

SUPPLEMENTAL INFORMATION

Figure S1. HFD pair-feeding does not impact glucose or food intake but deficiency of microglial inflammatory signaling tends to increase gluconeogenesis, related to Figure 2. (A)

Blood glucose and (B) food intake in IKK β -MGKO mice fed *ad libitum* (IKK β -MGKO AL) vs. pair-fed control (Ctl PF). Measurements were obtained 4 hours into the dark cycle. N = 7/group.

(C) Pyruvate tolerance test (2 g/kg pyruvate, i.p.) and (D) total glucose AUC at 11 weeks HFD in IKK β -MGKO mice fed *ad libitum* vs. pair-fed control.

Figure S2. Validation of TNF antibody (abcam, ab1793), related to Figure 4. Representative images of hypothalamic sections showing Iba1 (green) and TNF (red) staining in wild-type (top) and TNF KO (bottom) mice treated with LPS (5 mg/kg) for 6 hours. Scale bars = 20 μ m.

Figure S3. Pharmacological inhibition of microglia worsens insulin resistance during DIO in rats, related to Figure 5. Male Wistar rats (N = 5/group) were fed HFD for 4 weeks and treated with minocycline (10 μ g, i.c.v.) or saline vehicle administered 3 times over 2d. (A) Plasma insulin levels at t = 0 of ITT. (B) ITT (1.5 U/kg insulin, i.p.) performed 1h after the third dose of minocycline or vehicle. (C) Total glucose AOC for the ITT. Values are mean \pm SEM. 2-way ANOVA with Bonferroni post-hoc test and Student's *t*-test, * p < 0.05, ** p < 0.01.

Figure S4. Chemogenetic microglial activation in hM3D mice is associated with increased pro-inflammatory signaling, related to Figure 5, Figure 6, and Figure 7. Cx3cr1^{CreER/+}::LSL-HA-hM3Dq-citrine (hM3D) mice were generated as an inducible model of *in vivo* microglial activation. (A) Representative hypothalamic sections immunostained for the microglial marker

(Iba1, green) and hM3D DREADD receptor (anti-HA tag, red) in tamoxifen-treated control and hM3D mice. Scale bars = 100 μ m. (B) Quantification of hM3D transgene expression by qRT-PCR in CNS and peripheral tissues in hM3D mice normalized to control (Cx3cr1^{CreER/+}) levels. N = 3-4/group. Student's *t*-test, ** $p < 0.01$. (C) Fura Red was used to detect changes in intracellular calcium. Isolated hM3D primary microglia were treated with CNO (0.01 μ M or 0.1 μ M) or saline control over 150 minutes. Fura Red fluorescence of the microglial cells was measured using a plate reader and plotted normalized to 0 minute baseline. Mean traces shown for clarity. (D) 30 minute AUC bins of data in C. N = 6-7 replicates/treatment. Two-way ANOVA with Bonferroni post-hoc test, * $p < 0.05$, ** $p < 0.01$, *** $p < 0.001$, **** $p < 0.0001$. (E) Representative hypothalamic sections from hM3D mice immunostained for the microglial marker (Iba1, green), hM3D DREADD receptor (anti-HA tag, red), and DAPI nuclear stain in hM3D mice given CNO (1 mg/kg, i.p.) or saline and perfused 1 hour later. Scale bars = 100 μ m. (F-G) Quantification of Iba1⁺ microglia by (F) total number and (G) cellular size after CNO treatment. N = 6-12/group. Student's *t*-test, ** $p < 0.01$. (H) Cytokine expression in FACS-isolated microglia from hM3D normalized to control levels in mice treated with CNO (1 mg/kg, i.p.). N = 6-7/genotype. Student's *t*-test, * $p < 0.05$, ** $p < 0.01$, *** $p < 0.001$. All values are mean \pm SEM.

Figure S5. Acute CNO-mediated activation of microglial hM3D does not induce sickness behavior, related to Figure 5. (A) Dark cycle ambulatory activity was measured after saline or CNO injection (1 mg/kg, i.p.) in hM3D mice. Pretreatment line represents average of both groups of hM3D mice before injection (indicated by arrow). Dashed lines represent \pm SEM. (B) 30 min AUC bins of activity for 240 min after injection. N = 13-19/group. (C) Food intake and

(D) body weight were measured 24 hours after saline or CNO injection (0.3mg/kg or 1mg/kg, i.p.) using a within subjects crossover design. N = 14/group.

Figure S6. Subthreshold peripheral CNO administration does not improve glucose tolerance in hM3D mice, related to Figure 5. (A) GTT (2 g/kg glucose, i.p.) and (B) AUC in CD-fed hM3D and Ctl mice administered systemic CNO at the same dose used for i.c.v. experiments (1 µg, i.p., 2 hours prior). N = 8/group. All values are mean ± SEM.

Figure S7. Chemogenetic activation of microglia in a Tmem119 Cre-based DREADD model improves glucose tolerance, related to Figure 5. (A-D) GTT and AUC of CD-fed Tmem-Gq mice and littermate controls treated with (A-B) saline or (C-D) systemic CNO (10 mg/kg, i.p. 2 hr prior; 2 g/kg glucose, i.p.). N = 8-9/group. Student's *t*-test, *** $p < 0.001$.

Figure S8. Chemogenetic activation of microglia improves glucose tolerance independently of central IL-6, IL-1, or CCR5 signaling and increases POMC neuron glucose sensing, related to Figure 6. (A-F) CD-fed control and hM3D mice were injected with CNO (1 mg/kg, i.p.) and i.c.v. treatment listed below (indicated in figures by arrow labeled i.c.v.) 2 hours prior to all GTTs (2 g/kg glucose, i.p.). (A) GTT and (B) AUCs of mice administered IL-6ab (150 ng, i.c.v.) or vehicle. (C) GTT and (D) AUC of mice administered IL-1RA (100 ng, i.c.v.) or vehicle. (E) GTT and (F) AUC of mice administered CCR5 antagonist maraviroc (CCR5ant) (500 ng, i.c.v.) or vehicle. Two-way ANOVA with Šidák correction for multiple comparisons. (G) Representative images from the ARC of control and hM3D mice 2 hours after CNO (5 mg/kg, i.p.) and 1 hour after saline or glucose (2 g/kg, i.p.) stained for β-endorphin (green) and c-Fos (red).

Scale bars = 50 μm . (H) Quantification of c-Fos⁺ cells as a percent of β -endorphin cells from G, N = 7-8 mice/group, 5 sections/mouse. Two-way ANOVA with post-hoc comparison. * $p < 0.05$, ** $p < 0.01$, *** $p < 0.001$. All values are mean \pm SEM.

Figure S9. Microglial chemogenetic activation increases first-phase insulin release, which is not prevented by sympathetic agonist administration, related to Figure 7. (A-D) Acute-phase glucose-stimulated insulin secretion in CD-fed control and hM3D mice treated with systemic CNO (5 mg/kg, i.p., 2 hours prior; 2 g/kg glucose, i.p.) at (A-B) 4 min and (C-D) 8 min post-glucose injection, N = 9-11/group. Two-way ANOVA with Šidák correction for multiple comparisons. (E-G) GTT (2 g/kg glucose, i.p.) performed in CD-fed control and hM3D mice 45 min after administration of (E) saline or (F) clonidine (0.01 mg/kg, i.p.) and 20 min after administration of low-dose CNO (0.5 mg/kg, i.p.). (G) AUC quantification of E and F, N = 7-8/group. Two-way ANOVA with Sidak correction for multiple comparisons. * $p < 0.05$, ** $p < 0.01$, *** $p < 0.001$. All values are mean \pm SEM.

Table S1. Genotyping and real-time qPCR primers, related to STAR Methods.

REAGENT or RESOURCE	SOURCE	IDENTIFIER
Real-time qPCR Primers		
Primers for 18S; <i>Forward:</i> CGGACAGGATTGACAGATTG <i>Reverse:</i> CAAATCGCTCCAATAA		N/A
Primers for mouse TNF α ; <i>Forward:</i> CATCTTCTCAAACTCGAGTGACAA <i>Reverse:</i> TGGGAGTAGATAAGGTACAGCCC		N/A
Primers for mouse IL1 β ; <i>Forward:</i> TACAAGGAGAGACAAGCAACGACA <i>Reverse:</i> GATCCACACTCTCCAGCTGCA		N/A
Primers for mouse IL6; <i>Forward:</i> GTGGCTAAGGACCAAGACCA <i>Reverse:</i> GGTTTGCCGAGTAGACCTCA		N/A

Primers for mouse MIP1 α ; <i>Forward:</i> CTGCCCTTGCTGTTCTTCTC <i>Reverse:</i> GATGAATTGGCGTGGAATCT		N/A
Primers for mouse MIP1 β ; <i>Forward:</i> CTGCCCTCTCTCCTCTTG <i>Reverse:</i> AGGAAGTGGGAGGGTCAGAG		N/A
Primers for mouse MCP1; <i>Forward:</i> CTTCTGGGCCTGCTGTTCA <i>Reverse:</i> CAAGCCTACTCATGGGCTTC		N/A
Primers for mouse Slc2a1; <i>Forward:</i> CCCCGTCCTGCTGCTATTG <i>Reverse:</i> GCACCGTGAAGATGATGAAGAC		N/A
Primers for mouse UCP2; <i>Forward:</i> AAGCAGTTCTACACCAAGGGC <i>Reverse:</i> TCTCGTGCAATGGTCTTGTAG		N/A
Primers for mouse GCK; <i>Forward:</i> CAAGCTGCACCCGAGCTT <i>Reverse:</i> TGATTGATGAAGGTGATTTCG		N/A
Primers for mouse G6pc; <i>Forward:</i> TCAACCTCGTCTTCAAGTGGATT <i>Reverse:</i> CTGCTTTATTATAGGCACGGAGCT		N/A
Primers for mouse Pck1; <i>Forward:</i> GGC GGAGCATATGCTGATCC <i>Reverse:</i> CCACAGGCACTAGGGAAGGC		N/A
Primers for mouse Gq DREADD; <i>Forward:</i> TCTGGCAAGTGGTCTTCATC <i>Reverse:</i> TTACCAGGATGTTGCCGATG		N/A
Genotyping Primers		
Primers for Ikbkb floxed allele; <i>Forward:</i> CCTTGTCCCTATAGAAGCACAAC <i>Reverse:</i> GTCATTTCCACAGCCCTGTGA		N/A
Primers for CreER allele; <i>Common</i> <i>Forward:</i> AAGACTCACGTGGACCTGCT <i>Mutant</i> <i>Reverse:</i> CGGTTATTCAACTTGCACCA <i>Wild Type Reverse:</i> AGGATGTTGACTTCCGAGTTG		N/A
Primers for hM3D line; <i>Transgene</i> <i>Forward:</i> TCTGGCAAGTGGTCTTCATC <i>Transgene</i> <i>Reverse:</i> TTACCAGGATGTTGCCGATG <i>Int. Pos. Ctrl Forward:</i> CACGTGGGCTCCAGCATT <i>Int. Pos. Ctrl Reverse:</i> TCACCAGTCATTTCTGCCTTTG		N/A

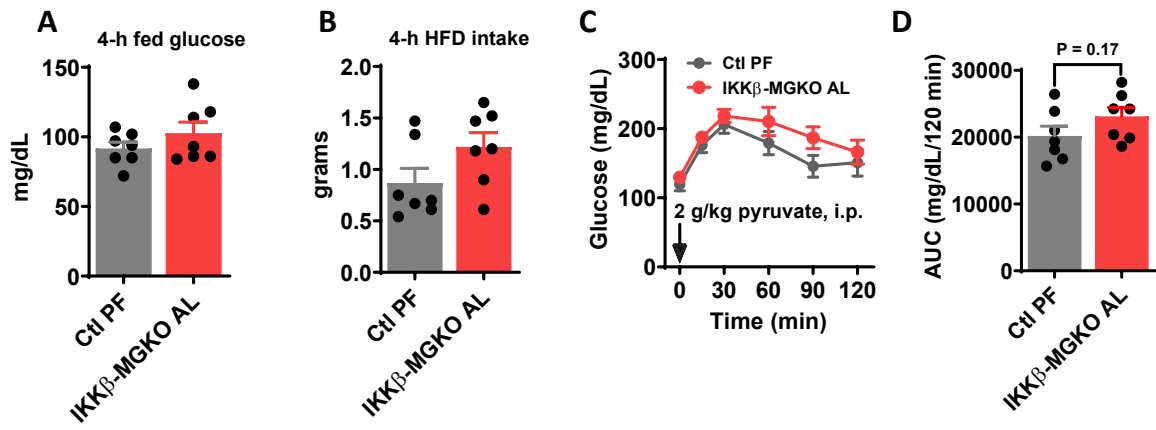


Figure S1

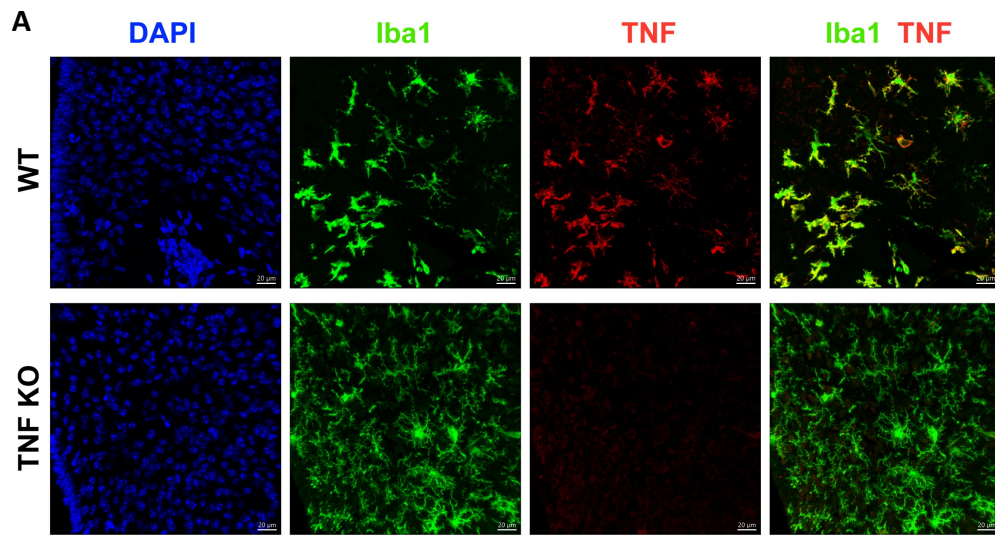


Figure S2

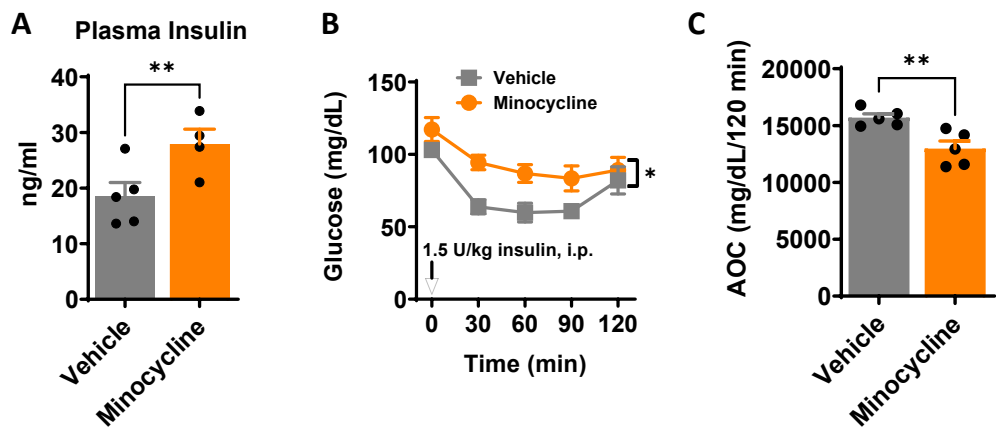


Figure S3

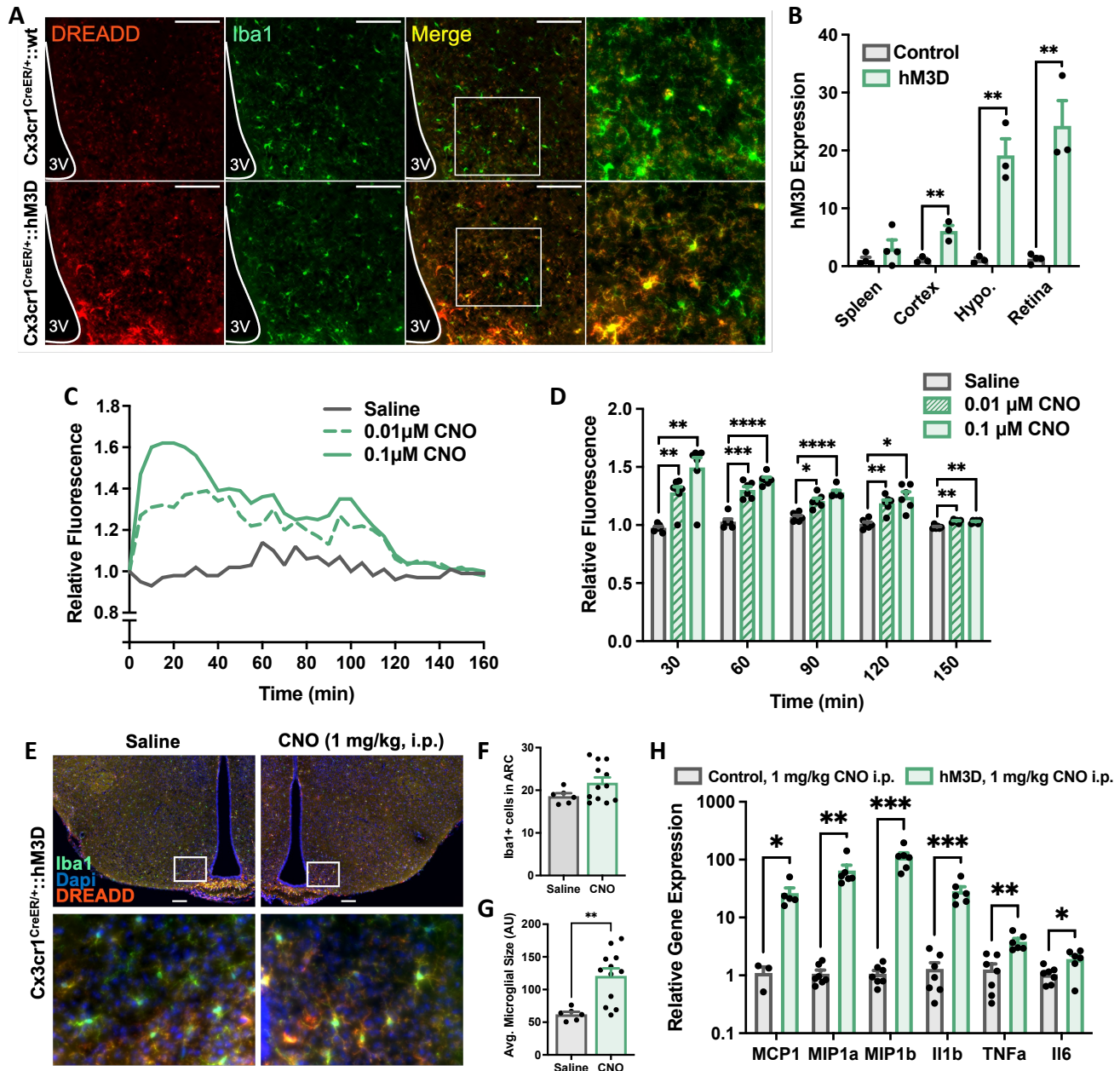


Figure S4

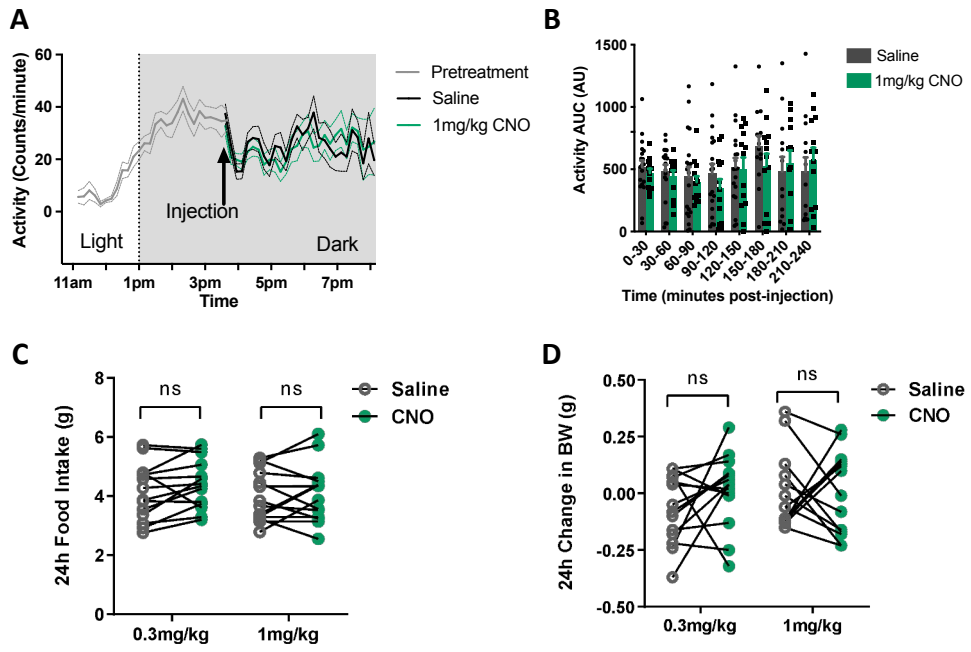


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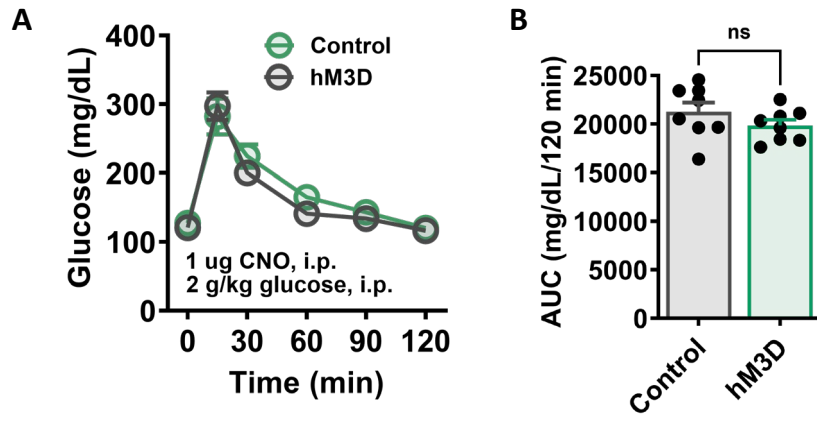


Figure S6

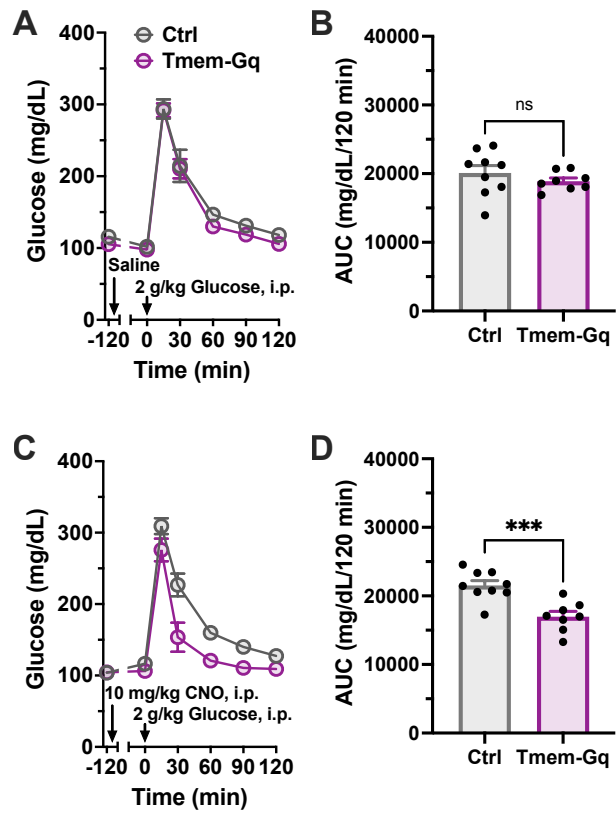


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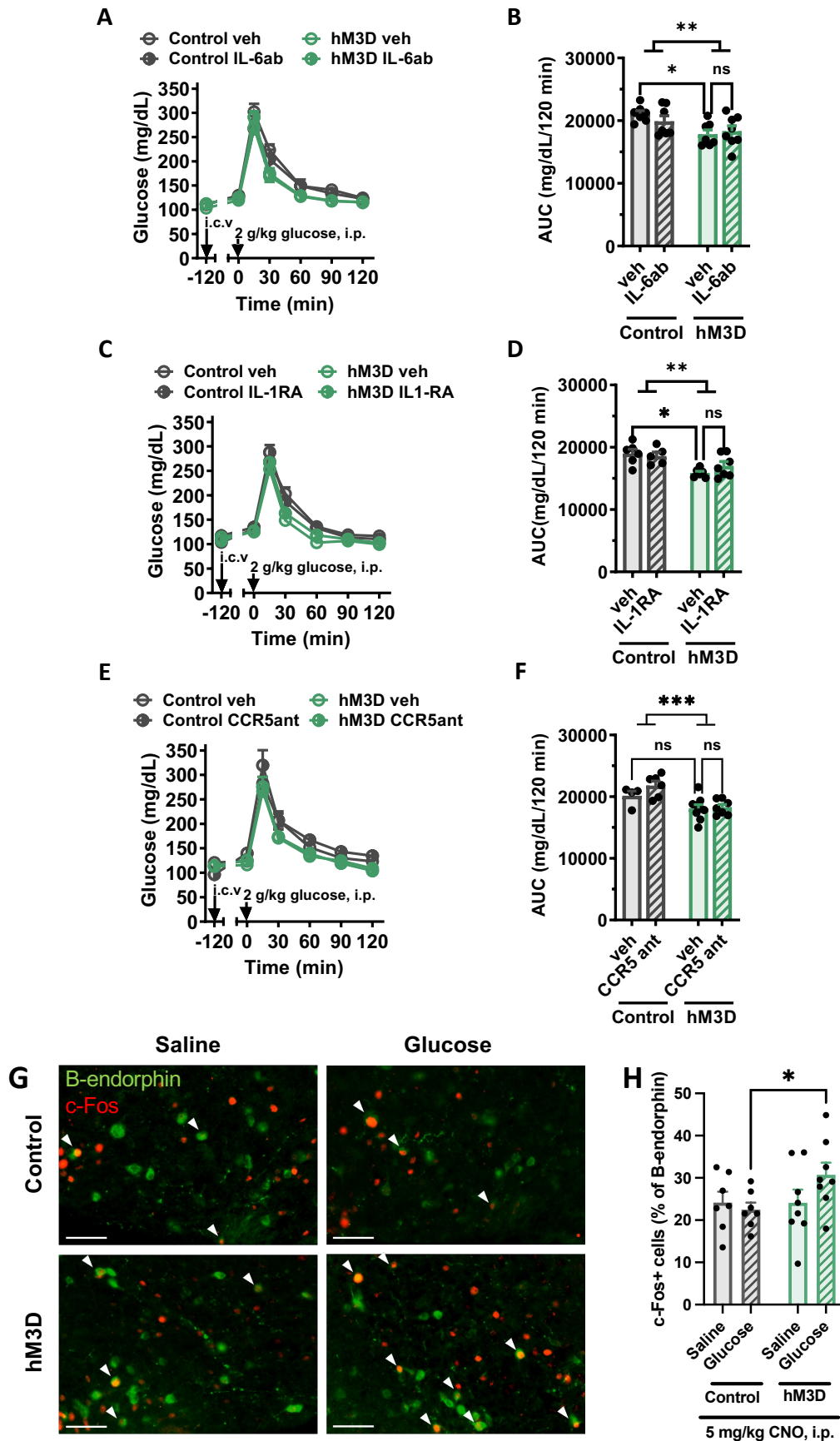


Figure S8

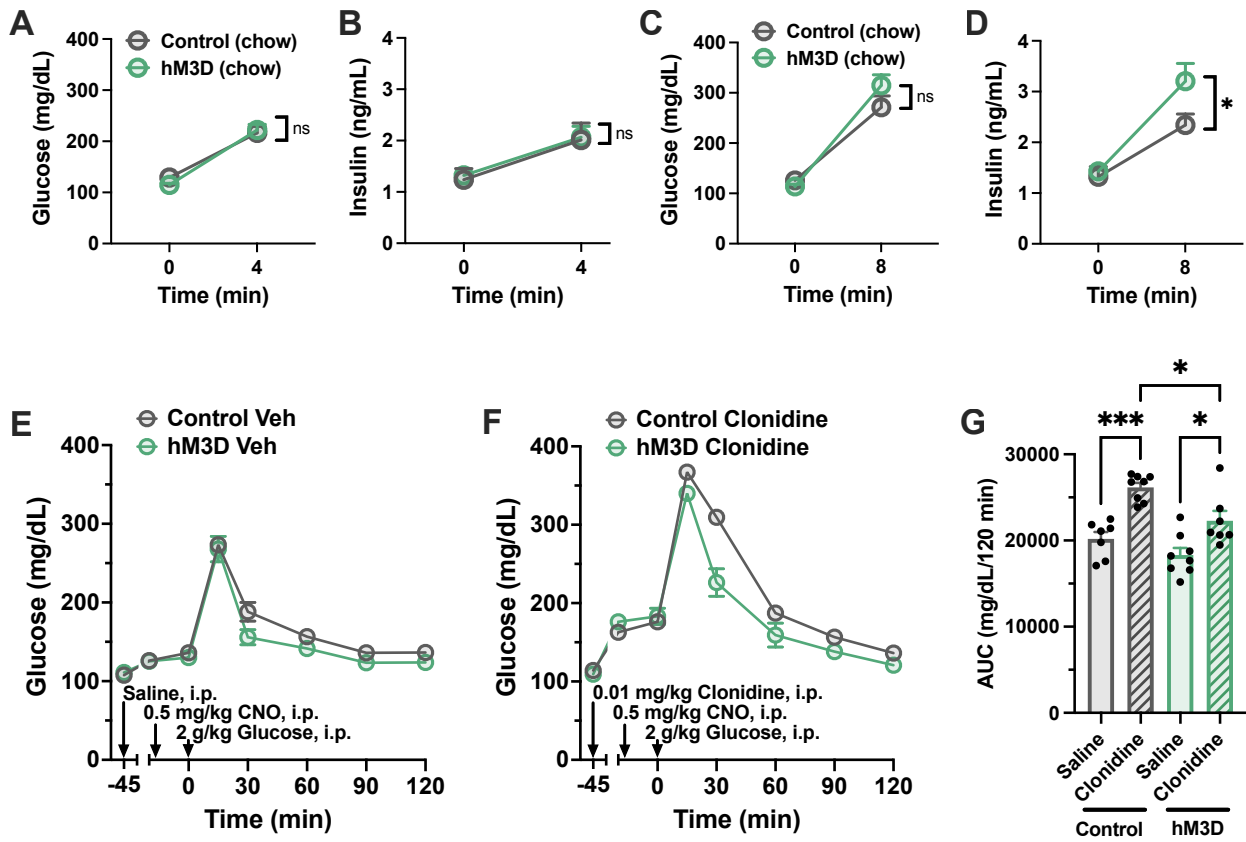


Figure S9