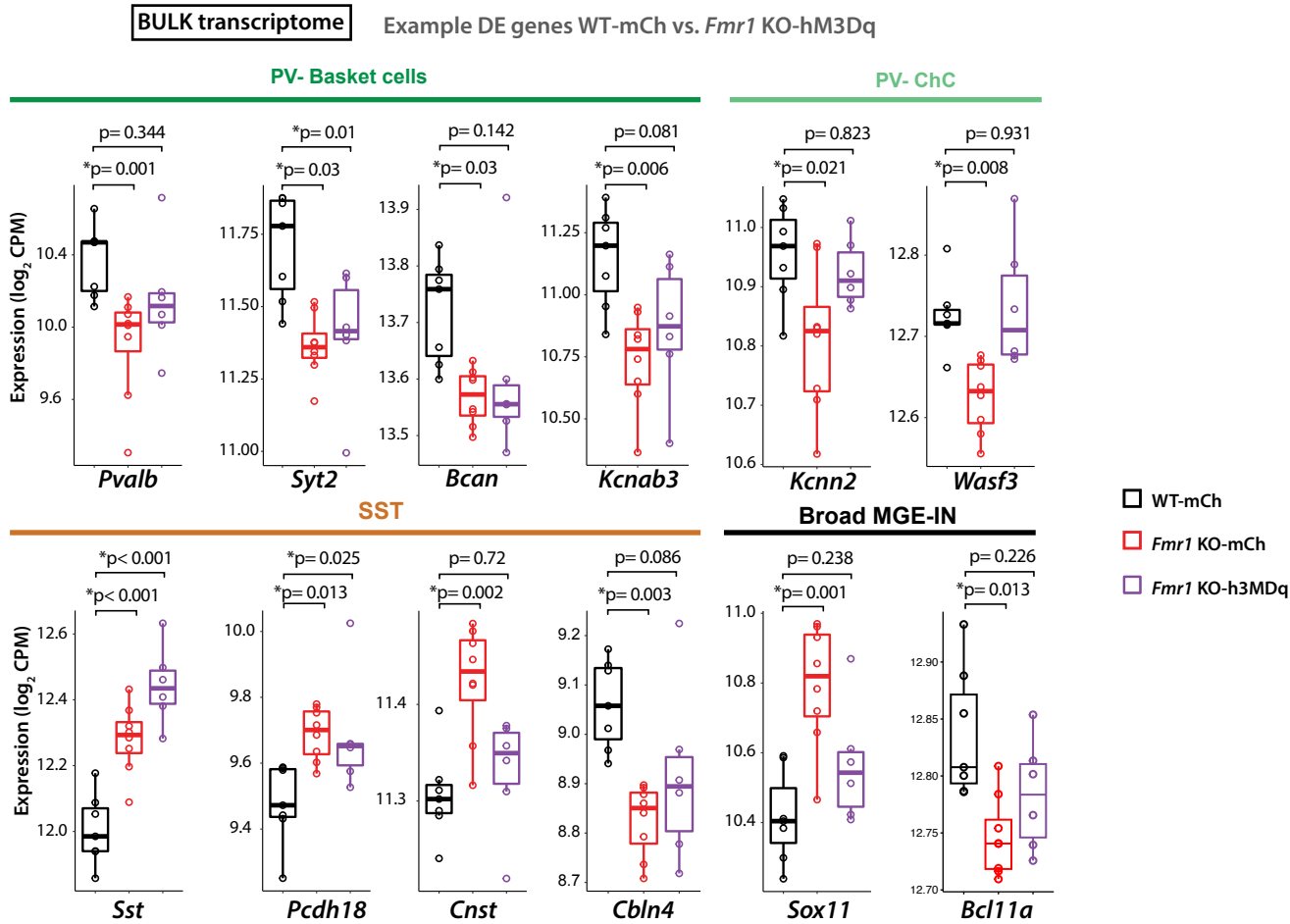


Supplementary Fig. S8

Supplementary Fig. S8: Chronic chemogenetic activation of Nkx2.1-INs (P5-P9) increases the density of PVALB⁺ cells at P21. (Related to Fig. 4)

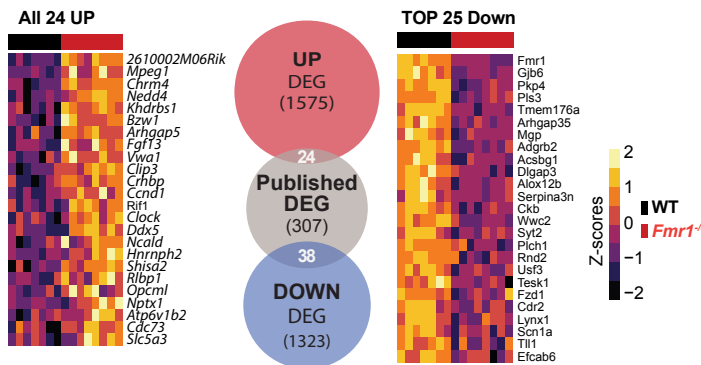
- A. Coronal section through the barrel field of S1 in a P15 Nkx2.1-Cre;WT mouse expressing mCherry in Nkx2.1⁺-IN (see Methods). Note the overlap in expression of mCherry (purple) in Nkx2.1⁺-IN and PVALB immunoreactivity (green) across cortical layers (white arrowheads). As expected based on the low efficiency of viral transduction, only a small proportion of PVALB⁺ cells co-express mCherry (and the Gq DREADD construct) (blue arrowheads). Scale= 50 μ m.
- B. Percentage of PVALB⁺ INs that also express mCherry. (25.8 ± 3.5 cells/mm², n=5 mice).
- C. Left: Quantification of PVALB⁺ INs density across cortical layers at P15. (161 ± 6 cells/mm² for WT-mCherry vs. 87 ± 6 for *Fmr1* KO-mCherry, p= 0.003; and 119 ± 9 for *Fmr1* KO-h3MDq, p= 0.040, n=6, 7 and 8, respectively). Right: the chronic DREADD manipulation did not significantly change the density of PV-INs in L4 or L5/6.

A

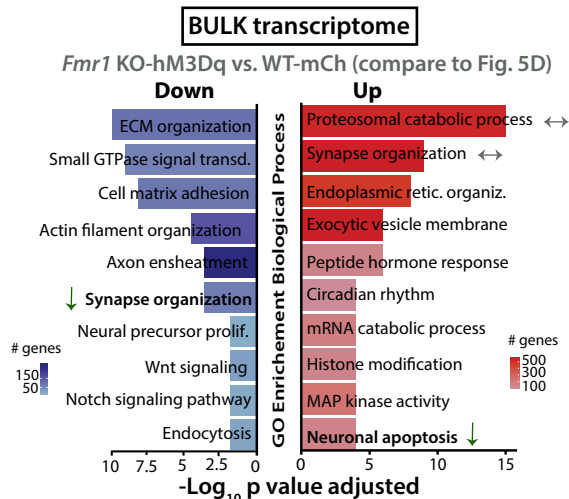


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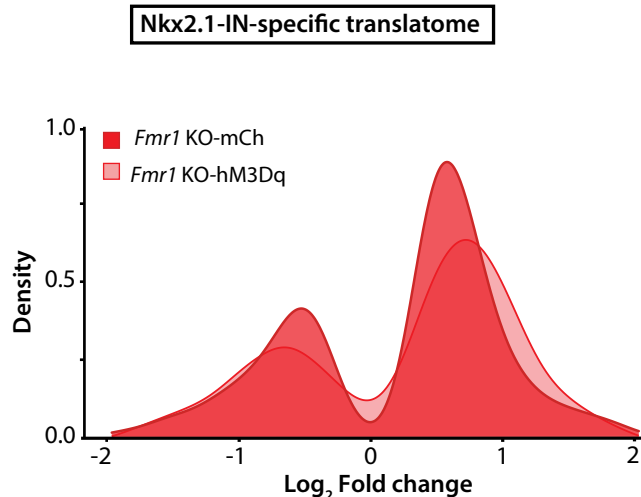
BULK transcriptome
Overlap of global cortical transcriptome with other RNAseq studies



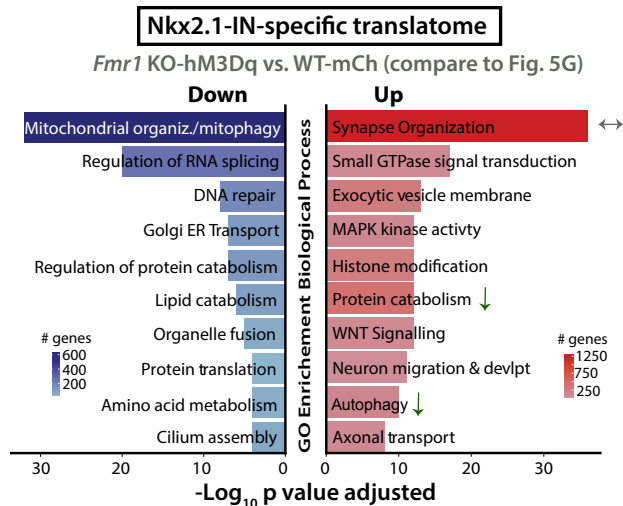
C



D



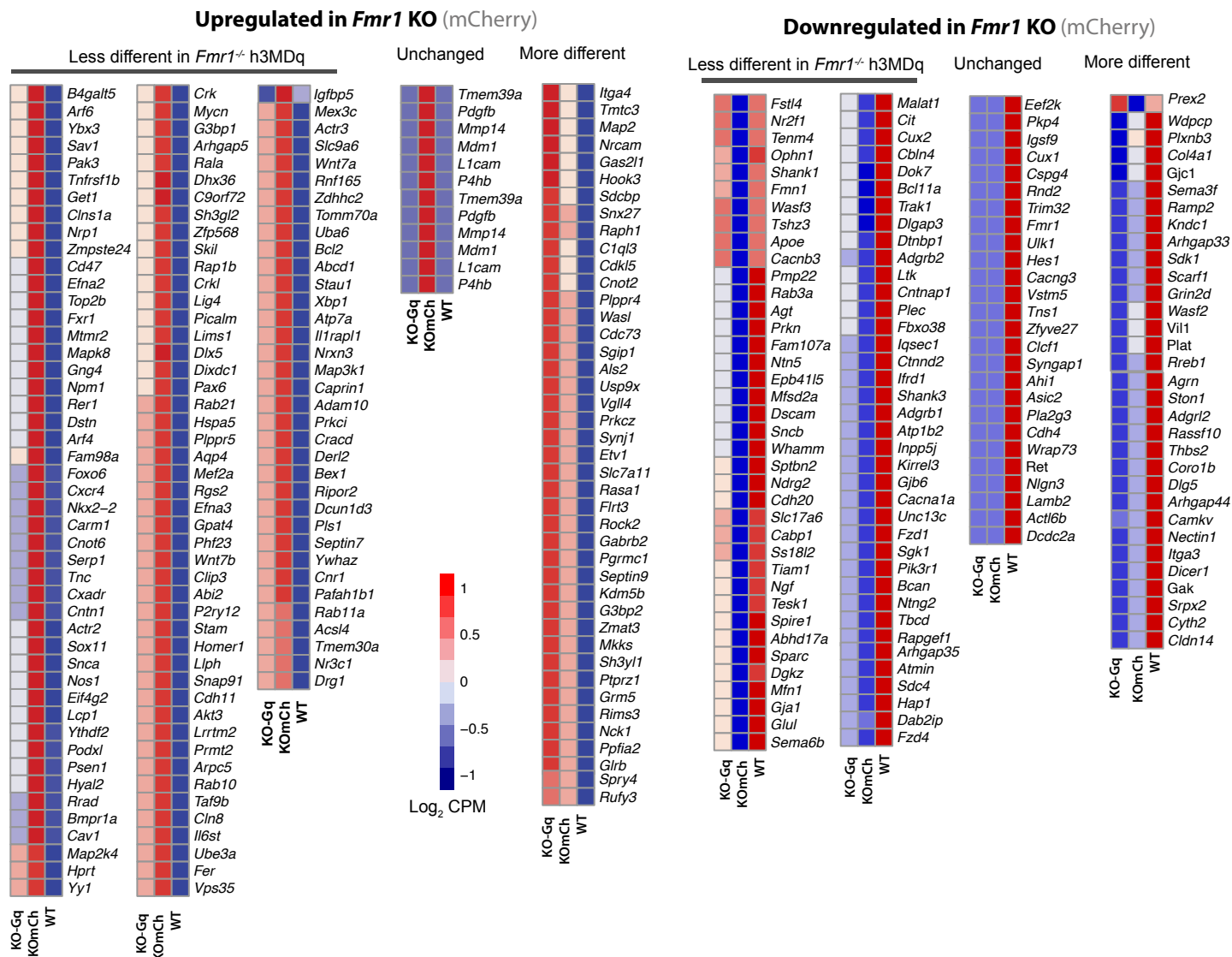
E



Supplementary Fig. S9: Gene ontology (GO) analysis for DE genes in *Fmr1* KO-hM3Dq mice compared to WT-mCherry mice, and effect of Gq DREADD manipulation. (Related to Fig. 5)

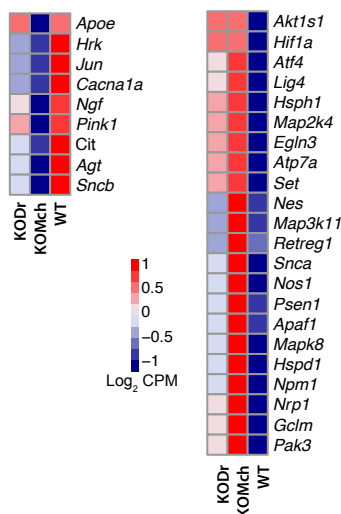
- A. Whisker plots comparing the expression of several MGE-derived markers in the bulk transcriptome. Markers shown for PV-INs, including fast-spiking basket cells and chandelier cells, and SST-INs markers; Wilcoxon signed-rank test.
- B. Number of DE genes in S1 cortex at P15 *Fmr1* KO-mCherry from the bulk transcriptome that overlap with previously reported differentially expressed in the hippocampus of adult *Fmr1* KO mice (see Methods). Heatmaps represent the changes of expression expressed as Z-scores among the up- and downregulated genes.
- C. Top 10 GO terms (using the biological process package) enriched among downregulated (blue) and upregulated (red) genes from the bulk cortical transcriptome in *Fmr1* -hM3Dq vs. WT-mCherry mice. Scale bars represent the number of genes in each category. Note that “*Synapse organization*” and “*Neuronal apoptosis*” categories are less different (reduced number of genes and adjusted p value) than in the comparison shown in Fig. 5D between *Fmr1* KO (mCherry) and WT (mCherry) mice, suggesting they were ‘improved’ by the chemogenetic activation of Nkx2.1-INs.
- D. Density plot showing how the log₂ fold change was affected by the DREADD manipulation in the Nkx2.1-IN specific transcriptome. Note that differences with WT mice were accentuated by the DREADD intervention in both down- and upregulated categories.
- E. Top 10 GO terms for DE genes from the Nkx2.1-specific transcriptome in *Fmr1* KO -hM3Dq vs. WT-mCherry mice. Only a few GO terms were modestly improved by DREADDs (e.g., *Protein catabolism* and *Autophagy*).

A "Synaptic organization" GO terms changed by DREADDs in *Fmr1* KO mice: Bulk transcriptome

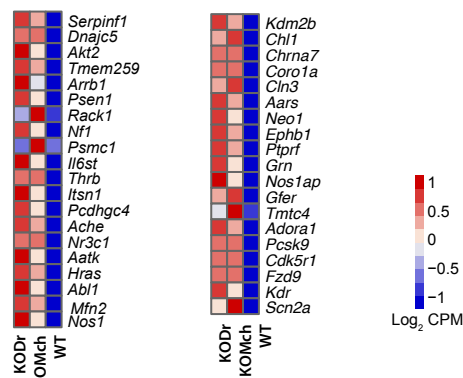


B "Neuron apoptosis" GO terms modified in *Fmr1* KO-h3MDq

BULK transcriptome

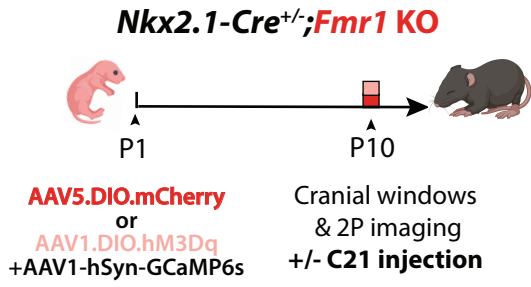
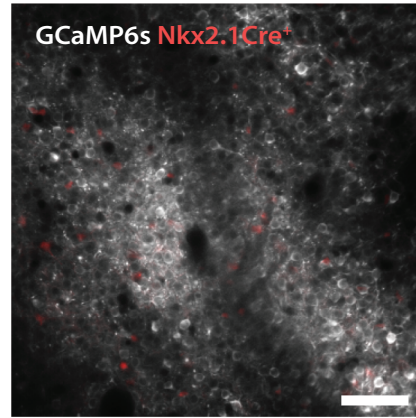
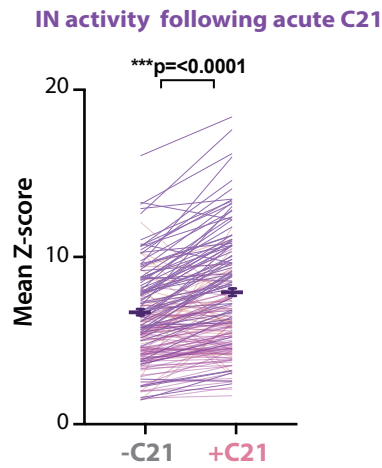
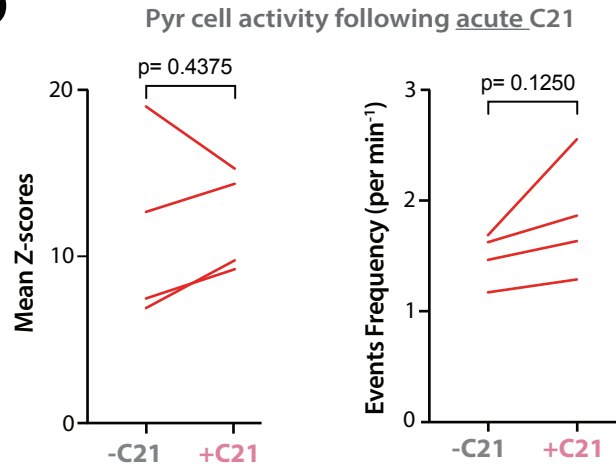
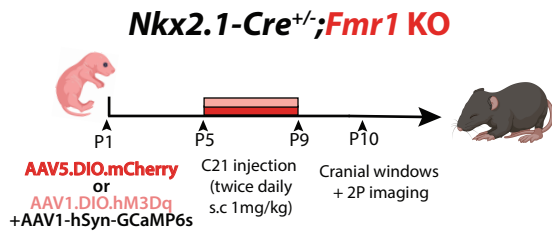
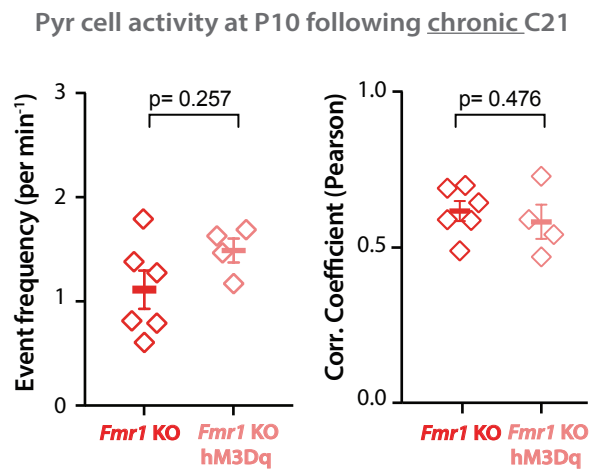


Nkx2.1-IN-specific transcriptome



Supplementary Fig. S10: Gene expression levels changed by C21 treatment in *Fmr1* KO-hM3Dq mice. (Related to Fig. 6)

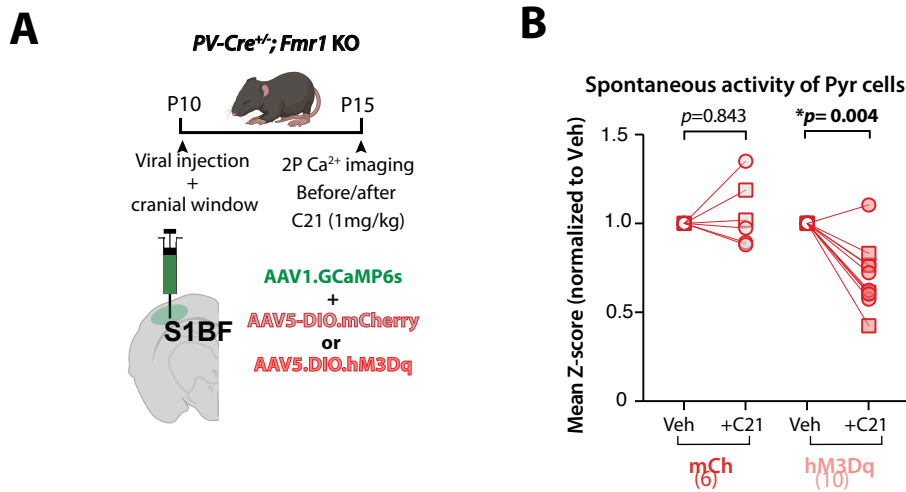
- A. List of upregulated or downregulated genes among the GO term *Synapse organization* in *Fmr1* KO-mCherry and hM3Dq groups (as compared to WT-mCherry) within the bulk cortical transcriptome. Heatmaps represent the average for each treatment/genotype group in (\log_2 CPM). Note that, while DREADD treatment reduced differences for many genes. The expression of other genes was either unchanged or worsened after C21 treatment.
- B. Same as in panel A but for GO term "*Neuronal apoptosis*." In both bulk transcriptome (left) and Nkx2.1-IN specific translome (right).

A**B****C****D****E****F**

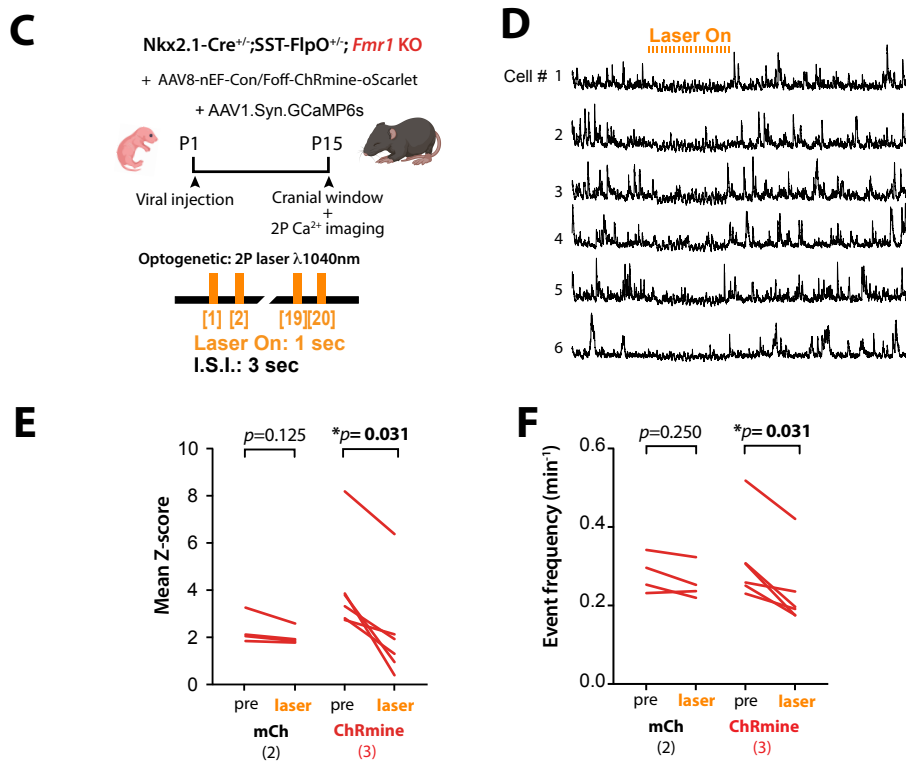
Supplementary Fig. S11: Acute chemogenetic activation of Nkx2.1-INs at P10 or chronic activation from P5 to P9, fails to modulate Pyr cell activity. (Related to Fig. 7)

- A. Experimental design for acute chemogenetic activation of Nkx2.1-INs at P10 in *Fmr1* KO mice to assess cortical circuit activity using in vivo calcium imaging at P10.
- B. Example FOV of Pyr cells expressing GCaMP6s and Nkx2.1-hM3Dq⁺ IN in S1 of *Nkx2.1-Cre; Fmr1* KO mouse (scale=100 μ m).
- C. Mean Z-scores of Nkx2.1-INs before (-C21) and 30-40min after s.c. injection of C21 (+C21) in P10 *Fmr1* KO-hM3Dq mice. Mean activity of Nkx2.1-INs is significantly higher following C21 injection (-C21: 6.61 ± 0.19 and +C21: 7.83 ± 0.23 , n= 182 cells from 3 mice; p<0.0001, Wilcoxon matched-pairs signed rank test).
- D. Left: Mean Z-scores of Pyr cells before (-C21) and 30-40min after (+C21) s.c. injection of C21 (1mg/kg) in P10 *Fmr1* KO-hM3Dq mice. Acutely increasing the activity of Nkx2.1-INs had no effect on Pyr cell activity (-C21: 11.52 ± 2.82 and +C21: 12.16 ± 1.55 , p=0.437, Wilcoxon matched-pairs signed rank test). Right: the mean frequency of synchronous network events (Pyr cells) remained unchanged after C21 in P10 *Fmr1* KO-hM3Dq mice. (-C21: 1.48 ± 0.11 and +C21: 1.83 ± 0.26 events/min, n= 4 mice; p=0.125, Wilcoxon matched-pairs signed rank test).
- E. Experimental design for chronic chemogenetic activation of Nkx2.1-INs (from P5 to P9) in *Fmr1* KO mice to assess cortical circuit activity using in vivo calcium imaging at P10.
- F. Left: Mean frequency of Pyr cell calcium transients at P10 was not significantly different *Nkx2.1-Cre; Fmr1* KO -hM3Dq mice after chronic C21 injections from P5 to P9 compared to *Fmr1* KO mice (1.11 ± 0.18 in *Fmr1* KO n=6 vs. 1.48 ± 0.11 events per min in *Fmr1* KO-hM3Dq mice n=4, p=0.257, MW test test). Right: Mean pair-wise correlation coefficients of Pyr cell calcium transients is unchanged in *Fmr1* KO-hM3Dq mice following chronic C21 injection from P5 to P9 (0.66 ± 0.03 in *Fmr1* KO n=6, and 0.62 ± 0.05 in *Fmr1* KO-hM3Dq mice n=4, p=0.476 MW test). Note that 1 animal was a new *Fmr1* KO-mCherry control, while the other 5 mice are *Fmr1* KO from Fig S4C-E.

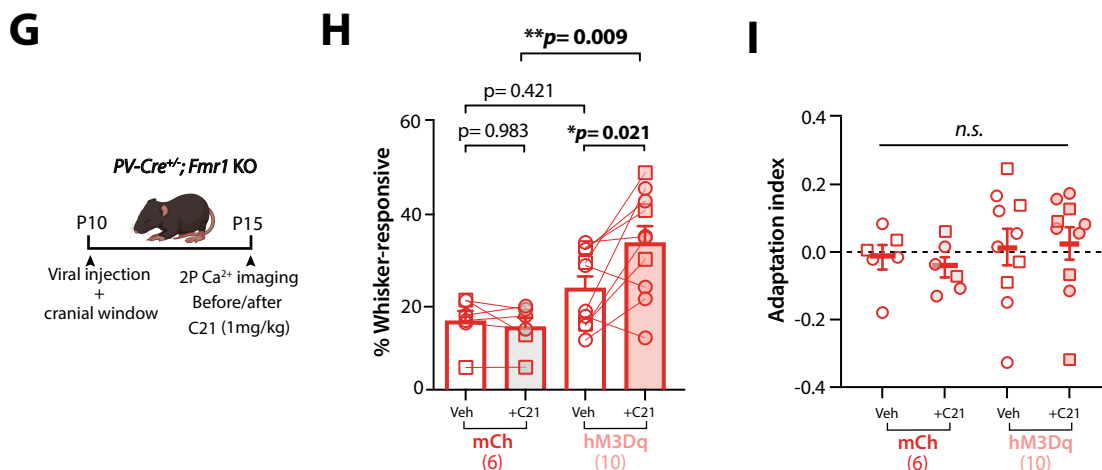
Acute DREADD manipulation of PV-IN reduces spontaneous Pyr cell activity in P15 Fmr1 KO mice



Acute optogenetic manipulation of MGE-IN reduces Pyr cell activity P15 Fmr1 KO mice



Acute DREADD manipulation of PV-IN modulates whisker-evoked responses



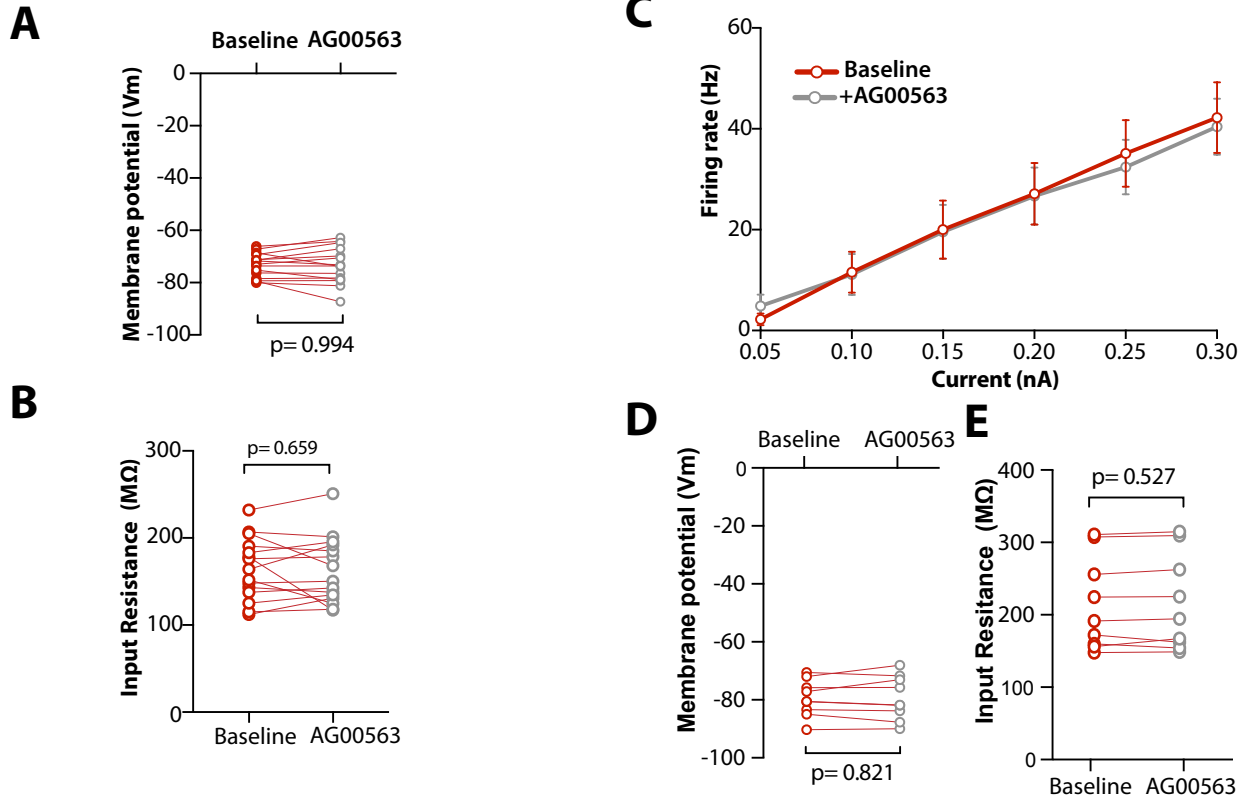
Supplementary Fig. S12

Supplementary Fig. S12: Acute chemogenetic or optogenetic activation of INs reduces Pyr cell activity at P15 in *Fmr1* KO mice. (Related to Figs. 7 and 8)

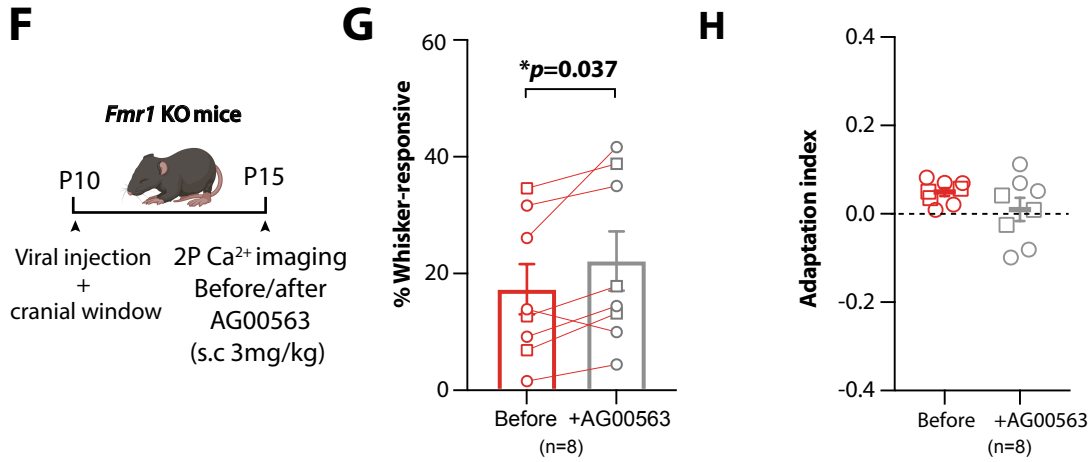
- A. Experimental design for in vivo calcium imaging recordings in Pyr cells after acute chemogenetic activation of PV-INs at P15. Calcium imaging was performed at P15 before and 30-40min after s.c injection of C21.
- B. Mean Z-score for spontaneous activity of L2/3 Pyr cells is reduced upon C21 injection in *Fmr1* KO -hM3Dq mice but not in *Fmr1* KO mCherry controls (before/after C21: 3.9 ± 0.7 vs. 4.0 ± 0.6 in mCherry group, $p=0.843$; and 2.3 ± 0.3 vs. 1.6 ± 0.3 , $p=0.004$ in hM3Dq group; $n=6$ and 10 mice, respectively; Wilcoxon matched pair signed rank test).
- C. Experimental design for optogenetic experiments. *Nkx2.1*-Cre mice (*Fmr1* KO or WT) were injected with a CreOn/FlpOff-ChRmine virus at P1 to express the opsin ChRmine in *Nkx2.1*-Cre⁺; Sst-FlpO-INs. Calcium imaging was done at P15 before, during, and after 20 laser pulses of orange light (1 s-long, 3 s I.S.I., $\lambda=1,040$ nm), just as in Fig. 2.
- D. Representative calcium traces for 6 Pyr cells (black) upon 2P laser stimulation.
- E. Mean Z-score of activity in Pyr cells before (pre) and during optogenetic stimulation (laser) in *Fmr1* KO mice. Each line in the panel represents an individual field of view. We observe a significant reduction of Pyr cell activity upon laser stimulation in ChRmine-expressing mice but not in control mCherry mice (2.32 ± 0.32 pre vs. 2.03 ± 0.18 with laser; $p=0.125$; $n=4$ FOV from 2 control mice); 4.12 ± 0.83 pre vs. 2.18 ± 0.87 with laser; $n=6$ FOV from 3 ChRmine-expressing mice; $p=0.019$. Wilcoxon matched-pairs signed rank test).
- F. Mean event frequency of calcium transients of Pyr cells before (pre) and during optogenetic stimulation (laser) in P15 *Fmr1* KO mice. We observe a significant reduction of Pyr cell activity upon laser stimulation in ChRmine-expressing mice but not in control m-Cherry mice (0.28 ± 0.02 pre vs. 0.25 ± 0.02 with laser; $p=0.250$ in control mice; 4.12 ± 0.83 pre vs. 2.18 ± 0.87 with laser for ChRmine-expressing mice; $p=0.031$, Wilcoxon matched-pairs signed rank test).
- G. Experimental design for calcium imaging recordings in Pyr cells after acute chemogenetic activation of PV-INs at P15 with the hM3Dq DREADD agonist C21 (same as Suppl. Fig. S12A-B).
- H. The percentage of whisker-responsive Pyr cells at P15 was significantly higher in upon C21 injection in *Fmr1* KO-hM3Dq mice ($n=10$) but not in *Fmr1* KO mCherry controls ($n=6$). (hM3Dq group before/after C21: $23.5 \pm 2.8\%$ vs. $33.9 \pm 3.9\%$; $p=0.021$; and mCherry group before/after C21: $15.9 \pm 2.3\%$ vs. $14.5 \pm 2.1\%$; $p=0.983$; two-way ANOVA with post-hoc Tukey).
- I. Neuronal adaptation was not affected by the DREADD manipulation (mCherry group before/after C21: -0.02 ± 0.09 vs. -0.05 ± 0.03 , hM3Dq group: 0.02 ± 0.05 vs. 0.03 ± 0.05 , $p>0.99$; two-way ANOVA with post-hoc Tukey).

Intrinsic properties of PV-Cre⁺;tdTom⁺ IN in P15 *Fmr1* KO mice

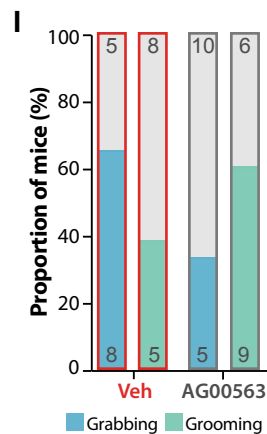
Firing rate and intrinsic properties of Pyramidal cells in P15 *Fmr1* KO mice



Acute AG00563 systemic injection modulates whisker-evoked responses



Proportion of mice grabbing or grooming during whisker stimulation at P21



Supplementary Fig. S13: Intrinsic properties of PV-INs and Pyr cells are unchanged by AG00563 (and relative proportions of grabbing/grooming in AG00563 treated mice).

(Related to Fig. 8)

- A. Resting membrane potential (V_m) of PV-INs is unchanged by bath application of AG00563 during current clamp recordings of *PV*-tdTom+ cells (-73.4 ± 1.2 mV vs. -73.2 ± 1.8 mV, $p = 0.805$, paired t -test, $n = 15$ cells from 6 *Fmr1* KO mice at P15-16).
- B. Input resistance (R_m) of PV-INs is unchanged by AG00563 (164.6 ± 9.2 M Ω vs. 161.2 ± 10.1 M Ω , $p = 0.608$, paired t -test).
- C. Cumulative input-output curves during baseline (red) or bath application of AG00563 (gray) ($n = 9$ Pyr cells from 6 *PV-Cre;tdTom+/-;Fmr1* KO mice, two-way RM ANOVA).
- D. V_m of Pyr cells is unchanged by AG00563 (-79.5 ± 2.1 mV vs -79.3 ± 2.5 mV, $p = 0.805$, paired t -test).
- E. R_m of Pyr cells is unchanged by AG00563 (214.0 ± 21.4 M Ω vs. 215.4 ± 22.0 M Ω , $p = 0.608$, paired t -test).
- F. Experimental design for the acute administration of AG00563 (3 mg/kg, s.c.) and calcium imaging at P15, before and 30 min after injection.
- G. The percentage of whisker-responsive Pyr cells in *Fmr1* KO mice was significantly higher after AG00563 injection compared to baseline ($17.1 \pm 4.3\%$ baseline vs. $21.9 \pm 5.1\%$ ~30-40 min after AG00563, $p = 0.033$; paired t -test, $n = 8$ mice).
- H. The neuronal adaptation index of Pyr cells was not changed by AG00563 (0.05 ± 0.01 baseline vs. 0.01 ± 0.03 after AG00563, $p = 0.033$; paired t -test, $n = 8$ mice).
- I. A smaller percentage of mice showed defensive behavior (grabbing) at least once during whisker stimulation in the AG00563-treated group than among vehicle controls (5/15 mice vs. 8/13, respectively). The opposite was true for adaptive healthy behavior (grooming) (9/15 mice vs. 5/13, respectively).

Case ID	Sex	Age	PMI (Hours)	Diagnosis	CGG Repeat Count	Hemisphere	Cause of Death
UCD 14-15	M	60	80	Control	NA	Right	Pulmonary Emboli
UCD 18-05	M	62	37	Control	NA	Left	Cardiopulmonary Arrest
UCD 15-07	F	64	NK	Control	NA	Left	NK
UCD 18-07	M	65	240	Control	NA	Left	Cardiac Arrest
UCD 14-01	M	66	48.5	Control	NA	Right	Acute Renal Failure
UCD 18-08	M	68	168	Control	NA	Left	Hypoxic Respiratory Failure
UCD 14-12	M	68	NK	Control	NA	Left	Cardiac Arrest
UCD 19-12	M	81	72	Control	NA	Left	NK
1031-08-GP	M	57	20	FXS	436	Left	Multiple System Organ Failure
1031-09-LZ	M	64	11.5	FXS	429	Left	NK
1061-19-JB	F	64	30	FXS	629,780	Left	NK
1005-14-JC	M	65	60	FXS	600-700	Right	Congestive Heart Failure
1013-10-SK	M	76	NK	FXS & FXTAS	447,540	Left	Respiratory Failure
1001-18-LD	M	78	6	FXS	235	Right	NK
1033-08-WS	M	79	17.5	FXS		Left	NK
1007-18-RF	M	80	NK	FXS	1,000	Right	NK

Table S1. Clinical characteristics of postmortem neurotypical and Fragile X cases.

(Related to Figure 3).

(NK: not known; NA: not applicable; M: male; F: female; PMI: post-mortem interval; FXS: Fragile X syndrome; FXTAS: Fragile X-associated tremor/ataxia syndrome).

SUPPLEMENTAL VIDEOS

Supplementary video 1 (Related to Fig. 2). Example of in vivo calcium imaging of Pyr cells before, during, and after optogenetic stimulation of presumed future PV-IN s in Nkx2.1-Cre;SST-Flp^{+/−} mice at P10. Data was acquired at 15 fps and is played back at 2x speed.

Supplementary video 2 (Related to Fig. 4). Example of in vivo calcium imaging of Pyr cells during whisker-evoked activity in Nkx2.1-Cre mice at P2. Data was acquired at 15 fps and is played back at 1x speed.

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2. Nigro, M.J., Hashikawa-Yamasaki, Y., and Rudy, B. (2018). Diversity and Connectivity of Layer 5 Somatostatin-Expressing Interneurons in the Mouse Barrel Cortex. *Journal of Neuroscience* 38, 1622–1633. 10.1523/jneurosci.2415-17.2017.
3. Daigle, T.L., Madisen, L., Hage, T.A., Valley, M.T., Knoblich, U., Larsen, R.S., Takeno, M.M., Huang, L., Gu, H., Larsen, R., et al. (2018). A Suite of Transgenic Driver and Reporter Mouse Lines with Enhanced Brain-Cell-Type Targeting and Functionality. *Elsevier Inc.* 174, 465-480.e22. 10.1016/j.cell.2018.06.035.