

Supplementary Figure 1. Histology of pituitary gland, adrenal gland, and testis.

Sections from the indicated tissues from WT and *FoxO1 KO*^{SF1} mice were analyzed using hematoxylin and eosin (H&E) staining. Ant, anterior lobe. Int, intermediate lobe. Post, posterior lobe. C, cortex. M, medulla. I, interstitium. S, spermatozoa. ST, seminiferous tubules. Scale bar = 200um.



Supplementary Figure 2. Effect of acute high fat diet on metabolic parameters in WT and *FoxO1 KO^{SF1}* mice.

Temporal changes of (a) heat generation, (b) CO_2 production, and (g) RER. Changes in (d) heat generation, (e) CO_2 production, and (i) RER between 2 days before and after HFD. (c) Cumulative food intake and (f) daily food intake between genotypes. (h) Ambulatory movement and (j) average in total movement. Numbers of animals examined are expressed in parenthesis in each graph. The data are expressed as either average (a-c, g, h) or mean±SEM (*P<0.05, Student's t-test). cnt, beam breaks counts.

Effect of FoxO1 KO in the VMH on biochemical parameters				
01 KO ^{SF1} Age (w	vks) Bleeding Time			
2±8.15 (5) 6-7	12:00-12:30			
2±9.03 (4) 5	08:00-08:30			
±30.47 (5) 5	08:00-08:30			
±22.72 (7) 13-16	6 14:00-15:30			
±26.60 (6) 13-16	6 14:00-15:30			
2±0.36 (5) 13-16	6 14:00-15:30			
4±0.07 (8) 7-9	14:00-15:30			
41±0.1 (6) 7-10				
	14:00-15:30			
67±1.2 (8) 7-9	14:00-15:30 14:00-15:30			
	±30.47 (5) 5 ±22.72 (7) 13-16 ±26.60 (6) 13-16 ±20.36 (5) 13-16 4±0.07 (8) 7-9			

Supplementary Table 1 Effect of FoxO1 KO in the VMH on biochemical parameters

The data were expressed as mean±SEM and numbers of experiemental animals were indicated in parenthesis. No statistical significance was detected between genotypes.

Supplementary Table 2 Body length in various food conditions

	WT	FoxO1 KO ^{SF1}	Age (wks)
Male, normal chow (cm)	9.36±0.05 (10)	9.38±0.05 (9)	10
Female, normal chow (cm)	9.34±0.05 (10)	9.4±0.04 (8)	17-18
Male, high fat (cm)	9.67±0.09 (8)	9.85±0.12 (6)	11-12
Female, high fat (cm)	9.02±0.21 (6)	9.15±0.21 (4)	12-13

The data were expressed as mean±SEM and numbers of experiemental animals were indicated in parenthesis. No statistical significance was detected between genotypes.