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Main Figures: 7

Supplementary Figures: 10

Supplementary Tables: 0

Supplementary Videos: 1

Reporting Checklist for Nature Neuroscience

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Please note that in the event of publication, it is mandatory that authors include all relevant methodological and statistical information in the manuscript.

► Statistics reporting, by figure

- Please specify the following information for each panel reporting quantitative data, and where each item is reported (section, e.g. Results, & paragraph number).
- Each figure legend should ideally contain an exact sample size (n) for each experimental group/condition, where n is an exact number and not a range, a clear definition of how n is defined (for example x cells from x slices from x animals from x litters, collected over x days), a description of the statistical test used, the results of the tests, any descriptive statistics and clearly defined error bars if applicable.
- For any experiments using custom statistics, please indicate the test used and stats obtained for each experiment.
- Each figure legend should include a statement of how many times the experiment shown was replicated in the lab; the details of sample collection should be sufficiently clear so that the replicability of the experiment is obvious to the reader.
- For experiments reported in the text but not in the figures, please use the paragraph number instead of the figure number.

Note: Mean and standard deviation are not appropriate on small samples, and plotting independent data points is usually more informative. When technical replicates are reported, error and significance measures reflect the experimental variability and not the variability of the biological process; it is misleading not to state this clearly.

		TEST USED		n			DESCRIPTIVE STATS (AVERAGE, VARIANCE)		P VALUE		DEGREES OF FREEDOM & F/t/z/R/ETC VALUE	
		WHICH TEST?	SECTION & PARAGRAPH #	EXACT VALUE	DEFINED?	SECTION & PARAGRAPH #	REPORTED?	SECTION & PARAGRAPH #	EXACT VALUE	SECTION & PARAGRAPH #	VALUE	SECTION & PARAGRAPH #
example	1a	one-way ANOVA	Fig. legend	9, 9, 10, 15	mice from at least 3 litters/group	Methods para 8	error bars are mean +/- SEM	Fig. legend	p = 0.044	Fig. legend	F(3, 36) = 2.97	Fig. legend
example	results, para 6	unpaired t-test	Results para 6	15	slices from 10 mice	Results para 6	error bars are mean +/- SEM	Results para 6	p = 0.0006	Results para 6	t(28) = 2.808	Results para 6
+ - +	1d	one-way ANOVA	Fig. legend	6, 7, 7, 8	brain sections from 4 animals	Fig. legend	mean (+), median, quartiles (boxes), range (whiskers)	Fig. legend	P < 0.0001	Fig. legend	F(3, 24) = 45.5	Fig. legend

		TEST USED		n			DESCRIPTIVE STATS (AVERAGE, VARIANCE)		P VALUE		DEGREES OF FREEDOM & F/t/z/R/ETC VALUE	
FIGURE NUMBER	WHICH TEST?	SECTION & PARAGRAPH #	EXACT VALUE	DEFINED?	SECTION & PARAGRAPH #	REPORTED?	SECTION & PARAGRAPH #	EXACT VALUE	SECTION & PARAGRAPH #	VALUE	SECTION & PARAGRAPH #	
+ -	1e (CCK)	unpaired t test	Fig. legend	7, 7	brain sections from 4 animals	Fig. legend	error bars are mean+/- SEM	Fig. legend	P < 0.0001	Fig. legend	t(12) = 6.85	Fig. legend
+ -	1e (LiCl)	unpaired t test	Fig. legend	7, 7	brain sections from 4 animals	Fig. legend	error bars are mean+/- SEM	Fig. legend	P < 0.0001	Fig. legend	t(12) = 7.03	Fig. legend
+ -	1e (LPS)	unpaired t test	Fig. legend	8, 8	brain sections from 4 animals	Fig. legend	error bars are mean+/- SEM	Fig. legend	P = 0.33	Fig. legend	t(14) = 1.00	Fig. legend
+ -	1f	unpaired t test	Fig. legend	15, 17	brain sections from 4-5 animals	Fig. legend	mean (+), median, quartiles (boxes), range (whiskers)	Fig. legend	p < 0.0001	Fig. legend	t(30) = 8.43	Fig. legend
+ -	1g refed	unpaired t test	Fig. legend	17, 17	brain sections from 4-5 animals	Fig. legend	error bars are mean+/- SEM	Fig. legend	p < 0.0001	Fig. legend	t(32) = 8.09	Fig. legend
+ -	1h	One-way ANOVA	Fig. legend	12, 15, 12	brain sections from 3 animals	Fig. legend	mean (+), median, quartiles (boxes), range (whiskers)	Fig. legend	P < 0.0001	Fig. legend	F(2, 36) = 73.2	Fig. legend
+ -	1i	unpaired t test	Fig. legend	15, 15	brain sections from 3 animals	Fig. legend	error bars are mean+/- SEM	Fig. legend	p = 0.0019	Fig. legend	t(28) = 3.42	Fig. legend
+ -	2b saline	Two-way ANOVA	Fig. legend	6, 6, 8, 8	mice from at least 3 litters	Fig.	error bars are mean+/- SEM	Fig. legend	p = 0.55	Fig. legend	F(1, 24) = 0.363	Fig. legend
+ -	2b CCK	Two-way ANOVA	Fig. legend	6, 10, 8, 11	mice from at least 3 litters	Fig.	error bars are mean+/- SEM	Fig. legend	p = 0.0018	Fig. legend	F(1, 31) = 11.7	Fig. legend
+ -	2b LiCl	Two-way ANOVA	Fig. legend	6, 10, 8, 10	mice from at least 3 litters	Fig.	error bars are mean+/- SEM	Fig. legend	p = 0.042	Fig. legend	F(1, 30) = 4.50	Fig. legend
+ -	2b LPS	Two-way ANOVA	Fig. legend	8, 8, 8, 12	mice from at least 3 litters	Fig.	error bars are mean+/- SEM	Fig. legend	P = 0.98	Fig. legend	F(3, 32) = 0.001	Fig. legend
+ -	2c	Two-way ANOVA	Fig. legend	5, 5, 5, 5	mice from at least 3 litters	Fig. legend	error bars are mean+/- SEM	Fig. legend	P = 0.0062	Fig. legend	F(1, 16) = 9.91	Fig. legend
+ -	2d	unpaired t test	Fig. legend	13, 14	mice from at least 3 litters	Fig. legend	mean (+), median, quartiles (boxes), range (whiskers)	Fig. legend	p = 0.045	Fig. legend	t(25) = 2.2	Fig. legend
+ -	3d	unpaired t test	Fig. legend	7, 7	mice from at least 3 litters	Fig. legend	mean (+), median, quartiles (boxes), range (whiskers)	Fig. legend	p = 0.026	Fig. legend	t(12) = 2.5	Fig. legend
+ -	3f	unpaired t test	Fig. legend	7, 7	mice from at least 3 litters	Fig. legend	mean (+), median, quartiles (boxes), range (whiskers)	Fig. legend	p = 0.18	Fig. legend	t(12) = 1.4	Fig. legend
+ -	3g	unpaired t test	Fig. legend	7, 7	mice from at least 3 litters	Fig. legend	mean (+), median, quartiles (boxes), range (whiskers)	Fig. legend	p = 0.0048	Fig. legend	t(12) = 3.5	Fig. legend
+ -	3h	unpaired t test	Fig. legend	7, 7	mice from at least 3 litters	Fig. legend	mean (+), median, quartiles (boxes), range (whiskers)	Fig. legend	p = 0.82	Fig. legend	t(12) = 0.24	Fig. legend
+ -	4b	One-way ANOVA	Fig. legend	4, 5, 4	mice from at least 3 litters	Fig. legend	mean (+), median, quartiles (boxes), range (whiskers)	Fig. legend	p < 0.0001	Fig. legend	F(2, 10) = 45.1	Fig. legend
+ -	4c	unpaired t test	Fig. legend	4, 5	mice from at least 3 litters	Fig. legend	mean (+), median, quartiles (boxes), range (whiskers)	Fig. legend	P < 0.0001	Fig. legend	t(7) = 8.02	Fig. legend
+ -	4e	unpaired t test	Fig. legend	8, 11	mice from at least 3 litters	Fig. legend	mean (+), median, quartiles (boxes), range (whiskers)	Fig. legend	p = 0.0058	Fig. legend	t(17) = 3.15	Fig. legend

+ -	4g	unpaired t test	Fig. legend	4, 5	mice from at least 3 litters	Fig. legend	mean (+), median, quartiles (boxes), range (whiskers)	Fig. legend	$p = 0.022$	Fig. legend	$t(7) = 2.9$	Fig. legend
+ -	4h	unpaired t test	Fig. legend	4, 5	mice from at least 3 litters	Fig. legend	mean (+), median, quartiles (boxes), range (whiskers)	Fig. legend	$p = 0.0083$	Fig. legend	$t(7) = 3.6$	Fig. legend
+ -	5a	unpaired t test	Fig. legend	4, 5	mice from at least 3 litters	Fig. legend	mean (+), median, quartiles (boxes), range (whiskers)	Fig. legend	$p = 0.0051$	Fig. legend	$t(7) = 4.01$	Fig. legend
+ -	5b	unpaired t test	Fig. legend	4, 5	mice from at least 3 litters	Fig. legend	mean (+), median, quartiles (boxes), range (whiskers)	Fig. legend	$p = 0.029$	Fig. legend	$t(7) = 2.74$	Fig. legend
+ -	5c	unpaired t test	Fig. legend	5, 7	mice from at least 3 litters	Fig. legend	mean (+), median, quartiles (boxes), range (whiskers)	Fig. legend	$p = 0.031$	Fig. legend	$t(10) = 2.50$	Fig. legend
+ -	5d	One-way ANOVA	Fig. legend	4, 5, 5	mice from at least 3 litters	Fig. legend	error bars are mean \pm SEM	Fig. legend	$p < 0.0001$	Fig. legend	$F(2, 13) = 44.5$	Fig. legend
+ -	5e saline vs 1mg/kg	unpaired t test	Fig. legend	4, 4	mice from at least 3 litters	Fig. legend	error bars are mean \pm SEM	Fig. legend	$p = 0.032$	Fig. legend	$t(6) = 2.78$	Fig. legend
+ -	7e	paired t test	Fig. legend	9, 9	mice from at least 3 litters	Fig. legend	mean (+), median, quartiles (boxes), range (whiskers)	Fig. legend	$p = 0.0030$	Fig. legend	$t(16) = 3.46$	Fig. legend
+ -	7g	unpaired t test	Fig. legend	10, 9	mice from at least 3 litters	Fig. legend	mean (+), median, quartiles (boxes), range (whiskers)	Fig. legend	$p = 0.65$	Fig. legend	$t(17) = 0.458$	Fig. legend
+ -	7c	Two-way ANOVA	Fig. legend	7, 7, 7, 7	mice from at least 3 litters	Fig. legend	error bars are mean \pm SEM	Fig. legend	$p = 0.00092$	Fig. legend	$t(12) = 4.37$	Fig. legend
+ -	7c Chr2	paired t test	Fig. legend	7, 7	mice from at least 3 litters	Fig. legend	error bars are mean \pm SEM	Fig. legend	$p = 0.0021$	Fig. legend	$t(12) = 3.85$	Fig. legend
+ -	6c	One-way ANOVA	Fig. legend	6, 6, 6	sections from two injections	Fig. legend	error bars are mean \pm SEM	Fig. legend	$p < 0.0001$	Fig. legend	$F(2, 15) = 23.1$	Fig. legend
+ -	6d	One-way ANOVA	Fig. legend	6, 6, 6	sections from two injections	Fig. legend	error bars are mean \pm SEM	Fig. legend	$p = 0.0008$	Fig. legend	$F(2, 15) = 12.0$	Fig. legend
+ -	6f	One-way ANOVA	Fig. legend	6, 6, 6	sections from two injections	Fig. legend	error bars are mean \pm SEM	Fig. legend	$p < 0.0001$	Fig. legend	$F(2, 15) = 42.4$	Fig. legend
+ -	6g	One-way ANOVA	Fig. legend	6, 6, 6	sections from two injections	Fig. legend	error bars are mean \pm SEM	Fig. legend	$p < 0.0001$	Fig. legend	$F(2, 15) = 25.1$	Fig. legend
+ -	6i	One-way ANOVA	Fig. legend	6, 6, 6	sections from two injections	Fig. legend	error bars are mean \pm SEM	Fig. legend	$p = 0.0003$	Fig. legend	$F(2, 15) = 14.6$	Fig. legend
+ -	6j	One-way ANOVA	Fig. legend	6, 6, 6	sections from two injections	Fig. legend	error bars are mean \pm SEM	Fig. legend	$p < 0.0001$	Fig. legend	$F(2, 15) = 18.8$	Fig. legend
+ -	supp 1a left	One-way ANOVA	Fig. legend	6, 7, 7, 8	brain sections from 4 animals	Fig. legend	error bars are mean \pm SEM	Fig. legend	$p < 0.0001$	Fig. legend	$F(3, 24) = 29.1$	Fig. legend
+ -	supp 1a right	One-way ANOVA	Fig. legend	6, 7, 7, 8	brain sections from 4 animals	Fig. legend	error bars are mean \pm SEM	Fig. legend	$p < 0.0001$	Fig. legend	$F(3, 24) = 39.8$	Fig. legend
+ -	supp 1b left	Unpaired t test	Fig. legend	15, 17	brain sections from 4-5 animals	Fig. legend	mean (+), median, quartiles (boxes), range (whiskers)	Fig. legend	$p < 0.0001$	Fig. legend	$t(30) = 7.88$	Fig. legend
+ -	supp 1b right	Unpaired t test	Fig. legend	15, 17	brain sections from 4-5 animals	Fig. legend	mean (+), median, quartiles (boxes), range (whiskers)	Fig. legend	$p = 0.0004$	Fig. legend	$t(30) = 3.95$	Fig. legend
+ -	supp 1c left	One-way ANOVA	Fig. legend	12, 15, 12	brain sections from 3 animals	Fig. legend	error bars are mean \pm SEM	Fig. legend	$p < 0.0001$	Fig. legend	$F(2, 36) = 34.1$	Fig. legend
+ -	supp 1c right	One-way ANOVA	Fig. legend	12, 15, 12	brain sections from 3 animals	Fig. legend	error bars are mean \pm SEM	Fig. legend	$p = 0.0028$	Fig. legend	$F(2, 36) = 6.97$	Fig. legend

+	supp - 2b total Fos	Unpaired t- test	Fig. legend	4, 4	mice from at least 3 litters	Fig. legend	error bars are mean+/- SEM	Fig. legend	p = 0.28	Fig. legend	t(6) = 1.2	Fig. legend
+	supp - 2b in PKC- δ+	Unpaired t- test	Fig. legend	4, 4	mice from at least 3 litters	Fig. legend	error bars are mean+/- SEM	Fig. legend	p < 0.0001	Fig. legend	t(6) = 8.6	Fig. legend
+	supp - 2b in PKC- δ-	Unpaired t- test	Fig. legend	4, 4	mice from at least 3 litters	Fig. legend	error bars are mean+/- SEM	Fig. legend	p = 0.0011	Fig. legend	t(6) = 5.9	Fig. legend
+	supp - 2b Fos/ PKC- δ+	Unpaired t- test	Fig. legend	4, 4	mice from at least 3 litters	Fig. legend	error bars are mean+/- SEM	Fig. legend	p = 0.0007	Fig. legend	t(6) = 6.4	Fig. legend
+	supp - 2b PKC- δ/ Fos	Unpaired t- test	Fig. legend	4, 4	mice from at least 3 litters	Fig. legend	error bars are mean+/- SEM	Fig. legend	p < 0.0001	Fig. legend	t(6) = 9.8	Fig. legend
+	supp - 2c total Fos	Unpaired t- test	Fig. legend	4, 4	mice from at least 3 litters	Fig. legend	error bars are mean+/- SEM	Fig. legend	p = 0.021	Fig. legend	t(6) = 3.1	Fig. legend
+	supp - 2c in PKC- δ+	Unpaired t- test	Fig. legend	4, 4	mice from at least 3 litters	Fig. legend	error bars are mean+/- SEM	Fig. legend	p < 0.0012	Fig. legend	t(6) = 5.8	Fig. legend
+	supp - 2c in PKC- δ-	Unpaired t- test	Fig. legend	4, 4	mice from at least 3 litters	Fig. legend	error bars are mean+/- SEM	Fig. legend	p < 0.0001	Fig. legend	t(6) = 12.7	Fig. legend
+	supp - 2c Fos/ PKC- δ+	Unpaired t- test	Fig. legend	4, 4	mice from at least 3 litters	Fig. legend	error bars are mean+/- SEM	Fig. legend	p < 0.0001	Fig. legend	t(6) = 8.9	Fig. legend
+	supp - 2c PKC- δ/ Fos	Unpaired t- test	Fig. legend	4, 4	mice from at least 3 litters	Fig. legend	error bars are mean+/- SEM	Fig. legend	p < 0.0001	Fig. legend	t(6) = 11.5	Fig. legend
+	supp - 2d total Fos	Unpaired t- test	Fig. legend	4, 4	mice from at least 3 litters	Fig. legend	error bars are mean+/- SEM	Fig. legend	p = 0.20	Fig. legend	t(6) = 1.5	Fig. legend
+	supp - 2d in PKC- δ+	Unpaired t- test	Fig. legend	4, 4	mice from at least 3 litters	Fig. legend	error bars are mean+/- SEM	Fig. legend	p < 0.0076	Fig. legend	t(6) = 3.9	Fig. legend
+	supp - 2d in PKC- δ-	Unpaired t- test	Fig. legend	4, 4	mice from at least 3 litters	Fig. legend	error bars are mean+/- SEM	Fig. legend	p = 0.0045	Fig. legend	t(6) = 4.4	Fig. legend
+	supp - 2d Fos/ PKC- δ+	Unpaired t- test	Fig. legend	4, 4	mice from at least 3 litters	Fig. legend	error bars are mean+/- SEM	Fig. legend	p = 0.0014	Fig. legend	t(6) = 3.4	Fig. legend
+	supp - 2d PKC- δ/ Fos	Unpaired t- test	Fig. legend	4, 4	mice from at least 3 litters	Fig. legend	error bars are mean+/- SEM	Fig. legend	p < 0.0001	Fig. legend	t(6) = 12.5	Fig. legend

+	supp -	3b	Two-way ANOVA	Fig. legend	7, 5	mice from at least 3 litters	Fig. legend	error bars are mean+/- SEM	Fig. legend	p = 0.82	Fig. legend	F(1, 35) = 0.051	Fig. legend
+	supp -	3c	Two-way ANOVA	Fig. legend	8, 7	mice from at least 3 litters	Fig. legend	error bars are mean+/- SEM	Fig. legend	p = 0.033	Fig. legend	F(1, 130) = 4.67	Fig. legend
+	supp -	3d	Unpaired t-test	Fig. legend	8, 7	mice from at least 3 litters	Fig. legend	mean (+), median, quartiles (boxes), range (whiskers)	Fig. legend	p = 0.28	Fig. legend	t(13) = 1.13	Fig. legend
+	supp -	3e	Unpaired t-test	Fig. legend	8, 7	mice from at least 3 litters	Fig. legend	mean (+), median, quartiles (boxes), range (whiskers)	Fig. legend	p = 0.031	Fig. legend	t(13) = 2.42	Fig. legend
+	supp -	3i	Unpaired t-test	Fig. legend	10, 8	stimulation events from 3 mice	Fig. legend	mean (+), median, quartiles (boxes), range (whiskers)	Fig. legend	p = 0.67	Fig. legend	t(16) = 0.429	Fig. legend
+	supp -	4a	Unpaired t-test	Fig. legend	18, 18	stimulation events from 11 mice	Fig. legend	error bars are mean+/- SEM	Fig. legend	p = 0.24	Fig. legend	t(34) = 1.21	Fig. legend
+	supp -	4b	Unpaired t-test	Fig. legend	18, 18	stimulation events from 11 mice	Fig. legend	error bars are mean+/- SEM	Fig. legend	p = 0.73	Fig. legend	t(34) = 0.342	Fig. legend
+	supp -	4d	Two-way ANOVA	Fig. legend	10, 8	mice from at least 3 litters	Fig. legend	error bars are mean+/- SEM	Fig. legend	p = 0.98	Fig. legend	F(1, 48) = 0.005	Fig. legend
+	supp -	4e	Two-way ANOVA	Fig. legend	10, 8	mice from at least 3 litters	Fig. legend	error bars are mean+/- SEM	Fig. legend	p = 1.0	Fig. legend	F(1, 48) = 0	Fig. legend
+	supp -	3h	One-way ANOVA	Fig. legend	5, 5, 5	mice from at least 3 litters	Fig. legend	error bars are mean+/- SEM	Fig. legend	P < 0.0001	Fig. legend	F(2, 14) = 99.5	Fig. legend
+	supp -	4f day1 &3	Two-way ANOVA	Fig. legend	6, 6, 6	mice from at least 3 litters	Fig. legend	error bars are mean+/- SEM	Fig. legend	p = 0.0004	Fig. legend	F(2, 15) = 13.5	Fig. legend
+	supp -	4f day7 &9	Two-way ANOVA	Fig. legend	6, 6	mice from at least 3 litters	Fig. legend	error bars are mean+/- SEM	Fig. legend	p = 0.011	Fig. legend	F(1, 10) = 9.76	Fig. legend
+	supp -	9b	Pearson correlation	Fig. legend	9 pairs	mice from at least 3 litters	Fig. legend	average	Fig. legend	p = 0.032	Fig. legend	r = -0.71	Fig. legend
+	supp -	10d	Unpaired t-test	Fig. legend	5, 7	mice from at least 3 litters	Fig. legend	error bars are mean+/- SEM	Fig. legend	p = 0.54	Fig. legend	t(10) = 0.634	Fig. legend
+	supp -	10e	Unpaired t-test	Fig. legend	5, 7	mice from at least 3 litters	Fig. legend	error bars are mean+/- SEM	Fig. legend	p = 0.77	Fig. legend	t(10) = 0.304	Fig. legend
+	supp -	10f	Unpaired t-test	Fig. legend	5, 5	mice from at least 3 litters	Fig. legend	error bars are mean+/- SEM	Fig. legend	p = 0.75	Fig. legend	t(8) = 0.335	Fig. legend
+	supp -	10g	Unpaired t-test	Fig. legend	5, 5	mice from at least 3 litters	Fig. legend	error bars are mean+/- SEM	Fig. legend	p = 0.64	Fig. legend	t(8) = 0.479	Fig. legend
+	supp -	7b	One-way ANOVA	Fig. legend	5, 8, 6	brain sections from 2 injections	Fig. legend	error bars are mean+/- SEM	Fig. legend	p = 0.0002	Fig. legend	F(2, 16) = 15.8	Fig. legend
+	supp -	7c	One-way ANOVA	Fig. legend	5, 8, 6	brain sections from 2 injections	Fig. legend	error bars are mean+/- SEM	Fig. legend	p = 0.0024	Fig. legend	F(2, 16) = 8.99	Fig. legend
+	supp -	7g	One-way ANOVA	Fig. legend	8, 8, 5	brain sections from 2 injections	Fig. legend	error bars are mean+/- SEM	Fig. legend	p < 0.0001	Fig. legend	F(2, 18) = 16.1	Fig. legend
+	supp -	7h	One-way ANOVA	Fig. legend	8, 8, 5	brain sections from 2 injections	Fig. legend	error bars are mean+/- SEM	Fig. legend	p < 0.0001	Fig. legend	F(2, 18) = 26.9	Fig. legend
+	supp -	7l	One-way ANOVA	Fig. legend	6, 5, 6	brain sections from 2 injections	Fig. legend	error bars are mean+/- SEM	Fig. legend	p < 0.0001	Fig. legend	F(2, 14) = 49.9	Fig. legend
+	supp -	7m	One-way ANOVA	Fig. legend	6, 5, 6	brain sections from 2 injections	Fig. legend	error bars are mean+/- SEM	Fig. legend	p = 0.0014	Fig. legend	F(2, 14) = 10.9	Fig. legend
+	fig 7i saline vs CCK activation		Unpaired t-test	Fig. legend	11, 9	mice from at least 3 litters	Fig. legend	error bars are mean+/- SEM	Fig. legend	p = 0.0093	Fig. legend	t(17) = 2.64	Fig. legend

+ -	fig 7i CCK contr ol vs activ ation	Unpaired t test	Fig. legend	9, 8	mice from at least 3 litters	Fig. legend	error bars are mean+/- SEM	Fig. legend	p = 0.014	Fig. legend	t(15) = 2.78	Fig. legend
+ -	supp 6b salin e vs CCK	Unpaired t test	Fig. legend	6, 6	brain sections from two injection experiments	Fig. legend	error bars are mean+/- SEM	Fig. legend	p = 0.0056	Fig. legend	t(10) = 3.37	Fig. legend
+ -	supp 6b salin e vs LiCl	Unpaired t test	Fig. legend	6, 6	brain sections from two injection experiments	Fig. legend	error bars are mean+/- SEM	Fig. legend	p = 0.028	Fig. legend	t(10) = 2.57	Fig. legend
+ -	supp 6c salin e vs CCK	Unpaired t test	Fig. legend	6, 6	brain sections from two injection experiments	Fig. legend	error bars are mean+/- SEM	Fig. legend	p = 0.0014	Fig. legend	t(10) = 4.14	Fig. legend
+ -	supp 6c salin e vs LiCl	Unpaired t test	Fig. legend	6, 6	brain sections from two injection experiments	Fig. legend	error bars are mean+/- SEM	Fig. legend	p = 0.011	Fig. legend	t(10) = 3.14	Fig. legend
+ -	supp 6d salin e vs CCK	Unpaired t test	Fig. legend	6, 6	brain sections from two injection experiments	Fig. legend	error bars are mean+/- SEM	Fig. legend	p = 0.30	Fig. legend	t(10) = 1.08	Fig. legend
+ -	supp 6d salin e vs LiCl	Unpaired t test	Fig. legend	6, 6	brain sections from two injection experiments	Fig. legend	error bars are mean+/- SEM	Fig. legend	p = 0.024	Fig. legend	t(10) = 2.66	Fig. legend
+ -	supp 7d	One-way ANOVA	Fig. legend	5, 8, 6	brain sections from two injection experiments	Fig. legend	error bars are mean+/- SEM	Fig. legend	p = 0.049	Fig. legend	F(2, 16) = 3.64	Fig. legend
+ -	supp 7i	One-way ANOVA	Fig. legend	8, 8, 5	brain sections from two injection experiments	Fig. legend	error bars are mean+/- SEM	Fig. legend	p = 0.0011	Fig. legend	F(2, 18) = 3.90	Fig. legend
+ -	supp 7n	One-way ANOVA	Fig. legend	6, 5, 6	brain sections from two injection experiments	Fig. legend	error bars are mean+/- SEM	Fig. legend	p = 0.045	Fig. legend	F(2, 14) = 10.9	Fig. legend

► Representative figures

1. Are any representative images shown (including Western blots and immunohistochemistry/staining) in the paper?

If so, what figure(s)?

Fig. 1a-c
Fig. 6k
Supplementary Fig. 2a
Supplementary Fig. 5a
Supplementary Fig. 6a-c
Supplementary Fig. 7e, j, o
Supplementary Fig. 8a
Supplementary Fig. 8h
Supplementary Fig. 9a
Supplementary Fig. 10a, b

2. For each representative image, is there a clear statement of how many times this experiment was successfully repeated and a discussion of any limitations in repeatability?

If so, where is this reported (section, paragraph #)?

Each representative figure is one of the figures that used for statistical analysis, the number of repeating is the n described in individual figure legends.

► Statistics and general methods

1. Is there a justification of the sample size?

If so, how was it justified?

Where (section, paragraph #)?

Even if no sample size calculation was performed, authors should report why the sample size is adequate to measure their effect size.

We chose the sample size based on literatures in the field.

2. Are statistical tests justified as appropriate for every figure?

Where (section, paragraph #)?

The statistics were used based on the properties of the data points, and described in individual figure legends

- a. If there is a section summarizing the statistical methods in the methods, is the statistical test for each experiment clearly defined?

Yes, we summarized in the final paragraph of the methods. each statistical test is defined in each fig legend.

- b. Do the data meet the assumptions of the specific statistical test you chose (e.g. normality for a parametric test)?

Where is this described (section, paragraph #)?

Based on previous literature, we assumed the data points have a normal distribution and used ANOVA or t-test.

- c. Is there any estimate of variance within each group of data? Is the variance similar between groups that are being statistically compared?

Where is this described (section, paragraph #)?

Yes. Either standard error of mean or Box-and-whisker plot was used to analyze our data. They were described in individual figure legends.

- d. Are tests specified as one- or two-sided?

Yes, in individual fig legend.

- e. Are there adjustments for multiple comparisons?

Yes, in indicated in individual fig legend

3. Are criteria for excluding data points reported?

Was this criterion established prior to data collection?

Where is this described (section, paragraph #)?

N/A

4. Define the method of randomization used to assign subjects (or samples) to the experimental groups and to collect and process data.

If no randomization was used, state so.

Where does this appear (section, paragraph #)?

N/A

5. Is a statement of the extent to which investigator knew the group allocation during the experiment and in assessing outcome included?
If no blinding was done, state so.
Where (section, paragraph #)?
- Yes, described in 5th paragraph of methods "Behavioral tests"
6. For experiments in live vertebrates, is a statement of compliance with ethical guidelines/regulations included?
Where (section, paragraph #)?
- Yes, described in methods, 1st paragraph
7. Is the species of the animals used reported?
Where (section, paragraph #)?
- Yes, described in methods "Mice"
8. Is the strain of the animals (including background strains of KO/transgenic animals used) reported?
Where (section, paragraph #)?
- Yes, described in methods "Mice"
9. Is the sex of the animals/subjects used reported?
Where (section, paragraph #)?
- Yes, described in methods "Mice"
10. Is the age of the animals/subjects reported?
Where (section, paragraph #)?
- Yes, described in methods "Animal surgery"
11. For animals housed in a vivarium, is the light/dark cycle reported?
Where (section, paragraph #)?
- Yes, described in methods "Mice"
12. For animals housed in a vivarium, is the housing group (i.e. number of animals per cage) reported?
Where (section, paragraph #)?
- Yes, described in methods "Animal surgery"
13. For behavioral experiments, is the time of day reported (e.g. light or dark cycle)?
Where (section, paragraph #)?
- Yes, described in methods "Behavioral test"
14. Is the previous history of the animals/subjects (e.g. prior drug administration, surgery, behavioral testing) reported?
Where (section, paragraph #)?
- N/A
- a. If multiple behavioral tests were conducted in the same group of animals, is this reported?
Where (section, paragraph #)?
- N/A
15. If any animals/subjects were excluded from analysis, is this reported?
Where (section, paragraph #)?
- N/A

a. How were the criteria for exclusion defined?

N/A

Where is this described (section, paragraph #)?

b. Specify reasons for any discrepancy between the number of animals at the beginning and end of the study.

N/A

Where is this described (section, paragraph #)?

▶ Reagents

1. Have antibodies been validated for use in the system under study (assay and species)?

Yes

a. Is antibody catalog number given?

Yes, in methods "Immunohistochemistry and histology"

Where does this appear (section, paragraph #)?

b. Where were the validation data reported (citation, supplementary information, Antibodypedia)?

Reported from the companies that provide the antibodies

Where does this appear (section, paragraph #)?

2. If cell lines were used to reflect the properties of a particular tissue or disease state, is their source identified?

N/A

Where (section, paragraph #)?

a. Were they recently authenticated?

N/A

Where is this information reported (section, paragraph #)?

▶ Data deposition

Data deposition in a public repository is mandatory for:

- Protein, DNA and RNA sequences
- Macromolecular structures
- Crystallographic data for small molecules
- Microarray data

Deposition is strongly recommended for many other datasets for which structured public repositories exist; more details on our data policy are available [here](#). We encourage the provision of other source data in supplementary information or in unstructured repositories such as [Figshare](#) and [Dryad](#).

1. Are accession codes for deposit dates provided?

N/A

Where (section, paragraph #)?

▶ Computer code/software

Any custom algorithm/software that is central to the methods must be supplied by the authors in a usable and readable form for readers at the time of publication. However, referees may ask for this information at any time during the review process.

1. Identify all custom software or scripts that were required to conduct the study and where in the procedures each was used.

N/A

2. Is computer source code/software provided with the paper or deposited in a public repository? Indicate in what form this is provided or how it can be obtained.

N/A

▶ Human subjects

1. Which IRB approved the protocol?

Where is this stated (section, paragraph #)?

N/A

2. Is demographic information on all subjects provided?

Where (section, paragraph #)?

N/A

3. Is the number of human subjects, their age and sex clearly defined?

Where (section, paragraph #)?

N/A

4. Are the inclusion and exclusion criteria (if any) clearly specified?

Where (section, paragraph #)?

N/A

5. How well were the groups matched?

Where is this information described (section, paragraph #)?

N/A

6. Is a statement included confirming that informed consent was obtained from all subjects?

Where (section, paragraph #)?

N/A

7. For publication of patient photos, is a statement included confirming that consent to publish was obtained?

Where (section, paragraph #)?

N/A

► fMRI studies

For papers reporting functional imaging (fMRI) results please ensure that these minimal reporting guidelines are met and that all this information is clearly provided in the methods:

1. Were any subjects scanned but then rejected for the analysis after the data was collected?
 - a. If yes, is the number rejected and reasons for rejection described?
Where (section, paragraph #)?
2. Is the number of blocks, trials or experimental units per session and/or subjects specified?

Where (section, paragraph #)?
3. Is the length of each trial and interval between trials specified?
4. Is a blocked, event-related, or mixed design being used? If applicable, please specify the block length or how the event-related or mixed design was optimized.
5. Is the task design clearly described?
Where (section, paragraph #)?
6. How was behavioral performance measured?
7. Is an ANOVA or factorial design being used?
8. For data acquisition, is a whole brain scan used?
If not, state area of acquisition.
 - a. How was this region determined?
9. Is the field strength (in Tesla) of the MRI system stated?
 - a. Is the pulse sequence type (gradient/spin echo, EPI/spiral) stated?
 - b. Are the field-of-view, matrix size, slice thickness, and TE/TR/flip angle clearly stated?
10. Are the software and specific parameters (model/functions, smoothing kernel size if applicable, etc.) used for data processing and pre-processing clearly stated?

11. Is the coordinate space for the anatomical/functional imaging data clearly defined as subject/native space or standardized stereotaxic space, e.g., original Talairach, MNI305, ICBM152, etc? Where (section, paragraph #)?

N/A

12. If there was data normalization/standardization to a specific space template, are the type of transformation (linear vs. nonlinear) used and image types being transformed clearly described? Where (section, paragraph #)?

N/A

13. How were anatomical locations determined, e.g., via an automated labeling algorithm (AAL), standardized coordinate database (Talairach daemon), probabilistic atlases, etc.?

N/A

14. Were any additional regressors (behavioral covariates, motion etc) used?

N/A

15. Is the contrast construction clearly defined?

N/A

16. Is a mixed/random effects or fixed inference used?

N/A

a. If fixed effects inference used, is this justified?

N/A

17. Were repeated measures used (multiple measurements per subject)?

N/A

a. If so, are the method to account for within subject correlation and the assumptions made about variance clearly stated?

N/A

18. If the threshold used for inference and visualization in figures varies, is this clearly stated?

N/A

19. Are statistical inferences corrected for multiple comparisons?

N/A

a. If not, is this labeled as uncorrected?

N/A

20. Are the results based on an ROI (region of interest) analysis?

N/A

a. If so, is the rationale clearly described?

N/A

b. How were the ROI's defined (functional vs anatomical localization)?

N/A

21. Is there correction for multiple comparisons within each voxel?

N/A

22. For cluster-wise significance, is the cluster-defining threshold and the corrected significance level defined?

N/A

▶ Additional comments

Additional Comments

