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## Supplementary Materials for

### **Development of a CRISPR-SaCas9 system for projection- and function-specific gene editing in the rat brain**

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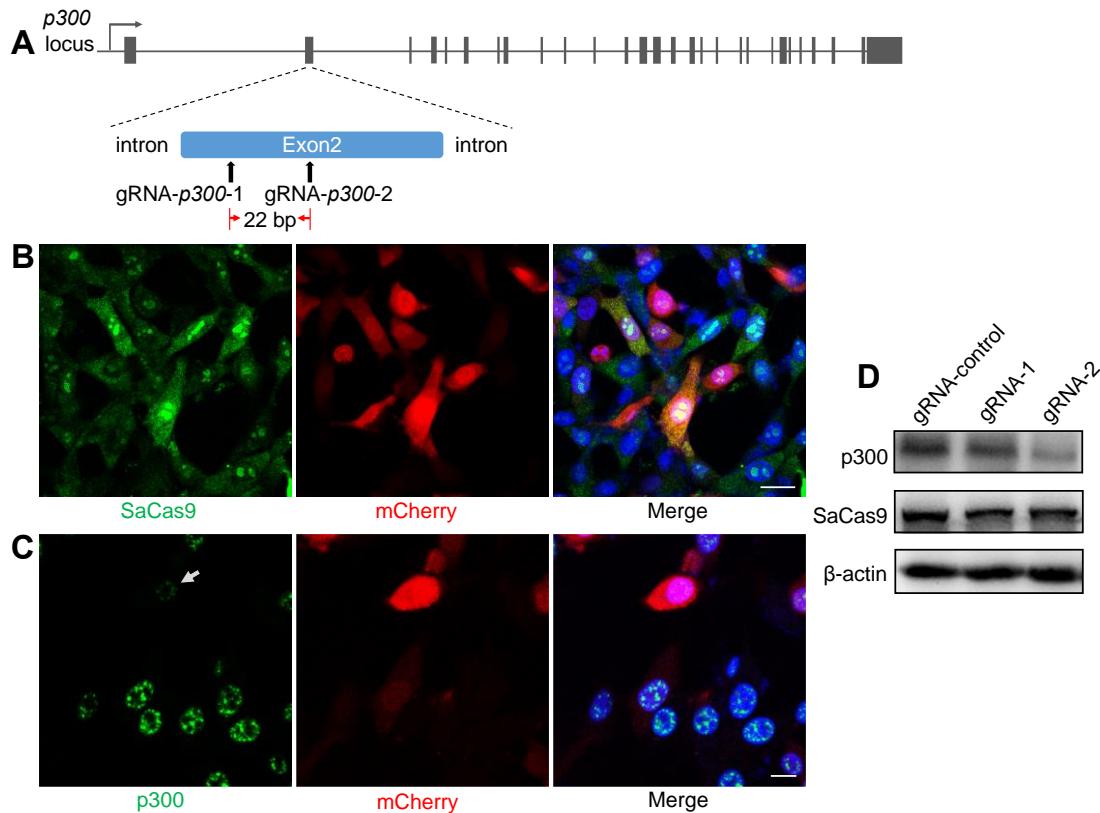
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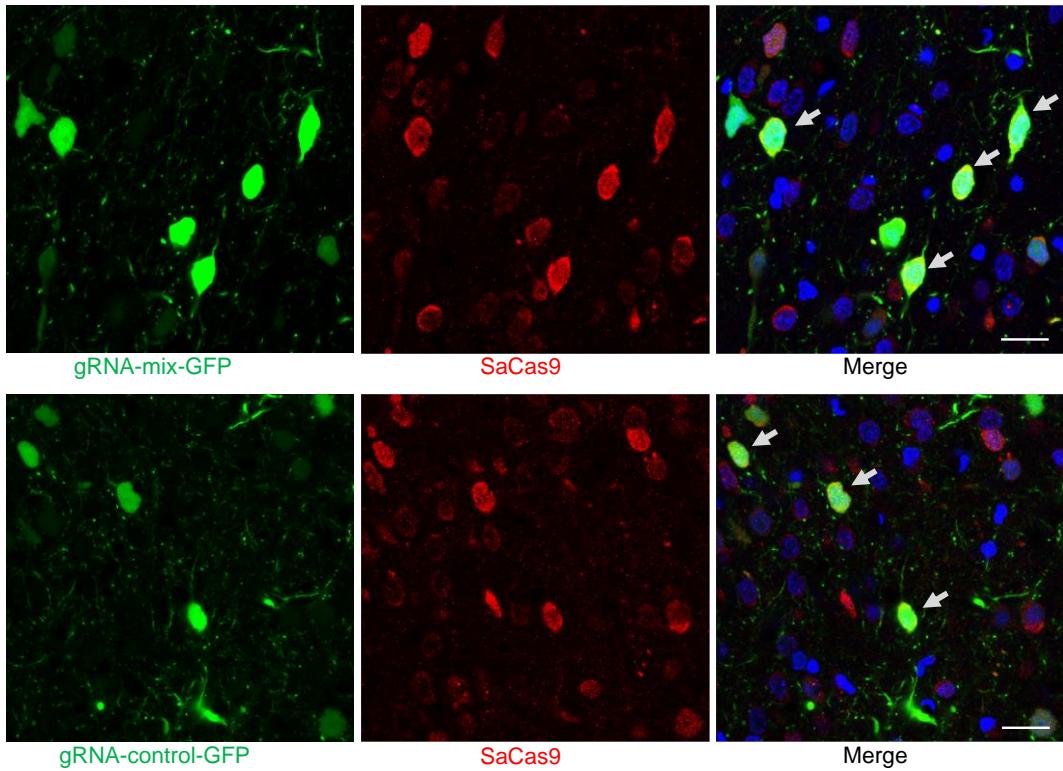
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## Supplementary Materials

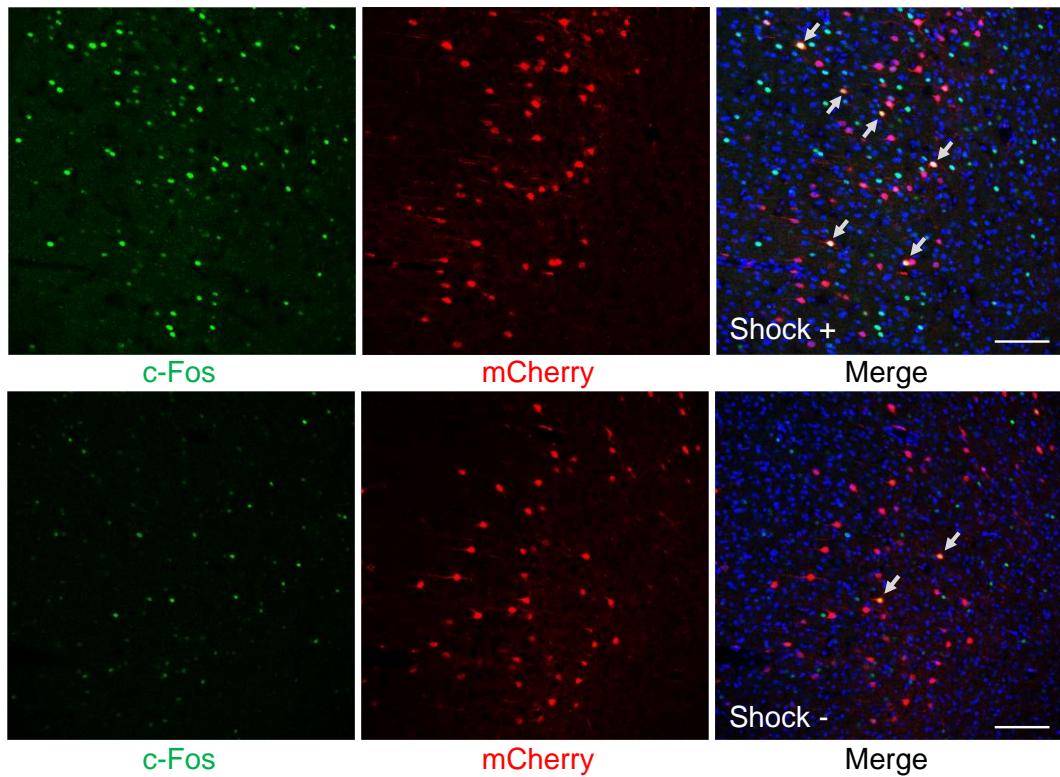


**Fig. S1. SaCas9 inactivates the *p300* gene in vitro with high efficiency.**

(A) Graphical representation of the rat *p300* gene locus showing gRNA target sites. (B) Elevated expression of SaCas9 shown by immunofluorescence three days after lentivirus transfection in F98 cells. Scale bars, 20  $\mu$ m. (C) Immunofluorescent staining of p300 one week after CRISPR-SaCas9 targeting of *p300* locus in F98 cells, with white arrows indicating decreased p300 expression in SaCas9- and gRNA-positive cells. Scale bars, 10  $\mu$ m. (D) Western blot analysis showing decreased p300 expression in F98 cells one week after lentivirus transfection.



**Fig. S2. Neuronal specificity for spine density analysis.** The presence of SaCas9 in gRNA-mix-GFP<sup>+</sup> (upper) and gRNA-control-GFP<sup>+</sup> (lower) neurons for morphology analysis was verified by immunofluorescence. Scale bars, 20  $\mu$ m.



**Fig. S3. Amygdala-projecting IL neurons are activated during extinction.**

Illustration of amygdala-projecting IL neurons ( $\text{mCherry}^+$ ) co-labeled with c-Fos 2 h after context exposure in the shock + (with shock during CFC training) and shock - (without shock during CFC training) groups. Scale bars, 100  $\mu\text{m}$ .

**Table S1. gRNA sequences targeting *cbp* gene or *p300* gene.**

	<b>gRNA sequence</b>	<b>PAM</b>	<b>Exon</b>	<b>Direction</b>
gRNA-control	GGAGACGAATAATGCGTCTCC	/	/	/
CBP-gRNA-1	CTGGCTGCCTGTTGGGCAGG	CTGGGT	2	-
CBP-gRNA-2	AAATGTCAGCGACGAGAGCAA	GCGAAT	4	+
CBP-gRNA-3	AGCAGCTGTGGAAGCAGGAGG	TGGAGT	14	-
CBP-gRNA-4	ATTCAGTGCTGGAGGCAGG	GGGAGT	2	-
CBP-gRNA-5	AGTAACCTGGCCATAGCTTA	ATGAAT	2	+
p300-gRNA-1	TAGTTCCCCAACCTCAATAT	GGGAGT	2	+
p300-gRNA-2	GTTATGCCAGCCAGGCCAA	CAGAAT	2	+

**Table S2. AAV vectors used.**

<b>AAV</b>	<b>Titer (v.g./ml)</b>	<b>Serotype</b>
AAV-hSyn-GFP	$9.3 \times 10^{12}$	9
AAV-CaMKII-Cre	$8.1 \times 10^{13}$	1
AAV-CaMKII-Cre-P2A-GFP	$5.2 \times 10^{13}$	1
AAV-hSyn-Cre-P2A-GFP	$4.5 \times 10^{13}$	1
AAV-hSyn-DIO-mCherry	$1.8 \times 10^{13}$	9
AAV-hSyn-Cre	$6.7 \times 10^{13}$	retro
AAV-hSyn-Cre-P2A-GFP	$1.7 \times 10^{13}$	retro
AAV-hSyn-mCherry	$5.4 \times 10^{13}$	retro
AAV-hSyn-DIO-SaCas9-3×flag	$3.3 \times 10^{13}$	9
AAV-hSyn-SaCas9-3×flag	$4.1 \times 10^{13}$	9
AAV-U6-gRNA1-U6-gRNA4-U6-gRNA5-CMV-GFP	$1.2 \times 10^{13}$	9
AAV-c-fos-rtTA-U6-gRNA1-U6-gRNA4-U6-gRNA5	$1.7 \times 10^{13}$	9
AAV-c-fos-rtTA	$3.7 \times 10^{13}$	9
AAV-TRE <sup>3G</sup> -SaCas9-3×flag	$2.2 \times 10^{13}$	9
AAV-TRE <sup>3G</sup> -DIO-SaCas9-3×flag	$3.2 \times 10^{13}$	9

**Table S3. Primer sequences for T7 endonuclease assays.**

gRNA-1-F	GATTGGATCATGGTT
gRNA-1-R	CCTGTGCAAAGTCTCA
gRNA-2-F	GCTGGTGCTGTTCTGG
gRNA-2-R	CAAATCACCTGCCTTCT
gRNA-3-F	TAAGGTGCCCTGTATGA
gRNA-3-R	GTCGGTAAACTAAGTAAATGTG

gRNA-4-F	TGTGCCGTATTCAGTCC
gRNA-4-R	CTGCGTCCAAACATAAAC
gRNA-5-F	CCAGATGCTGCGTCAA
gRNA-5-R	CAACAACCCGTTCTTC
gRNA-1+4+5-F	CGGGATAGGCAATGTG
gRNA-1+4+5-R	CTGGAGCAGGGTAGGGCATT

**Table S4. Primer sequences for off-target analysis.**

gRNA-1-F	CTGTGATGTGGCAGGACTAC
gRNA-1-R	CGGGATAGGCAATGTG
off-target 1-F	TTGCTTCAGACTCCACATTG
off-target 1-R	TGAAGAACTTGAGAAGAAGGG
off-target 2-F	AATTATGTCCCACCCAATGA
off-target 2-R	AGACTGATTTCCCTTAAG
off-target 3-F	GCCCCCTTAAATCGAACTC
off-target 3-R	TCACCCGAGACCATGTAGAG
off-target 4-F	ACGTTAAGAGATGGCATCTG
off-target 4-R	AGTTTGCACTCATAGCCAA
off-target 5-F	AGTTAAAAACCAAGGCAC
off-target 5-R	TCAGTTGATTAGTCGC
off-target 6-F	TGTTCTGCTAAACTCGAGGC
off-target 6-R	AAGAAGTCAGGCGTGCT
off-target 7-F	ACAGCATGACACCGTGTGAG
off-target 7-R	GGAAAGGAAGGCAGGTAGAC
off-target 8-F	CAGAAAGGAAGTCAGGGTAT
off-target 8-R	AACAGCAGCATTAGCCA
off-target 9-F	TTCTGTGTTGGCATGGTATT
off-target 9-R	TGTTTCGTGGAGCTATTCA
gRNA-4-F	AAGCTATGCCAGAGTTACT
gRNA-4-R	GCATGCCAGTTAGG
off-target 10-F	AAGCCACTGCCTGTATTCC
off-target 10-R	CAATGAGCAGTTGGTCCC
off-target 11-F	GGGACCATGAAAGCATTGA
off-target 11-R	GCCCTGAGTCTAGAGCATGT
off-target 12-F	AAAATGGAGCATACTTAGAG
off-target 12-R	CCAACATCATGCAAAGTGTG
off-target 13-F	TGATAACCTACTCCCTAAC
off-target 13-R	GCGAACAGTAAGGACT
off-target 14-F	CCAAATTAAAATCCAGTTAG
off-target 14-R	GATGACCTGGAACTCTT
off-target 15-F	CTTCCCAAAGAGCAGATTAG
off-target 15-R	GAGGCACATTAAAGTCCC

off-target 16-F	GCTGAAACAGTATAAGTCGG
off-target 16-R	GCCTGCTATTAAGACCTT
off-target 17-F	GGACCGTTCCCTCAGACC
off-target 17-R	ACCGATAATCCCAGCTCTAC
off-target 18-F	TTAAAGGGTTGCAGCGTTAG
off-target 18-R	TGGTTAAGTGGGAATGTC
off-target 19-F	CCCTTGAGTTATTGTTGAG
off-target 19-R	CACCAAAGGTTCTGGTTAGA
off-target 20-F	GTCTGTGAGACATAACCTGAGG
off-target 20-R	ATGTGCAGGCAGCCATTAC
off-target 21-F	AAAACAAGACAACTTGGC
off-target 21-R	TTAGCAAGACTACACGGAGA
off-target 22-F	TCCAATCTGAGGGAGTGTA
off-target 22-R	TCCCATTCCAATGATTAC
off-target 23-F	ACGATATTAAAGGTGGTTAG
off-target 23-R	TGTTAGGACTCCCACATAA
off-target 24-F	AGACTAGCCTCAAACATC
off-target 24-R	ATAAAGTGAATTCAAGGACTC
off-target 25-F	CGGCCAATGGATGCTA
off-target 25-R	GTCCTCCCCACTTGATCACT
off-target 26-F	AATTAAGAGAAAAGAACAC
off-target 26-R	ATGGTTAAATTAGCAAT
off-target 27-F	CTCGTCAATAAAAGACAGTA
off-target 27-R	AGAAGAGGAAAATCTCAT
off-target 28-F	AAGGGTAAGAAACTTCC
off-target 28-R	AAAACTAAGGTAACATTG
off-target 29-F	CTACCAGCTGGAAATCCTC
off-target 29-R	ACTGCTGGAGTGAGCTCTG
gRNA-5-F	AAGCACTGAATCCGCAAGC
gRNA-5-R	CTGGAGCAGGGTAGGGCATT
off-target 30-F	TCCACCTCCTGTGAACCC
off-target 30-R	AATTCTAGAGGCTGATATCTGC
off-target 31-F	TAGGGCTTCATACTTGC
off-target 31-R	AGAAAAGTGTGGGACCTAT
off-target 32-F	GATGGGTGAGGAAGGTGATA
off-target 32-R	GGCTTCTGCTCTATGGC
off-target 33-F	AGATGGCACAACTACTCACCT
off-target 33-R	TTGGGTTCCATCCCTAGTAC