

## Supplementary figure legends

### **Figure S1. The glutamatergic neurons of PVT are activated during CPP-Test**

(A) Total distance traveled in CPP apparatus during CPP-Test of cohort 3 mice. Two-tailed unpaired t test.  $n = 20$  mice per group.  $t = 1.4370$ ,  $p = 0.1589$ .

### **Figure S2. Chemogenetic inhibition of PVT glutamatergic neurons during CPP-Training does not block the acquisition of METH CPP**

(A) Experimental design and timeline of cohort S1 mice.

(B) Schematic diagram of viral transfection in WT mice and representative image of *rAAV2/9-CaMKII $\alpha$ -hM4D(Gi)-mCherry* injection sites in the PVT of cohort S1 mice. Scale bar, 2 mm / 200  $\mu$ m.

(C) The percentage of virus transfected efficiency in CaMKII-positive neurons of PVT in cohort S1 mice. Two-tailed unpaired t test.  $n = 4$  mice per group.  $t = 0.3114$ ,  $p = 0.7661$ . Scale bar, 200  $\mu$ m / 50  $\mu$ m.

(D) The percentage of c-Fos-positive neurons on *mCherry*-transfected neurons in the PVT of cohort S1 mice. Two-tailed unpaired t test.  $n = 4$  mice per group.  $t = 3.4940$ ,  $p = 0.0129$ . Scale bar, 200  $\mu$ m / 50  $\mu$ m.

(E) Heatmap of spent duration by mice in CPP apparatus and CPP analysis of cohort S1 mice. Left panel, Representative heatmap. Middle panel, CPP score during the Pre-Test and CPP-Test. Two-way ANOVA with Sidak's multiple comparisons test.  $n = 10$  mice per group. CPP-Test of METH-Veh group,  $t = 6.3140$ ,  $p < 0.0001$  vs corresponding Pre-Test. CPP-Test of METH-CNO group,  $t = 3.6610$ ,  $p = 0.0036$  vs corresponding Pre-Test. Right panel,  $\Delta$ CPP score (CPP-Test score minus Pre-Test score). Two-tailed unpaired t test.  $n = 10$  mice per group.  $t = 1.8760$ ,  $p = 0.0770$ .

(F) Total distance traveled in CPP apparatus during CPP-Test of cohort S1 mice. Two-tailed unpaired t test.  $n = 10$  mice per group.  $t = 0.0381$ ,  $p = 0.9700$ .

### **Figure S3. Chemogenetic inhibition of PVT glutamatergic neurons during CPP-Test disrupts the acquisition of METH CPP**

(A) The percentage of virus transfected efficiency in CaMKII-positive neurons of PVT in cohort 4 mice. Two-way ANOVA with Sidak's multiple comparisons test.  $n = 4$  mice per group.  $F_{(1, 12)} = 0.0131$ ,  $p = 0.9108$ . Saline-CNO group,  $t = 0.0961$ ,  $p > 0.9999$  vs Saline-Veh group. METH-CNO group,  $t = 0.2580$ ,  $p > 0.9999$  vs METH-Veh group. Scale bar, 200  $\mu\text{m}$  / 50  $\mu\text{m}$ .

(B) The percentage of c-Fos-positive neurons on *mCherry*-transfected neurons in the PVT of cohort 4 mice. Two-way ANOVA with Sidak's multiple comparisons test.  $n = 4$  mice per group.  $F_{(1, 12)} = 5.8370$ ,  $p = 0.0326$ . Saline-CNO group,  $t = 0.0091$ ,  $p > 0.9999$  vs Saline-Veh group. METH-CNO group,  $t = 3.4070$ ,  $p = 0.0308$  vs METH-Veh group. METH-Veh group,  $t = 3.2950$ ,  $p = 0.0378$  vs Saline-Veh group. Scale bar, 200  $\mu\text{m}$  / 50  $\mu\text{m}$ .

(C) Total distance traveled in CPP apparatus during CPP-Test 1 and CPP-Test 2 of cohort 4 mice. Two-way ANOVA with Sidak's multiple comparisons test.  $n = 10$  mice per group. CPP-Test 1,  $F_{(1, 36)} = 0.0078$ ,  $p = 0.9297$ . Saline-CNO group,  $t = 0.6667$ ,  $p = 0.9860$  vs Saline-Veh group. METH-CNO group,  $t = 0.7923$ ,  $p = 0.9669$  vs METH-Veh group. METH-Veh group,  $t = 1.0310$ ,  $p = 0.8915$  vs Saline-Veh group. CPP-Test 2,  $F_{(1, 36)} = 3.0030$ ,  $p = 0.0917$ . Saline-CNO group,  $t = 0.6742$ ,  $p = 0.9852$  vs Saline-Veh group. METH-CNO group,  $t = 1.7770$ ,  $p = 0.4096$  vs METH-Veh group. METH-Veh group,  $t = 2.5240$ ,  $p = 0.0930$  vs Saline-Veh group.

**Figure S4. The paired pulse ratio was increased within PVT neurons during the acquisition of METH CPP**

(A) Experimental design and timeline of cohort S2 mice.

(B) Schematic diagram and representative image of whole-cell voltage-clamp recording site in the PVT of cohort S2 mice. Scale bar, 200  $\mu\text{m}$  / 50  $\mu\text{m}$ .

(C) Paired pulse ratio (PPR) recordings. Representative traces (left) and quantification of PPR in PVT neurons (right). Two-tailed unpaired t test.  $n = 13$  cells from 3 mice per group.  $t = 2.2150$ ,  $p = 0.0365$ .

**Figure S5. The spike frequency of AP decreased in mPFC *eNpHR3.0-mCherry*-transfected neurons by turning-on the yellow lights**

(A) Schematic diagram of viral transfection in WT mice and representative image of whole-cell current-clamp recording site in the mPFC of cohort 9 mice. Scale bar, 200  $\mu\text{m}$  / 50  $\mu\text{m}$ .

(B) The spike frequency of AP of mPFC *eNpHR3.0-mCherry*-transfected neurons under whole-cell current-clamp configuration and sample traces for the spike frequency of AP following optogenetic stimulation in naive mice (cohort 9 mice). Left, quantification of AP spike frequency. Middle, sample trace. Right, percentage of AP changes in the mPFC *eNpHR3.0-mCherry*-transfected neurons (25 out of 25 cells from 4 mice, 100%). One-way ANOVA with Sidak's multiple comparisons test.  $n = 25$  cells from 4 mice per group.  $F_{(1.206, 28.94)} = 80.0100$ ,  $p < 0.0001$ . ON,  $t = 9.8570$ ,  $p < 0.0001$  vs OFF 1;  $t = 8.4170$ ,  $p < 0.0001$  vs OFF 2.

**Figure S6. The mPFC glutamatergic neurons are activated during CPP-Test**

(A) Immunofluorescence for c-Fos/CaMKII in the mPFC layer II/III and layer V and VI following METH-induced CPP-Test of cohort 3 mice. Two-tailed unpaired t test.  $n = 6$  mice per group. Layer II/III,  $t = 2.4740$ ,  $p = 0.0329$ . Layer V and VI,  $t = 5.1660$ ,  $p = 0.0004$ . Scale bar, 200  $\mu\text{m}$  / 50  $\mu\text{m}$ .

(B) Total distance traveled in CPP apparatus during CPP-Test of cohort 10 mice. Two-tailed unpaired t test.  $n = 6$  mice per group.  $t = 0.6627$ ,  $p = 0.5225$ .

**Figure S7. Suppressing mPFC-PVT pathway disrupts the acquired METH CPP**

(A) The percentage of virus transfected efficiency in CaMKII-positive neurons of mPFC in cohort 11 mice. Two-tailed unpaired t test.  $n = 4$  mice per group.  $t = 0.0265$ ,  $p = 0.9797$ . Scale bar, 200  $\mu\text{m}$  / 50  $\mu\text{m}$ .

(B) The percentage of c-Fos-positive neurons on *mCherry*-transfected neurons following optogenetic stimulation in the PVT of cohort 11 mice. Two-tailed unpaired t test.  $n = 4$  mice per group.  $t = 5.7540$ ,  $p = 0.0012$ . Scale bar, 200  $\mu\text{m}$  / 50  $\mu\text{m}$ .

(C) Total distance traveled in CPP apparatus during CPP-Test 1 and CPP-Test

2 of cohort 11 mice. CPP-Test 1, Two-tailed unpaired t test.  $n = 10$  mice per group.  $t = 0.7133$ ,  $p = 0.4848$ . CPP-Test 2, Two-tailed unpaired t test.  $n = 10$  mice per group.  $t = 1.9790$ ,  $p = 0.0633$ .

**Figure S8. Suppression of PVT glutamatergic neurons caused a reduction in wakefulness**

(A) Schematic diagram of viral transfection in WT mice and representative image of *rAAV2/9-CaMKII $\alpha$ -hM4D(Gi)-mCherry* injection sites in the PVT of cohort S3 mice. Scale bar, 2 mm / 200  $\mu$ m.

(B) Percentage of time spent and number in each state within 3 h after injection of saline or CNO of cohort S3 mice. Two-way ANOVA with Sidak's multiple comparisons test.  $n = 6$  mice per group. Time spent in each state within 3 h,  $F_{(2, 20)} = 43.1300$ ,  $p < 0.0001$ . Wake,  $t = 6.8580$ ,  $p = 0.0008$ ; NREM,  $t = 6.3610$ ,  $p = 0.0013$ ; REM,  $t = 2.0800$ ,  $p = 0.2154$ . Number in each state within 3 h,  $F_{(2, 20)} = 28.8800$ ,  $p < 0.0001$ . Wake,  $t = 3.3330$ ,  $p = 0.0489$ ; NREM,  $t = 3.3650$ ,  $p = 0.0307$ ; REM,  $t = 0.1832$ ,  $p = 0.9973$ .

(C) EEG/EMG recording representative hypnograms during 3 h after saline or CNO injection of cohort S3 mice.

(D) Heatmap of relative EEG power and example traces of EEG/EMG in different sleep states defined by Sirenia Sleep.

(E) Percentage of time spent and number in each state within 24 h, 12 h (dark phase) and 12 h (light phase) after injection of saline or CNO in cohort S3 mice. Two-way ANOVA with Sidak's multiple comparisons test.  $n = 6$  mice per group. Time spent in each state within 24 h,  $F_{(2, 20)} = 26.5500$ ,  $p < 0.0001$ . Wake,  $t = 5.1120$ ,  $p = 0.0015$ ; NREM,  $t = 5.2820$ ,  $p = 0.0011$ ; REM,  $t = 1.6340$ ,  $p = 0.3758$ . Number in each state within 24 h,  $F_{(2, 20)} = 31.7900$ ,  $p < 0.0001$ . Wake,  $t = 3.2690$ ,  $p = 0.0261$ ; NREM,  $t = 3.1800$ ,  $p = 0.0302$ ; REM,  $t = 0.1406$ ,  $p = 0.9987$ . Time spent in each state within 12 h (dark phase),  $F_{(2, 20)} = 31.6900$ ,  $p < 0.0001$ . Wake,  $t = 5.6120$ ,  $p = 0.0009$ ; NREM,  $t = 5.6990$ ,  $p = 0.0008$ ; REM,  $t = 1.2230$ ,  $p = 0.5956$ . Number in each state within 12 h (dark phase),  $F_{(2, 20)} = 28.1100$ ,  $p < 0.0001$ . Wake,  $t = 3.1860$ ,  $p = 0.0298$ ; NREM,  $t$

= 3.3300,  $p = 0.0230$ ; REM,  $t = 0.2447$ ,  $p = 0.9933$ . Time spent in each state within 12 h (light phase),  $F_{(2, 20)} = 0.1017$ ,  $p = 0.9038$ . Wake,  $t = 0.3544$ ,  $p = 0.9807$ ; NREM,  $t = 0.2485$ ,  $p = 0.9931$ ; REM,  $t = 1.7320$ ,  $p = 0.3410$ . Number in each state within 12 h (light phase),  $F_{(2, 20)} = 12.6600$ ,  $p = 0.0003$ . Wake,  $t = 2.0860$ ,  $p = 0.1801$ ; NREM,  $t = 2.1280$ ,  $p = 0.1686$ ; REM,  $t = 0.2187$ ,  $p = 0.9952$ .

**Figure S9. Suppressing mPFC-PVT pathway has no influence on wakefulness**

(A) Percentage of time spent and number in each state within 24 h, 12 h (dark phase) and 12 h (light phase) after injection of saline or CNO in cohort 11 mice. Two-way ANOVA with Sidak's multiple comparisons test.  $n = 6$  mice per group. Time spent in each state within 24 h,  $F_{(2, 20)} = 2.6210$ ,  $p = 0.0975$ . Wake,  $t = 1.5160$ ,  $p = 0.4568$ ; NREM,  $t = 1.7520$ ,  $p = 0.3580$ ; REM,  $t = 0.8095$ ,  $p = 0.8239$ . Number in each state within 24 h,  $F_{(2, 20)} = 0.0213$ ,  $p = 0.9789$ . Wake,  $t = 0.2565$ ,  $p = 0.9925$ ; NREM,  $t = 0.1834$ ,  $p = 0.9972$ ; REM,  $t = 0.5401$ ,  $p = 0.9366$ . Time spent in each state within 12 h (dark phase),  $F_{(2, 20)} = 0.5591$ ,  $p = 0.5804$ . Wake,  $t = 0.7563$ ,  $p = 0.8486$ ; NREM,  $t = 0.7479$ ,  $p = 0.8527$ ; REM,  $t = 0.3548$ ,  $p = 0.9805$ . Number in each state within 12 h (dark phase),  $F_{(2, 20)} = 0.4269$ ,  $p = 0.6583$ . Wake,  $t = 0.8301$ ,  $p = 0.7978$ ; NREM,  $t = 0.9122$ ,  $p = 0.7487$ ; REM,  $t = 0.0912$ ,  $p = 0.9996$ . Time spent in each state within 12 h (light phase),  $F_{(2, 20)} = 2.6330$ ,  $p = 0.0966$ . Wake,  $t = 1.5020$ ,  $p = 0.4368$ ; NREM,  $t = 1.7740$ ,  $p = 0.3027$ ; REM,  $t = 0.8538$ ,  $p = 0.8036$ . Number in each state within 12 h (light phase),  $F_{(2, 20)} = 0.1004$ ,  $p = 0.9049$ . Wake,  $t = 0.5258$ ,  $p = 0.9421$ ; NREM,  $t = 0.4109$ ,  $p = 0.9710$ ; REM,  $t = 0.7163$ ,  $p = 0.8695$ .

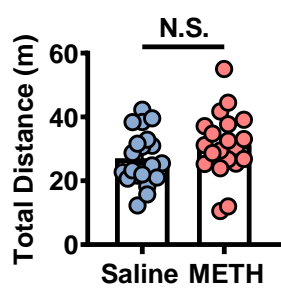
**Figure S10. Postsynaptic GluN2A is a key molecule along mPFC-PVT pathway that involves in the acquisition of METH CPP**

(A) Total distance traveled in CPP apparatus of GNE-5729 (Left) and S-AMPA (Right) mice (cohort 13 mice). GNE-5729, One-way ANOVA with Sidak's multiple comparisons test.  $n = 7$  mice.  $F_{(2.068, 12.41)} = 3.2800$ ,  $p = 0.0706$ . CPP-

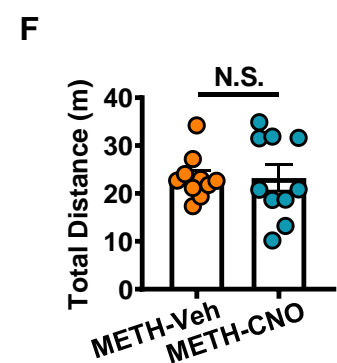
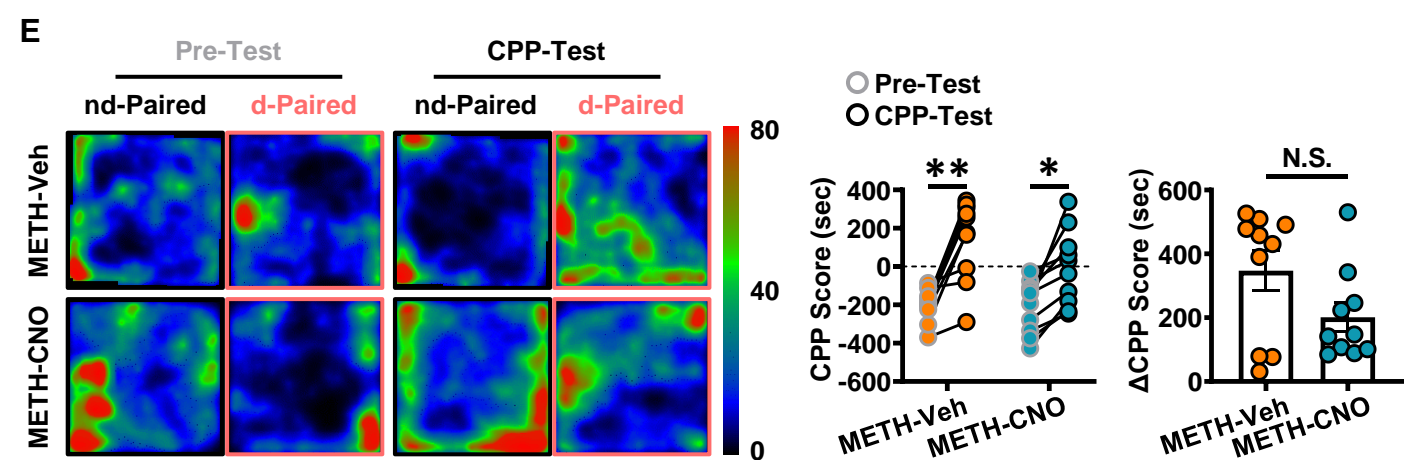
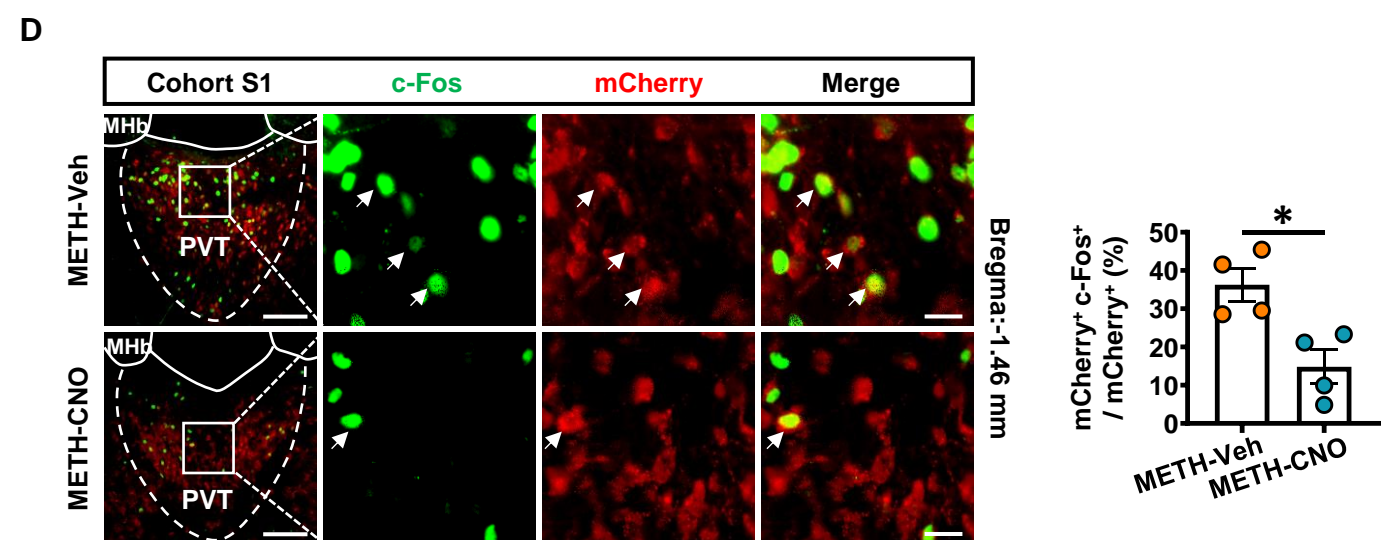
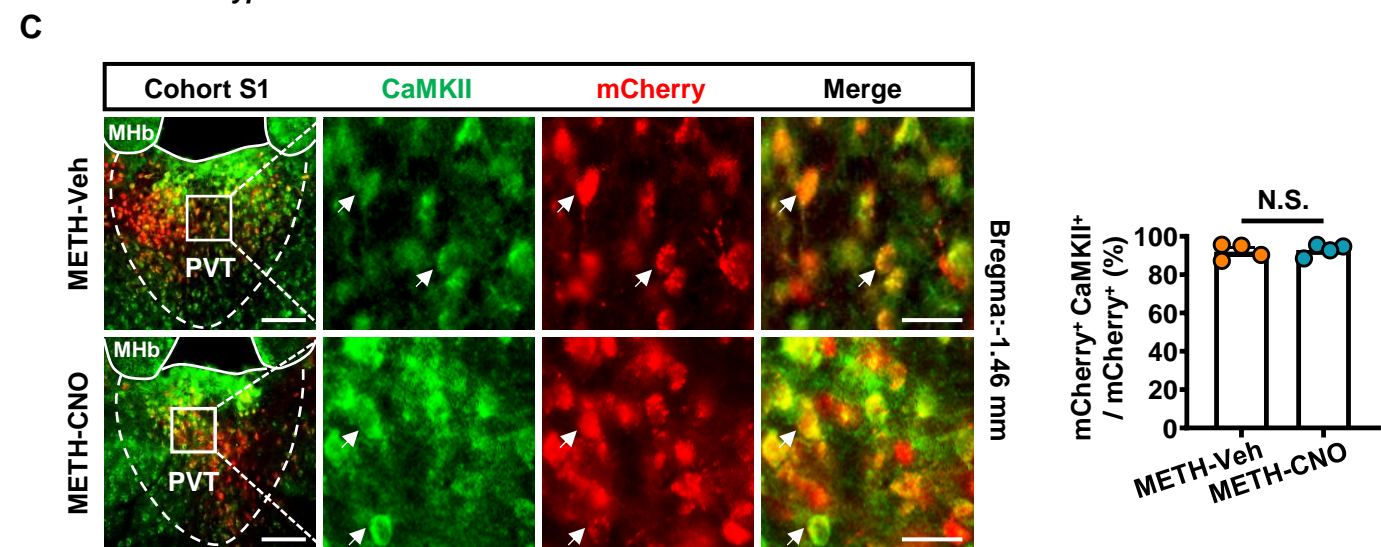
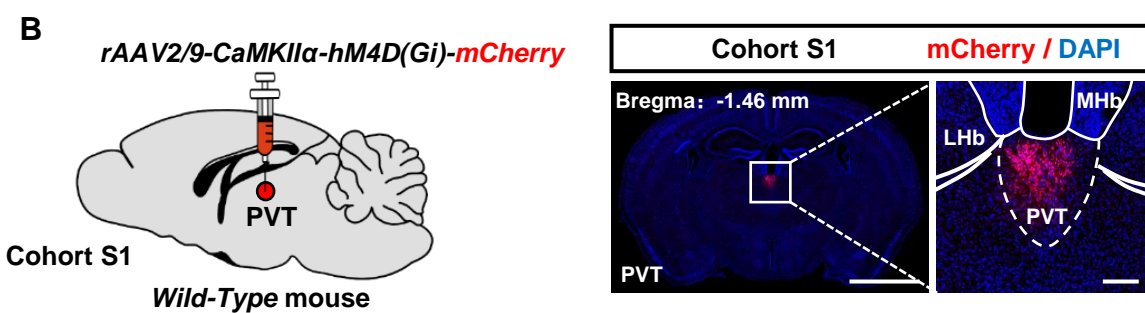
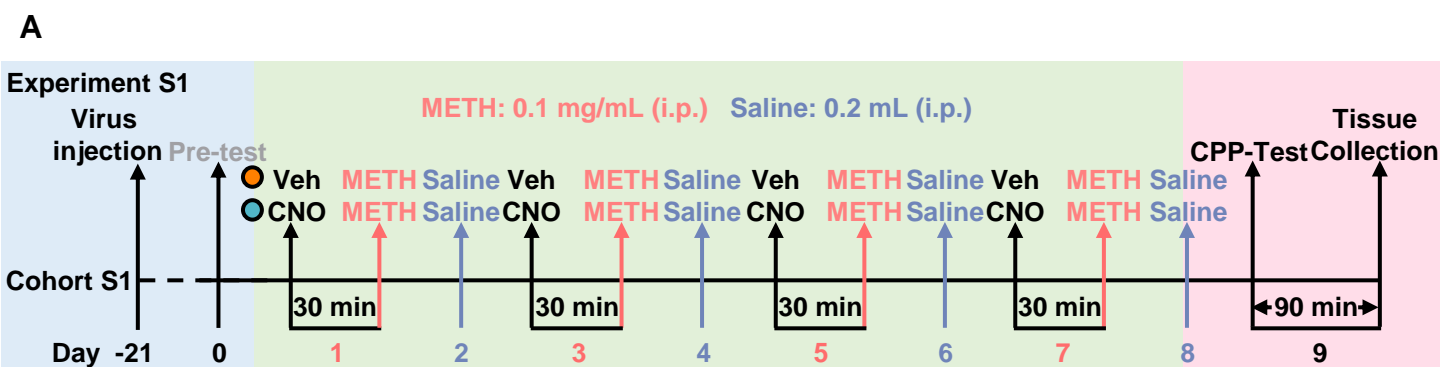
Test 1,  $t = 3.2880$ ,  $p = 0.0958$  vs Pre-Test. CPP-Test 2,  $t = 1.0290$ ,  $p = 0.9196$  vs Pre-Test. CPP-Test 3,  $t = 0.3908$ ,  $p = 0.9994$  vs Pre-Test. S-AMPA, One-way ANOVA with Sidak's multiple comparisons test.  $n = 7$  mice.  $F_{(2.028, 12.17)} = 3.6800$ ,  $p = 0.0558$ . CPP-Test 1,  $t = 0.1041$ ,  $p > 0.9999$  vs Pre-Test. CPP-Test 2,  $t = 3.1040$ ,  $p = 0.1196$  vs Pre-Test. CPP-Test 3,  $t = 1.9430$ ,  $p = 0.4688$  vs Pre-Test.

Figure S1

A



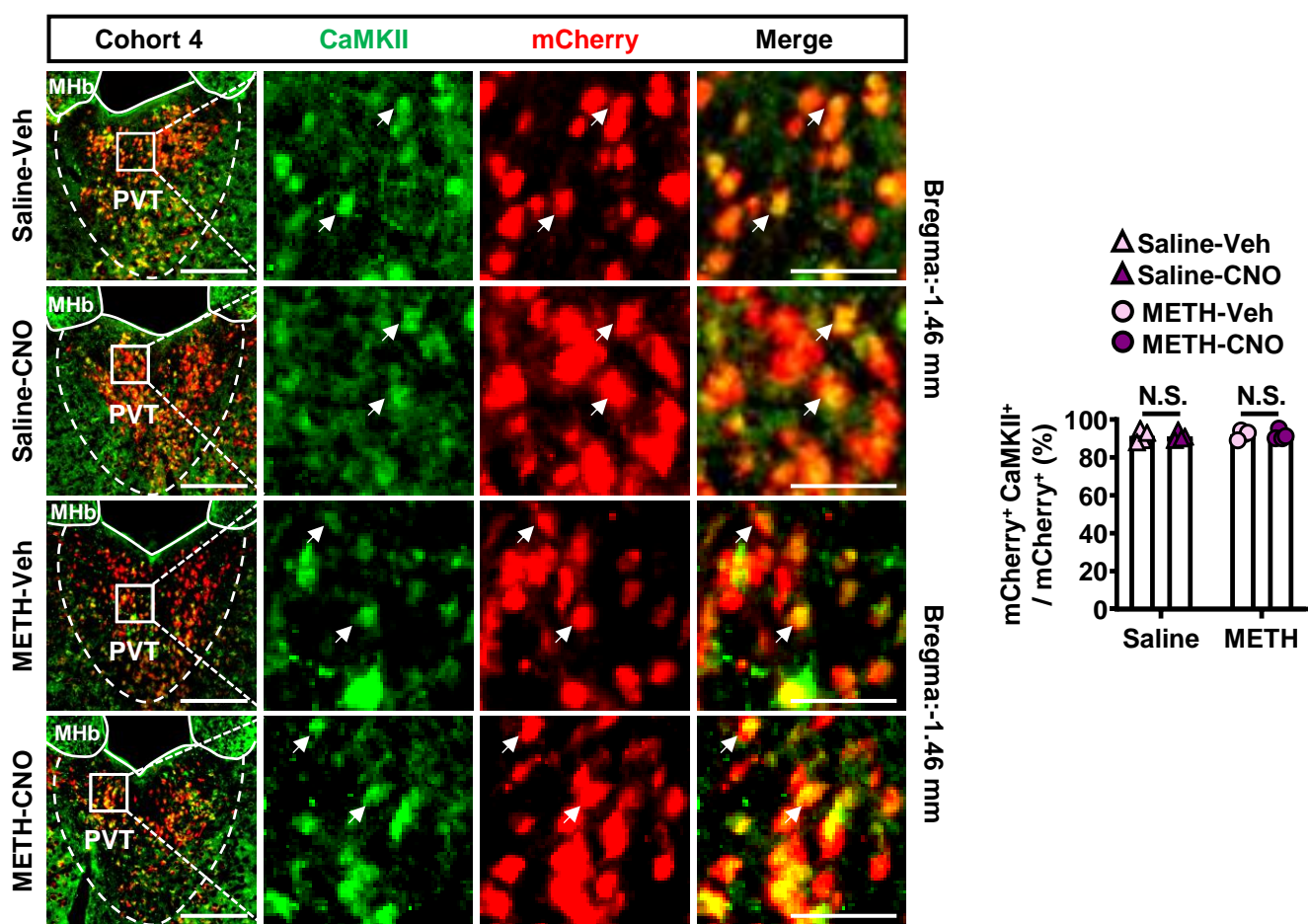
# Figure S2



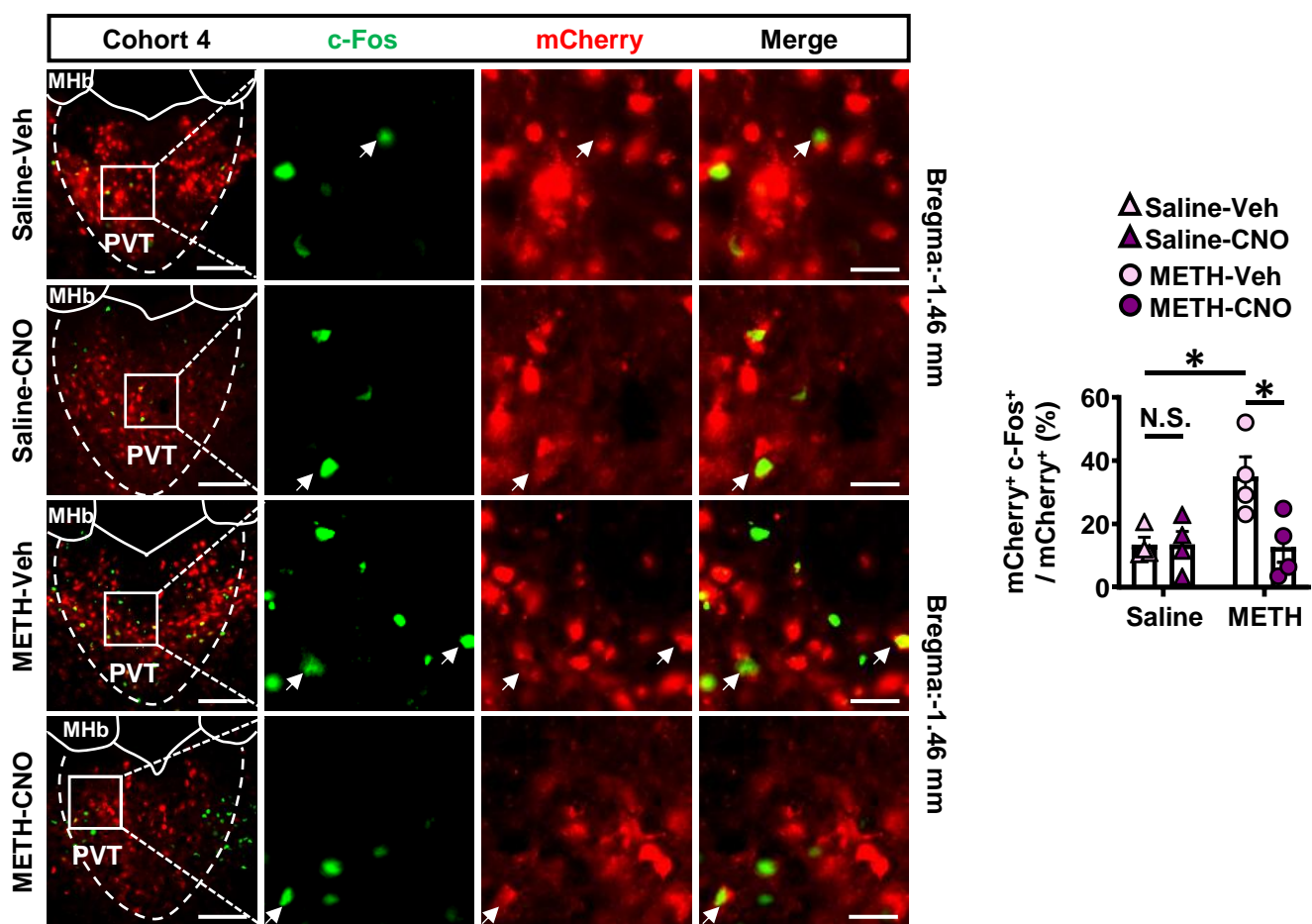


**Figure S3**

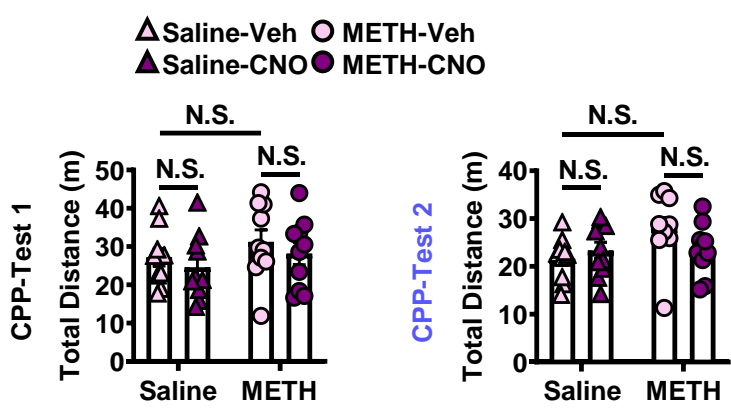
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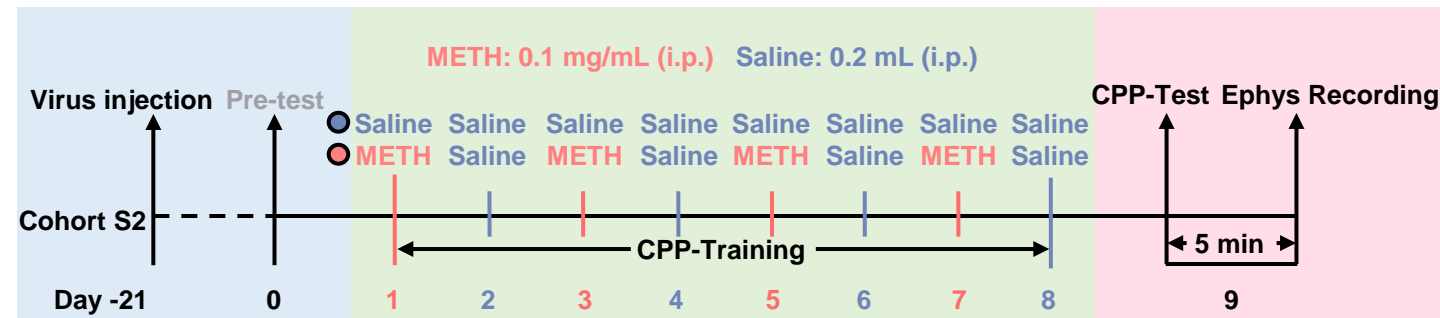
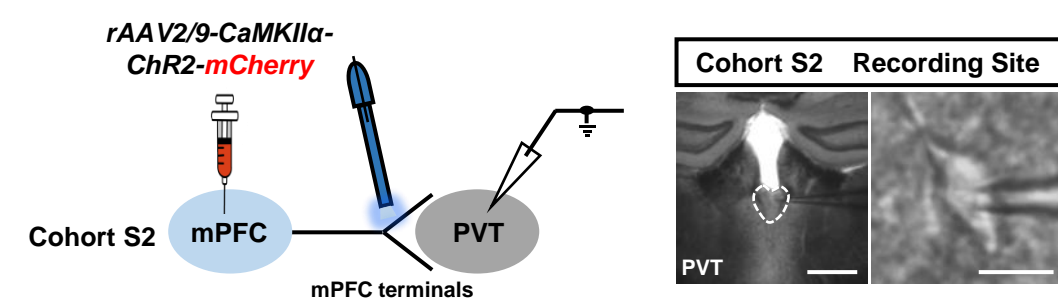
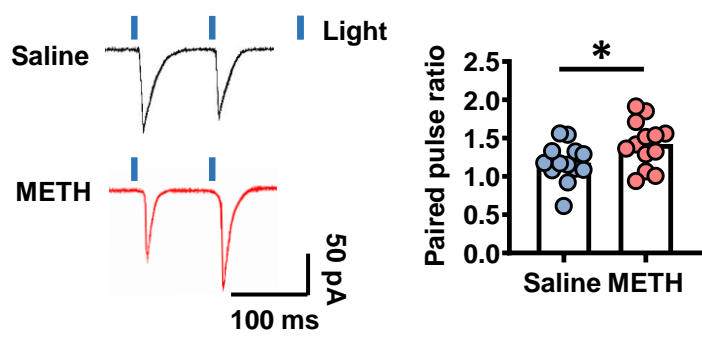


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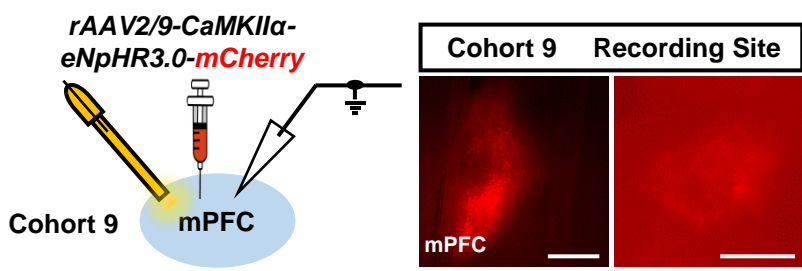
**C**



**Figure S4****A****B****C**

**Figure S5**

**A**



**B**

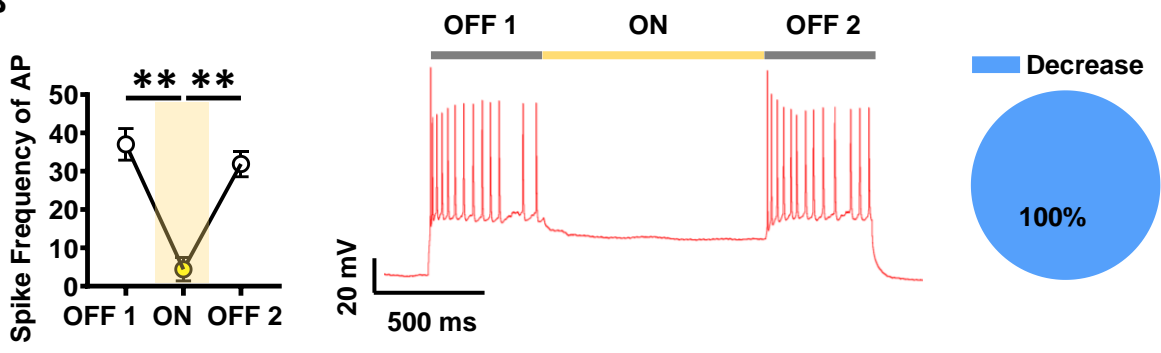
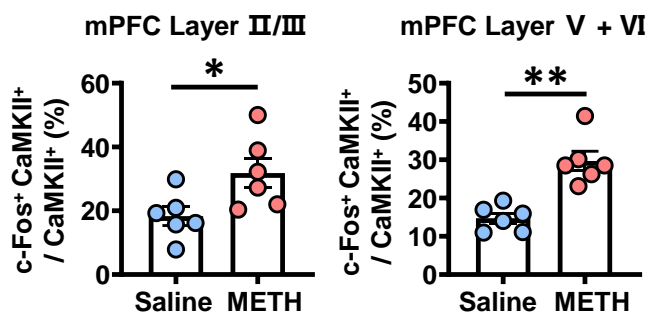
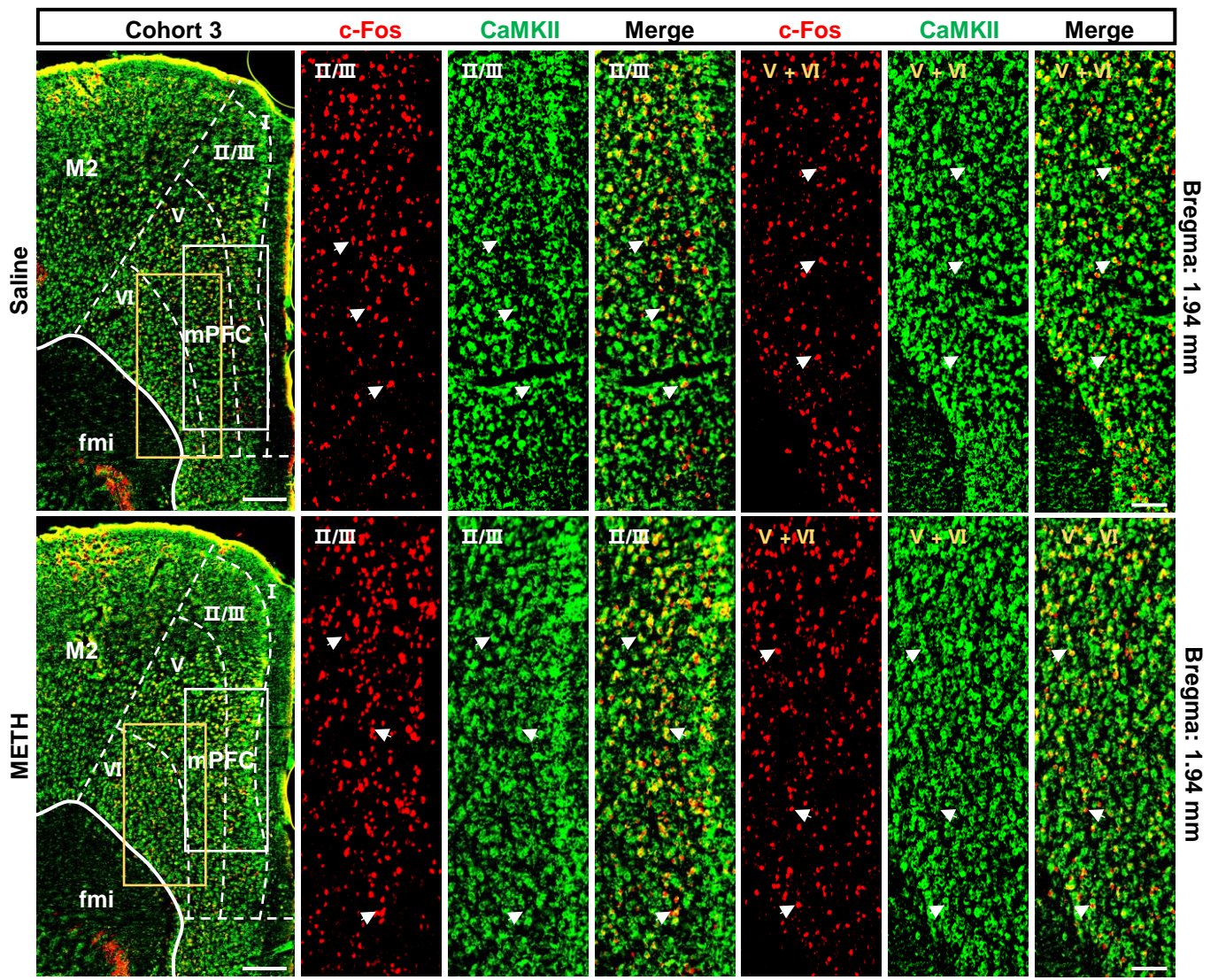
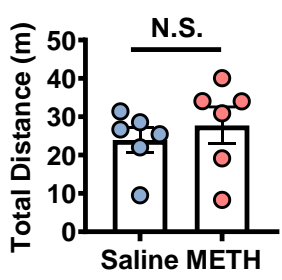


Figure S6

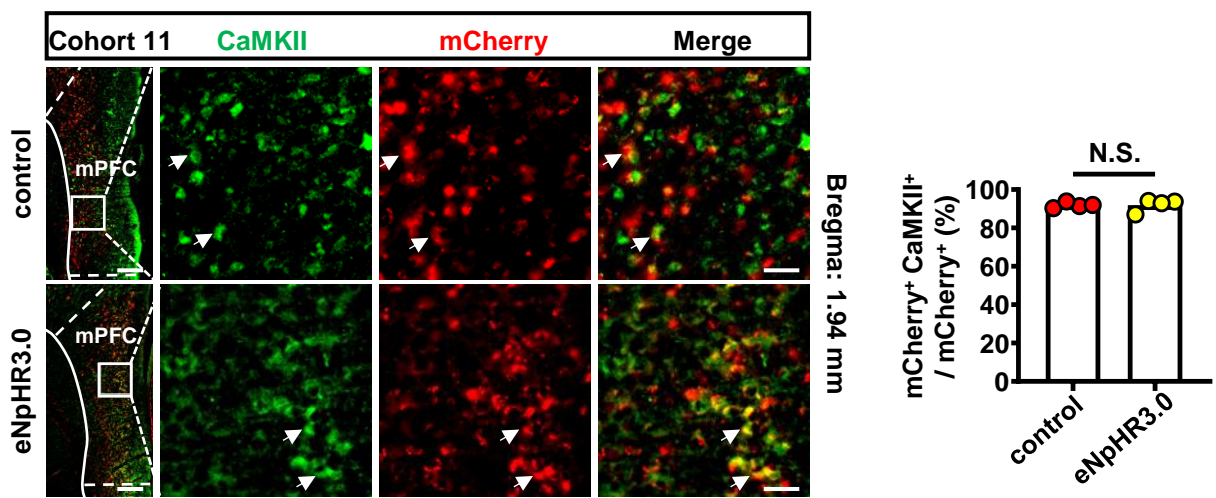
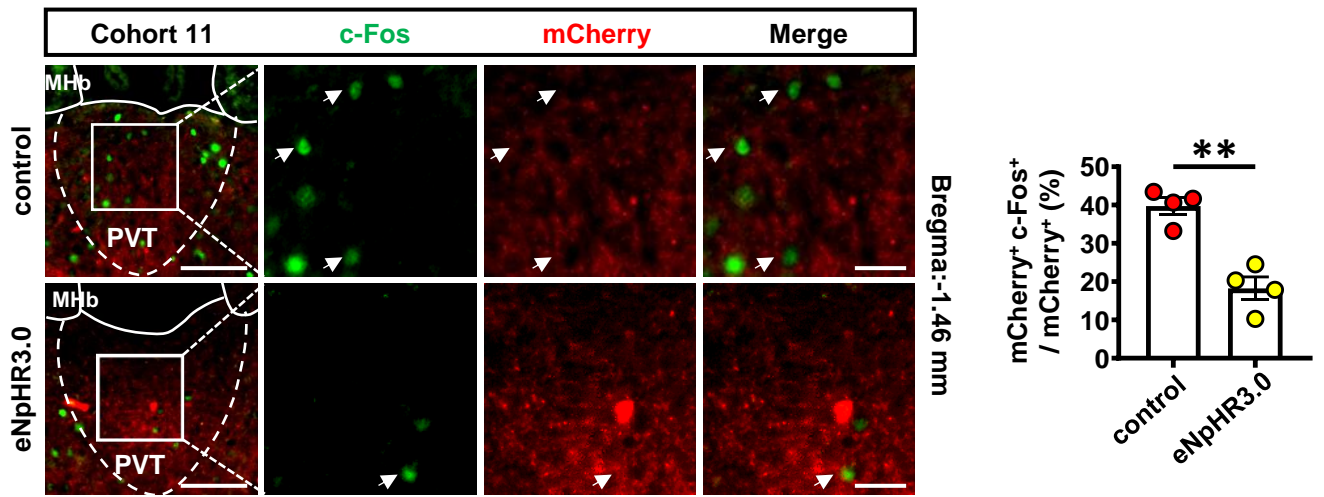
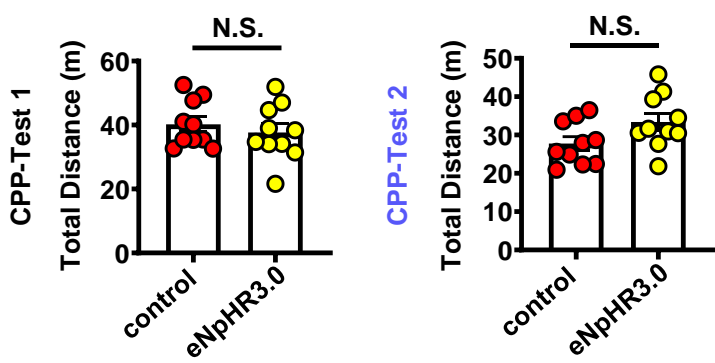
A



B

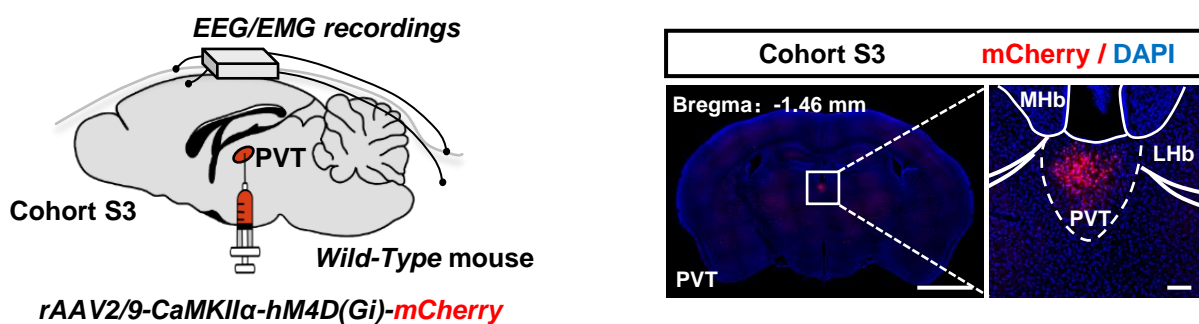




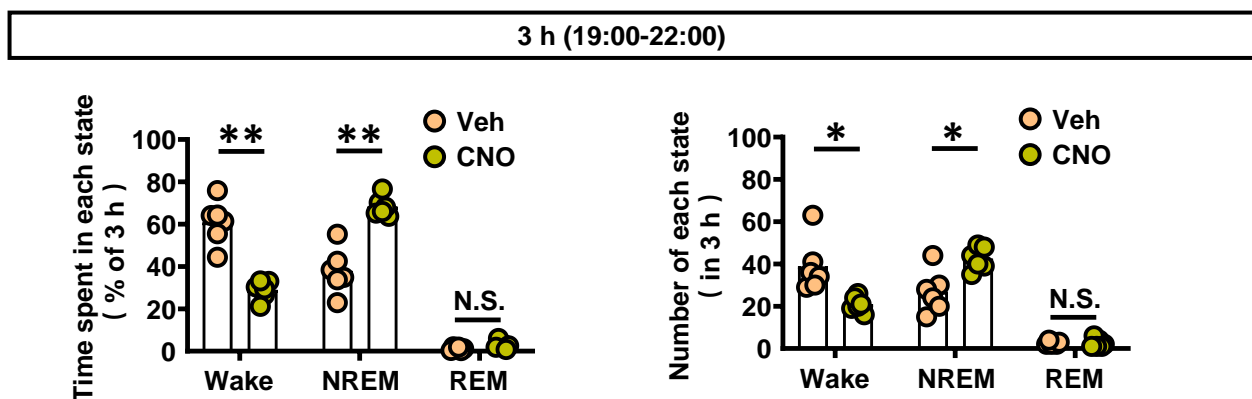
**Figure S7****A****B****C**

**Figure S8**

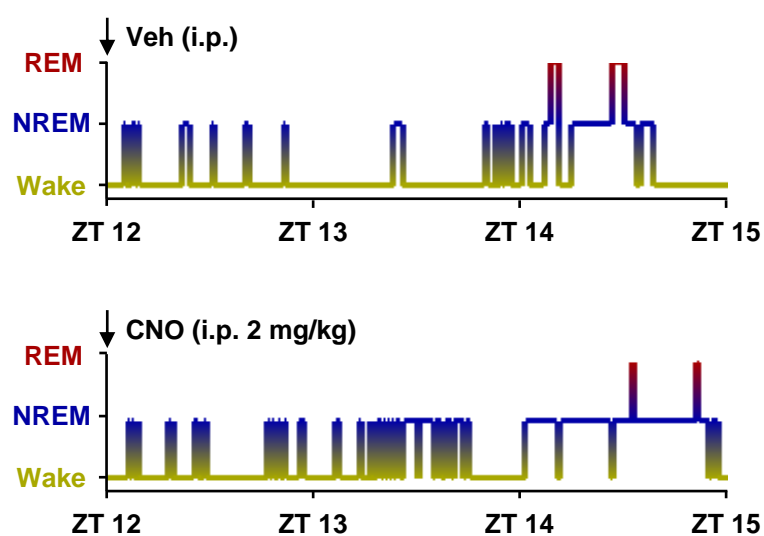
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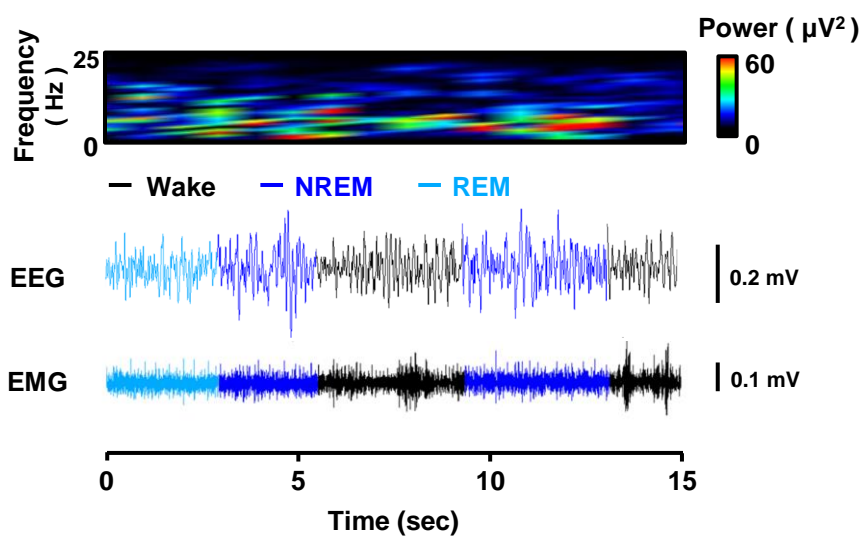
**B**



**C**



**D**



**E**

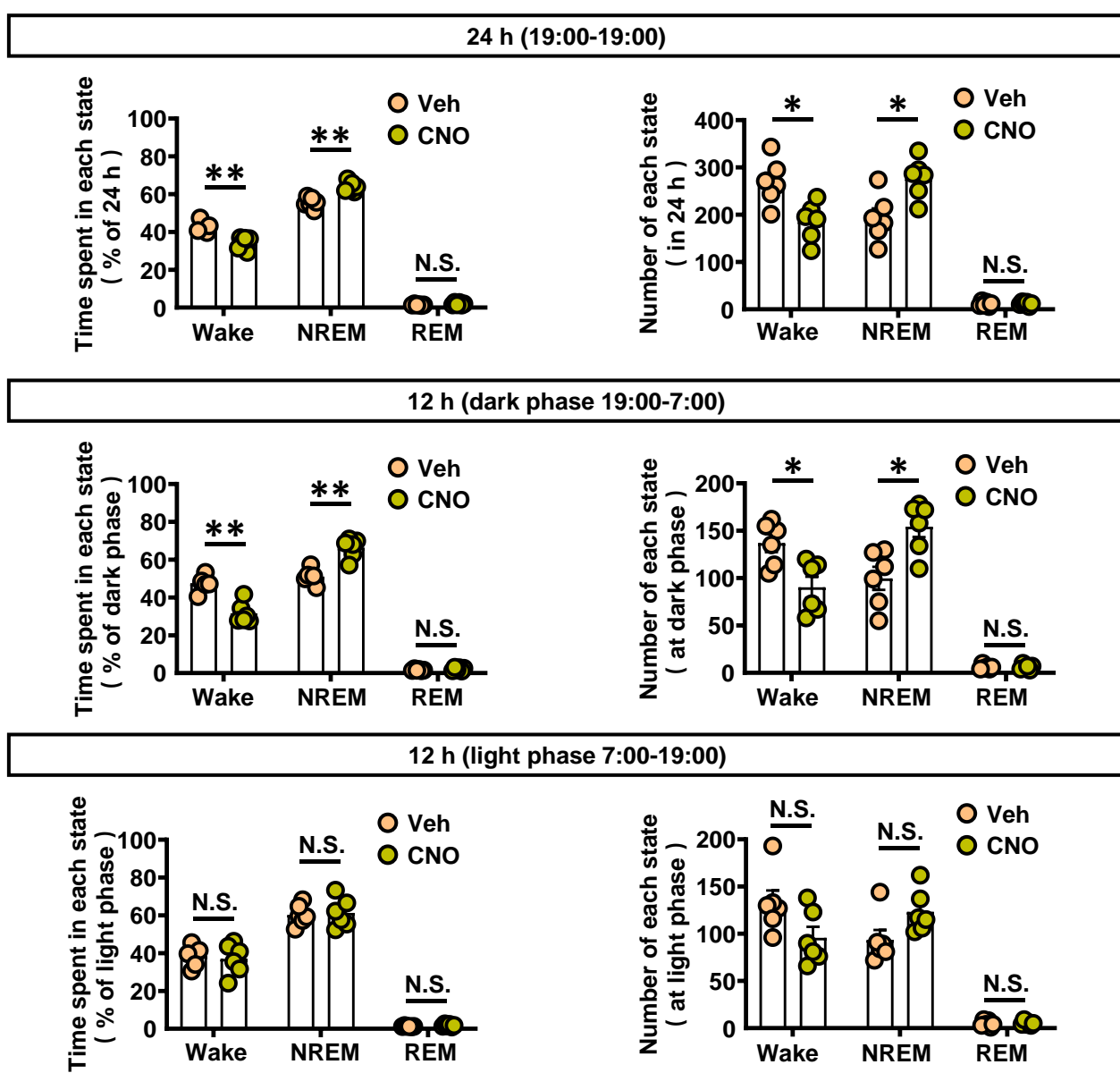


Figure S9

A

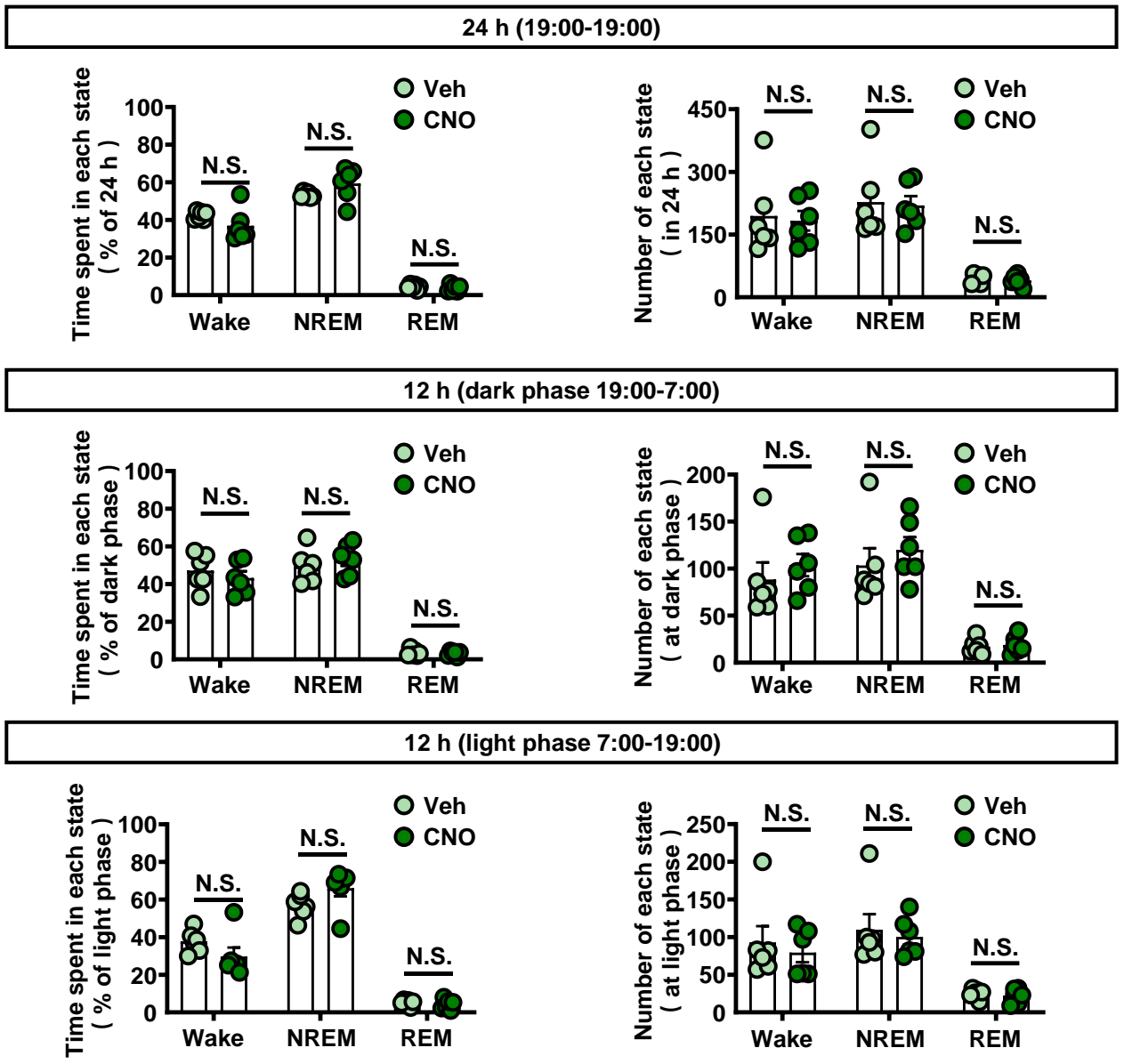
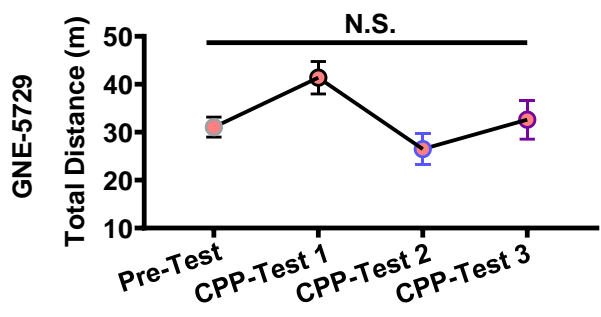


Figure S10

A

○ Pre-Test    ○ CPP-Test 2  
○ CPP-Test 1    ○ CPP-Test 3



○ Pre-Test    ○ CPP-Test 2  
○ CPP-Test 1    ○ CPP-Test 3

