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Supplemental information

Central FGF21 production regulates

memory but not peripheral metabolism

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Supplemental Figures

Figure S1, related to Figure 1, FGF21 is expressed in the retrosplenial cortex but not the hypothalamus.

(A-B) Relative Fgf21(A) and Cre (B) mRNA levels from the liver of 10-12 week old male Fgf21-P2A-CRE mice and CRE negative littermate *ad libitum* fed or following a 24-hour fast (n=5-6/group). Statistical analyses were conducted using a two-way ANOVA with multiple comparisons (* = p < 0.05).

(C) Representative fluorescence imaging for tdTomato-positive cells in the pancreas of 12-weekold male Fgf21-P2A-CRE;Ai-tdTomato mice. Scale bar = 500 µm.

(D) Representative fluorescence imaging for tdTomato-positive cells in the medial hypothalamus of female Fgf21-P2A-CRE;Ai14-tdTomato and CRE negative control mice following a 24-hour fast. Scale bars = 250 µm.

(E) RNAscope *in situ* hybridization for *Fgf21* and *tdTomato* mRNA in retrosplenial cortex (RSC) and hypothalamus from *Fgf21*-P2A-CRE;Ai-tdTomato mice. 3V = third ventricle. Scale bars = 75 μ m.

(F) Representative immunofluorescence imaging for FGF21 and vimentin in the median eminence of male wild-type mice using primary (Rabbit anti-FGF21; Chicken anti-vimentin) and secondary (FGF21, Goat anti-Rabbit IgG (H+L) Cross-Absorbed Secondary Antibody Texas Red; Vimentin, Goat anti-Chicken IgY (H+L) Secondary Antibody Alexa Fluor 488) antibodies or FGF21 secondary antibody only.

(G) In silico single cell analysis of Fgf21 expression in the hypothalamus in the indicated cell types.

(H) In silico single cell analysis of Fgf21-expressing cells in the thalamus based on the indicated marker genes.

Values are mean \pm SEM.

Figure S2, related to Figure 2, Neuron-derived FGF21 does not regulate energy homeostasis.

(A) Representative fluorescence imaging for GFP-positive cells in 12-week-old $FGF21^{fl/fl}$; PHP.eB-eGFP-CRE mice. Scale bar = 1 mm.

(B) Daily measurement of sucrose intake during two-bottle choice of 10% sucrose versus water in male FGF21^{fl/fl} mice injected with AAV-TBG-Con (null) or AAV-TBG-CRE virus (n=10/group). Statistical analyses were conducted using a two-way ANOVA with multiple comparisons (* = p < 0.05).

(C-D) Plasma glucose (C) and plasma non-esterified fatty acid (NEFA) (D) levels in *ad libitum* fed and 24 hour fasted male FGF21^{fl/fl} mice injected with AAV-TBG-Con (null) or AAV-TBG-CRE virus (n=7-10/group). Statistical analyses were conducted using a two-way ANOVA with multiple comparisons (* = p < 0.05).

(E-F) Linear modeling of energy expenditure of 12-14 week old male FGF21^{fl/fl};PHP.eB-GFP and FGF21^{fl/fl};PHP.eB-CRE mice (n=7-9/group) during the light (E) or dark (F) cycles.

Values are mean \pm SEM.

Figure S3, related to Figure 3, Role of FGF21 in learning and memory.

(A-B) Average of the four trials for latency to the escape chamber during training (A) and primary error to the escape chamber during training (B) in male FGF21^{fl/fl} mice injected with PHP.eB-GFP

or PHP.eB-CRE virus (n=7-9/group). Statistical analyses were conducted using a two-way ANOVA with multiple comparisons (* = p < 0.05).

(C-D) Average of the four trials for latency to the escape chamber during training (C) and primary error to the escape chamber during training (D) in male FGF21^{fl/fl} mice injected with AAV-TBG-Con (null) or AAV-TBG-CRE virus (n=15/group). Statistical analyses were conducted using a two-way ANOVA with multiple comparisons (* = p < 0.05).

(E) Representative fluorescent image of viral targeting of AAV-CMV-CRE to the retrosplenial cortex (RSC) of FGF21^{fl/fl} mice. Scale bar = 1 mm.

(F-G) Average of the four trials for latency to the escape chamber during training (F) and primary error to the escape chamber during training (G) in male FGF21^{fl/fl} mice injected with AAV-CMV-CRE and AAV-CMV-GFP virus in the RSC (n=15/group). Statistical analyses were conducted using a two-way ANOVA with multiple comparisons (* = p < 0.05).

(H-J) Escape latency on test trials 24 hours (H), 7 days (I), and 14 days (J) after the training of male FGF21^{fl/fl} mice which received stereotaxic injection of AAV-CMV-CRE and AAV-CMV-GFP virus compared to wild-type mice which received AAV-CMV-CRE virus into the retrosplenial cortex (RSC) (n=9-15group). Statistical analyses were conducted using a two-tailed t test (* = p < 0.05).

(K) Representative fluorescent image of viral targeting of AAV-hSyn-CRE-EGFP to the retrosplenial cortex (RSC) of FGF21^{fl/fl} mice. Scale bar = 1 mm.

(L-M) Average of the four trials for latency to the escape chamber during training (L) and primary error to the escape chamber during training (M) in male FGF21^{fl/fl} mice injected with AAV-hSyn-CRE and control virus in the RSC (n=15-16/group). Statistical analyses were conducted using a two-way ANOVA with multiple comparisons (* = p < 0.05).

(N-O) Average of the four trials for latency to the escape chamber during training (N) and primary error to the escape chamber during training (O) in wild-type male mice administered vehicle or FGF21 (1 mg/kg) each day of training (n=14-15group). Statistical analyses were conducted using a two-way ANOVA with multiple comparisons (* = p < 0.05).

(P-R) Escape latency on test trials 24 hours (P), 7 days (Q), and 14 days (R) after the training of 11-13 week old female C57BL/6J mice treated with FGF21 (1 mg/kg) or saline for 4 days via i.p. injection (n=13-15/group). Statistical analyses were conducted using a two-tailed t test (* = p < 0.05).

Values are mean \pm SEM.

Figure S4, related to Figure 3, FGF21 administration consolidates spatial and fear memory.

Total freezing time of cued-dependent fear (A) and contextual (B) memory of 10-12 week old male C57BL/6 mice administered FGF21 (1mg/kg) or saline treatment via i.p. injection (n=10/group). Values are mean \pm SEM.. Statistical analyses were conducted using a two-tailed t test (* = p < 0.05).

Video S1, related to Figure 1, Comprehensive evaluation of FGF21-expressing cells in the CNS.

Light sheet imaging of a brain from *Fgf21*-CRE;Ai14-tdTomato mice processed by DISCO clearing.





Fig. S2





Figure S4