

NPY and MC4R Signaling Regulate Thyroid Hormone Levels during Fasting through Both Central and Peripheral Pathways

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SUMMARY

Fasting-induced suppression of the hypothalamic-pituitary-thyroid (HPT) axis is an adaptive response to decrease energy expenditure during food deprivation. Previous studies demonstrate that leptin communicates nutritional status to the HPT axis through thyrotropin-releasing hormone (TRH) in the paraventricular nucleus (PVN) of the hypothalamus. Leptin targets TRH neurons either directly or indirectly via the arcuate nucleus through pro-opiomelanocortin (POMC) and agouti-related peptide/neuropeptide Y (AgRP/NPY) neurons. To evaluate the role of these pathways *in vivo*, we developed double knockout mice that lack both the melanocortin 4 receptor (MC4R) and NPY. We show that NPY is required for fasting-induced suppression of *Trh* expression in the PVN. However, both MC4R and NPY are required for activation of hepatic pathways that metabolize T₄ during the fasting response. Thus, these signaling pathways play a key role in the communication of fasting signals to reduce thyroid hormone levels both centrally and through a peripheral hepatic circuit.

INTRODUCTION

Circulating thyroid hormone (TH) levels are critical regulators of energy expenditure in rodents and man (Barker, 1951; Du Bois, 1936). Thus, their acute regulation in periods of nutritional stress or illness is an important adaptive mechanism for survival. In humans, long-term caloric restriction leads to a fall in TH levels, which is secondary to a fall in leptin levels (Rosenbaum et al., 2005; Rosenbaum et al., 2002). Similar physiology exists in rodents during an acute fast where a drop in leptin levels leads to a fall in both the predominant form of thyroid hormone, thyroxine (T₄), and the active form, triiodothyronine (T₃), over 24–48 hr (Ahima et al., 1996; Connors et al., 1985; Légrádi et al., 1997). Additionally, suppression of the hypothalamic-pituitary-thyroid (HPT) axis during nutritional stress extends to both thyrotropin-releasing hormone (TRH) production in the paraven-

tricular nucleus (PVN) of the hypothalamus and thyroid-stimulating hormone (TSH) production from the pituitary (Blake et al., 1991; Blake et al., 1992; Spencer et al., 1983). To further emphasize the role of leptin in the HPT axis, leptin replacement during food restriction prevented decreases of TH levels in both humans and rodents (Ahima et al., 1996; Légrádi et al., 1997; Rosenbaum et al., 2002). Thus, the nutritional regulation of the HPT axis by leptin becomes an important paradigm to mechanistically understand the importance of this axis in regulating energy expenditure and ultimately body weight.

Two possible mechanisms have been proposed to explain leptin's actions on the HPT axis. These two mechanisms both involve hypothalamic neurocircuitry that regulates TRH production in the PVN: (1) Leptin acts directly through its receptors on hypophysiotropic TRH neurons that project to the median eminence to regulate TSH production in the pituitary (Harris et al., 2001; Nillni et al., 2000; Perello et al., 2006); or (2) leptin regulates TRH neurons indirectly via its actions on pro-opiomelanocortin (POMC) and agouti-related peptide/neuropeptide Y (AgRP/NPY) neurons in the arcuate nucleus (Bjørbaek and Hollenberg, 2002; Fekete et al., 2001; Fekete et al., 2000a; Fekete et al., 2002b; Legradi et al., 1998). Although these pathways are not mutually exclusive, genetic data suggest that leptin signaling is absolutely required for normal function of the HPT axis as mice with leptin receptor mutations have central hypothyroidism whereas mice that lack the MC4R or NPY have normal T₄ levels at baseline (Bates et al., 2004; Erickson et al., 1997; Fekete et al., 2004). Contrary to these observations, the POMC-derived peptide α -melanocyte stimulating hormone (α -MSH) can stimulate *Trh* expression in the PVN, and both AgRP and NPY can suppress *Trh* expression in the PVN, suggesting that neurons within the arcuate nucleus could play a major role in the regulation of the HPT axis through their action on *Trh* (Fekete et al., 2001; Fekete et al., 2000a; Fekete et al., 2002b). Indeed, ablation of the arcuate nucleus by monosodium *L*-glutamate in rats prevented a fasting-induced decrease in *Trh* mRNA, serum TSH, and T₄ levels, strengthening the case for the role of the arcuate nucleus in metabolic regulation of *Trh* (Legradi et al., 1998). However, because MSG may have more widespread effects, including increased leptin insensitivity, this experiment did not rule out other pathways for leptin action (Nillni, 2010). Despite these insights, the exact pathway that regulates leptin's actions on the HPT axis is not clear.

To definitively identify the pathway that allows for the fasting-induced fall in TH levels, we developed a genetic approach using

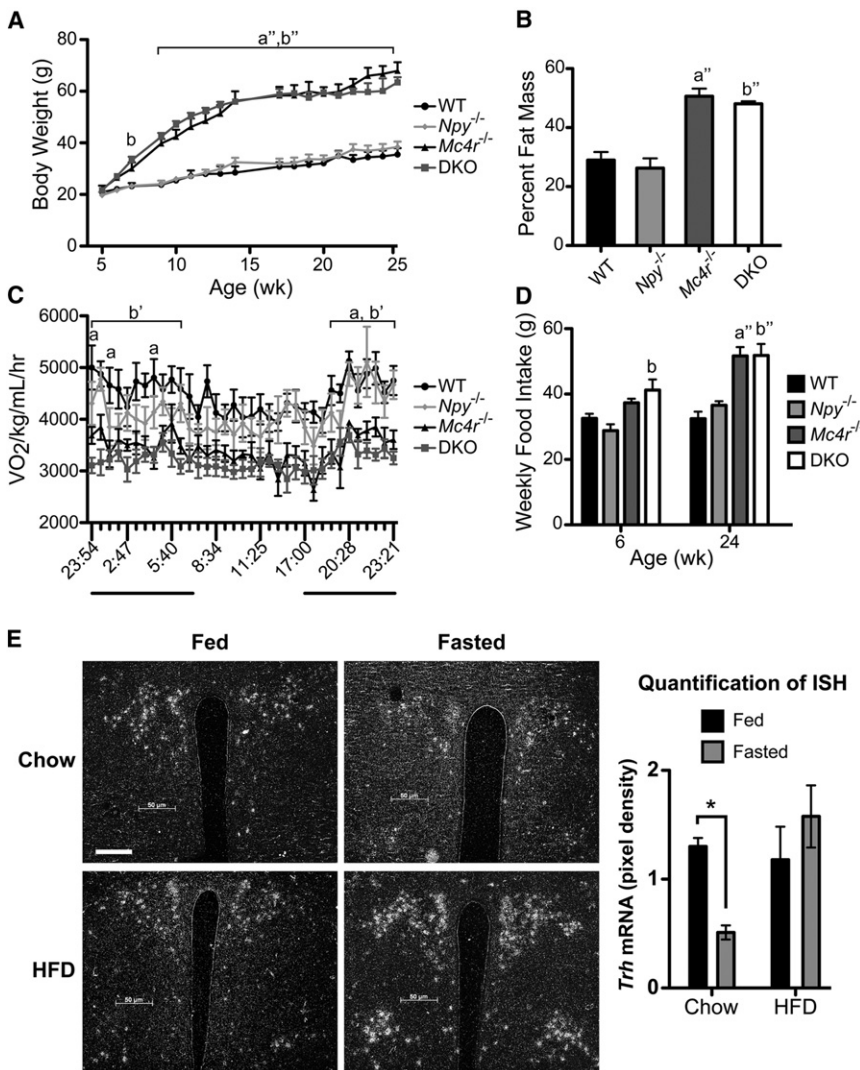


Figure 1. DKO Mice Have a Similar Metabolic Phenotype as *Mc4r*^{-/-} Mice

(A) Average body weight of male WT, *Npy*^{-/-}, *Mc4r*^{-/-} and DKO mice from 5–25 weeks of age. (B) Percent fat mass as measured by DEXA. (C) CLAMS: Mice were placed in a CLAMS apparatus and monitored for oxygen consumption (VO_2) over a 24 hr period. The dark bars indicate dark period in the light cycle. (D) Weekly food intake at 6 and 24 weeks of age. (A)–(D), data are presented as mean \pm SEM, $n = 6$ per genotype. (A) and (C)–(D) were measured with repeated-measures two-way ANOVA with Bonferroni post hoc test. (B) was measured with one-way ANOVA with Tukey's multiple comparison post hoc test. $a = p < 0.05$ *Mc4r*^{-/-} versus WT and *Npy*^{-/-}; $a'' = p < 0.001$ *Mc4r*^{-/-} versus WT and *Npy*^{-/-}; $b = p < 0.05$ DKO versus WT and *Npy*^{-/-}; $b' = p < 0.01$ DKO versus WT and *Npy*^{-/-}; $b'' = p < 0.001$ DKO versus WT and *Npy*^{-/-}. (E) ISH was performed on brain sections using a ³⁵S-labeled riboprobe against mouse *Trh* mRNA. Representative images of fed and fasted C57BL/6 mice on chow or HFD are shown of the PVN at original magnification, $\times 10$; scale bar, 50 μ m. Quantification of *Trh* expression (right panel). Data are presented as relative pixel densities. Significance was tested by two-way ANOVA with Bonferroni post hoc test ($n = 4$ per diet and fed status). Data are presented as mean \pm SEM. * = $p < 0.05$. See also Figures S1 and S2.

mice that lacked both MC4R and NPY and thus would be unable to communicate α -MSH, AgRP, and NPY signals to the PVN. Importantly, we pair-fed these animals to prevent obesity and to equalize leptin levels. Remarkably, we show that signaling through NPY and MC4R controls the fasting-induced suppression of thyroid hormone levels through two mechanisms: (1) Fasting-induced suppression of the central axis requires NPY; and (2) a second pathway based in the liver that enhances the metabolism of TH requires MC4R and NPY. Thus, MC4R and NPY signaling engage both central and peripheral targets to control TH levels during a fast. This engagement of multiple targets underscores the importance of this pathway in the adaptation to nutritional stress.

RESULTS

Obesity Prevents Suppression of the HPT Axis during Fasting

To develop a genetic approach that would determine the importance of POMC and AgRP/NPY signaling in the regulation of the

via CLAMS apparatus (Figure 1C), and hyperphagia (Figure 1D) (Balthasar et al., 2005; Huszar et al., 1997). Like their male counterparts, female DKO mice were similar in phenotype to *Mc4r*^{-/-} mice (data not shown). Interestingly, obese DKO mice had further impaired glucose tolerance compared to *Mc4r*^{-/-} mice (Figure S1).

Since both *Mc4r*^{-/-} and DKO mice were extremely obese, we were concerned with their ability to sense a fast and suppress their HPT axis given their high leptin levels. To test this hypothesis, C57BL/6 mice were placed either on a chow diet or a high fat diet (HFD) for 14 weeks. As expected, HFD-fed mice had increased body weight and leptin levels that were 20-fold higher than WT mice (Figures S2A–S2B). We then subjected these mice to a 36 hr fast. While leptin levels fell during a fast in HFD-fed mice, they were over 20 times higher than fasted leptin levels in chow-fed mice (Figure S2B). Whereas total thyroxine (bound T_4 plus free T_4 in serum: TT_4) and total triiodothyronine (bound T_3 plus free T_3 in serum: TT_3) levels fell significantly in chow-fed mice, they did not change in HFD-fed mice as previously demonstrated in rats (Figures S2C–S2D) (Goodman

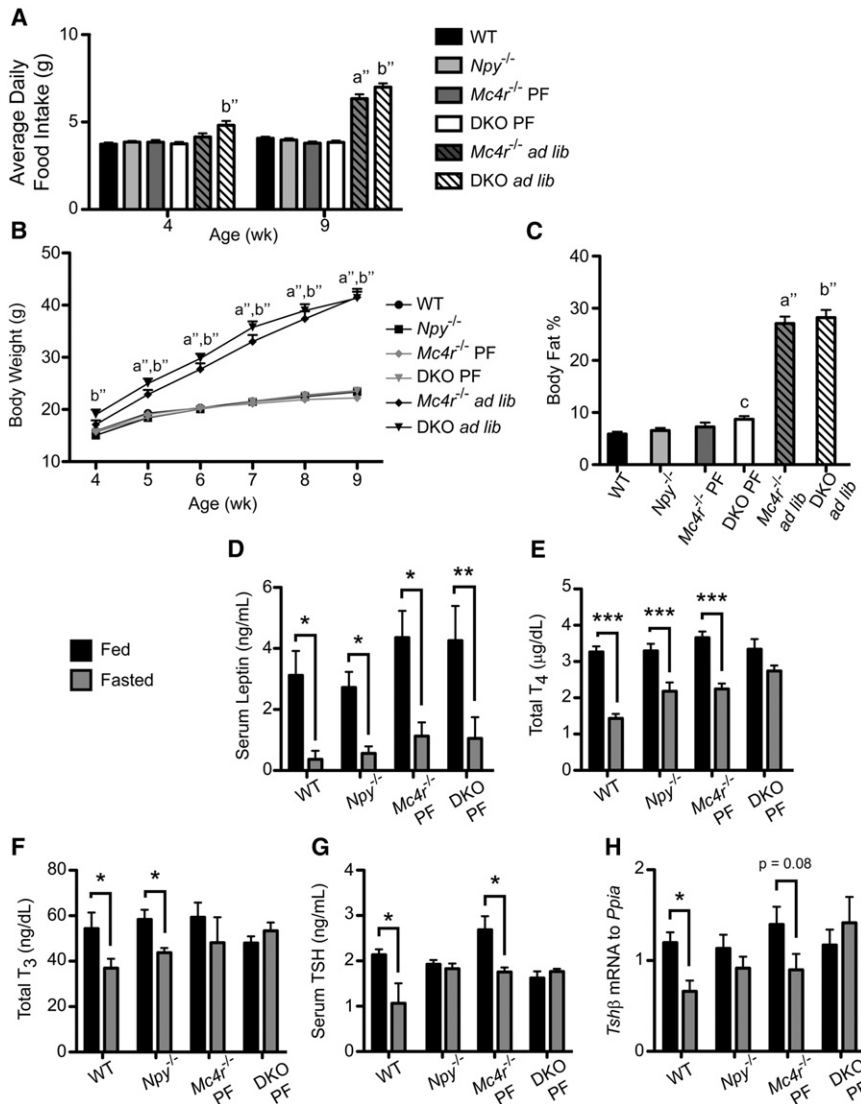


Figure 2. Pair-Fed Mice Lacking Both NPY and MC4R Fail to Suppress Thyroid Hormone and TSH Levels

(A–C) Average daily food intake (A), Body Weight (B) and Body Fat percentage (C) of combined NPY/MC4R cohorts 2 and 3: WT (n = 40), *Npy*^{-/-} (n = 38), pair-fed *Mc4r*^{-/-} (*Mc4r*^{-/-} PF; n = 20), and DKO (DKO PF; n = 28), and ad libitum *Mc4r*^{-/-} (*Mc4r*^{-/-} ad lib; n = 11) and DKO (DKO ad lib; n = 12) mice. (A) and (B) were analyzed by repeated-measures two-way ANOVA with Bonferroni post hoc test. (C) was analyzed via one-way ANOVA with Tukey’s multiple comparison post hoc test. For (A)–(C), data are presented as mean ± SEM. a” = p < 0.001 *Mc4r*^{-/-} ad lib versus WT, *Npy*^{-/-}, *Mc4r*^{-/-} PF, and DKO PF; b” = p < 0.001 DKO ad lib versus WT, *Npy*^{-/-}, *Mc4r*^{-/-} PF, and DKO PF; c = p < 0.05 DKO PF versus WT.

(D) Serum leptin levels were measured in cohort 2 fed and fasted WT (n = 5, n = 7), *Npy*^{-/-} (n = 10, n = 10), *Mc4r*^{-/-} PF (n = 4, n = 4), and DKO PF (n = 6, n = 5) mice.

(E) Combined cohort 2 and 3 Total T₄ (TT₄) levels of fed and fasted WT (n = 23, n = 17), *Npy*^{-/-} (n = 21, n = 17), *Mc4r*^{-/-} PF (n = 10, n = 10), and DKO PF (n = 16, n = 12) mice.

(F) Total T₃ (TT₃) concentration of cohort 2 WT (n = 5, n = 7), *Npy*^{-/-} (n = 10, n = 10), *Mc4r*^{-/-} PF (n = 4, n = 4), and DKO PF (n = 6, n = 5) mice.

(G) Serum TSH concentration of cohort 2 WT (n = 4, n = 4), *Npy*^{-/-} (n = 10, n = 10), *Mc4r*^{-/-} PF (n = 3, n = 3), and DKO PF (n = 6, n = 5) mice.

(H) *Tshβ* levels relative to the control gene *Ppia* (cyclophilin) were measured via QPCR in cohort 3 WT (n = 18, n = 11), *Npy*^{-/-} (n = 10, n = 8), *Mc4r*^{-/-} PF (n = 7, n = 7), and DKO PF (n = 9, n = 7) mice. For (D)–(H), data are presented as mean ± SEM. Significance was measured with two-way ANOVA with Bonferroni post hoc test. * = p < 0.05; ** = p < 0.01; *** = p < 0.001. See also Figure S3.

et al., 1980). Consistent with these observations, *Trh* mRNA decreased 60% in the PVN of fasted chow-fed animals but did not fall in the PVN of HFD-fed animals (Figure 1E). *Trh* mRNA did not change in the lateral hypothalamic area (LHA), anterior hypothalamic area (AHA), or the dorsomedial hypothalamus (DMH) regardless of nutritional status in chow or HFD-fed mice (Figures S2E–S2J). Taken together these data demonstrate that obesity prevents fasting-induced suppression of the HPT axis at 36 hr fasting. Therefore, to properly interpret the roles of the MC4R and NPY in DKO mice we would need to prevent obesity in these models.

Pair-Feeding Rescues Obese Phenotype of *Mc4r*^{-/-} and DKO Mice

To prevent obesity in *Mc4r*^{-/-} and DKO animals we developed a pair-feeding paradigm, such that the food intake of *Mc4r*^{-/-} and DKO mice was adjusted to WT levels on a daily basis. Importantly, using this method we were able to deliver

identical amounts of food to all genotypes through 9 weeks of age (Figure 2A), which allowed body weight (Figure 2B) and fat mass (Figure 2C) to be relatively equalized across genotypes in the pair-fed groups compared to the ad libitum-fed *Mc4r*^{-/-} and DKO mice. Furthermore, pair-fed *Mc4r*^{-/-} and DKO mice with a body fat percentage of greater than 15% as measured by MRI were excluded from the study.

We next fasted groups of WT, *Npy*^{-/-}, and pair-fed *Mc4r*^{-/-} and DKO mice to determine the response of the HPT axis. As shown in Figure 2D, leptin levels of fed mice were slightly but not significantly higher in pair-fed *Mc4r*^{-/-} and DKO mice compared to WT and *Npy*^{-/-} mice but fell equivalently after the fast. A comparison of the leptin levels of pair-fed *Mc4r*^{-/-} and DKO mice to levels in fed and fasted ad libitum-fed *Mc4r*^{-/-} and DKO mice (Figure S3A) confirms the effectiveness of the pair-feeding paradigm in establishing normal leptin levels in *Mc4r*^{-/-} and DKO animals.

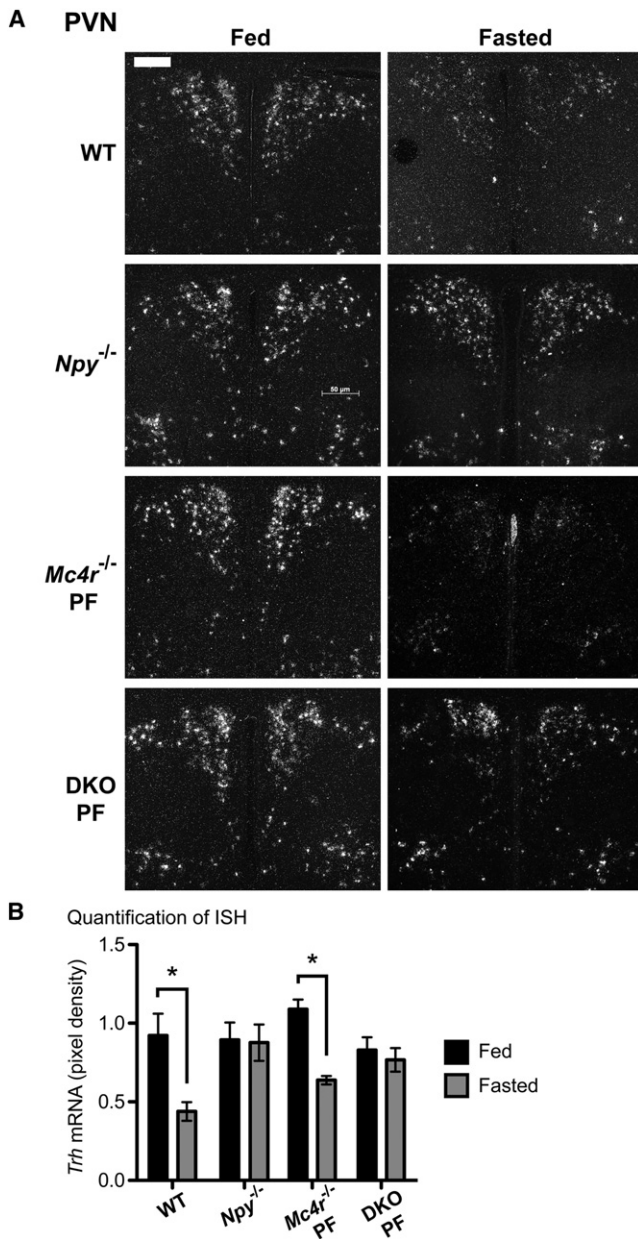


Figure 3. NPY Is Required for Fasting-Induced Suppression of *Trh* Expression

ISH was performed on brain sections using a ³⁵S-labeled riboprobe against mouse *Trh* mRNA. (A) Representative images of fed and fasted WT (n = 5, n = 6), *Npy*^{-/-} (n = 8, n = 9), and pair-fed *Mc4r*^{-/-} PF (n = 4, n = 6) and DKO PF (n = 4, n = 5) mice are shown of the PVN, ×10; scale bar, 50 μm. (B) Quantification of *Trh* expression. Data are presented as mean pixel density ± SEM. Significance was measured with two-way ANOVA with Bonferroni post hoc test. * = p < 0.05. See also Figure S4.

Fasting-Induced Suppression of the HPT Axis Is Impaired in *Npy*^{-/-} Mice and Prevented in DKO Mice

To determine the contribution of POMC and AgRP/NPY signaling in fasting-induced suppression of the HPT axis, we measured several factors in the HPT axis in fed and fasted WT, *Npy*^{-/-},

and pair-fed *Mc4r*^{-/-} and DKO mice. Following 36 hr of fasting, the concentration of TT₄ dropped significantly, by 56%, in WT mice (Figure 2E). As previously described, TT₄ fell significantly by 36% in *Npy*^{-/-} mice, although the drop was not as great as in WT mice with 36 hr of food deprivation (Erickson et al., 1997). TT₄ concentrations also fell in pair-fed *Mc4r*^{-/-} mice, but did not change in pair-fed DKO mice with fasting. As expected, TT₄ concentrations measured in ad libitum-fed *Mc4r*^{-/-} and DKO mice did not change with fasting, consistent with the impact of very high levels of leptin on the HPT axis (Figure S3B). Concentrations of TT₃ and serum TSH remained unchanged in fasting pair-fed DKO mice compared to the respective 32% and 50% drop seen in WT mice (Figures 2F–2G). Whereas TT₃ levels fell by 25% in *Npy*^{-/-} mice, serum TSH did not. Conversely, pair-fed *Mc4r*^{-/-} mice suppressed serum TSH by 31% but not TT₃ with fasting. No changes in TT₃ or serum TSH were observed in ad libitum-fed *Mc4r*^{-/-} and DKO mice (Figures S3C–S3D).

Since serum TSH does not always accurately reflect TRH action, we confirmed the serum TSH data by measuring *Tshβ* mRNA levels from the pituitaries of a separate cohort of fed and fasted WT, *Npy*^{-/-}, and pair-fed *Mc4r*^{-/-} and DKO mice via quantitative RT-PCR (QPCR). Fasting *Npy*^{-/-} and pair-fed DKO mice failed to suppress *Tshβ* compared to WT mice (1.8-fold) while *Tshβ* expression trended to suppression with fasting (1.5-fold, p = 0.08) in *Mc4r*^{-/-} mice (Figure 2H). Thus, at the level of the pituitary, NPY appears critical for suppression of *Tshβ* and serum TSH during fasting, which suggests that NPY may mediate this effect through communication of fasting signals to TRH neurons in the PVN.

To test this hypothesis, we performed in situ hybridization for *Trh* expression in the PVN of the hypothalamus, the location of hypophysiotropic neurons that project from the PVN to the median eminence to release TRH peptide. As has been previously demonstrated, *Trh* expression was significantly decreased, by 53%, in fasted WT mice when compared to fed controls (Figures 3A–3B) (Blake et al., 1991). Similarly, fasted pair-fed *Mc4r*^{-/-} mice also repressed *Trh* expression by 42% in the PVN when compared to fed controls. Importantly, *Trh* was only regulated during fasting in WT and *Mc4r*^{-/-} mice in the PVN and not in other areas of the hypothalamus where *Trh* is expressed, which includes the AHA, the LHA, and the DMH (Figures S4A–S4C). Remarkably, fasting did not regulate *Trh* levels in the PVN of *Npy*^{-/-} mice. Similar results were seen in pair-fed DKO mice, validating the key requirement for NPY in mediating the fasting-induced suppression of *Trh* (Figure 3). Furthermore, *Trh* expression did not change in fasting ad libitum-fed *Mc4r*^{-/-} and DKO mice, again highlighting the impact of obesity on the HPT axis (Figure S4D). Taken together, NPY signaling is the major contributor to communicating fasting signals to the central HPT axis, as *Trh*, *Tshβ*, and serum TSH remained unchanged in fasted *Npy*^{-/-} and pair-fed DKO mice. Despite this, TT₄ and TT₃ levels still drop in fasted *Npy*^{-/-} mice, but not in pair-fed DKO mice. This suggests that the MC4R plays a role in suppression of thyroid hormone levels during fasting through a TRH-independent pathway. This is further supported by the failure of pair-fed *Mc4r*^{-/-} mice to fully suppress TT₄ levels and TT₃ levels during a fast.

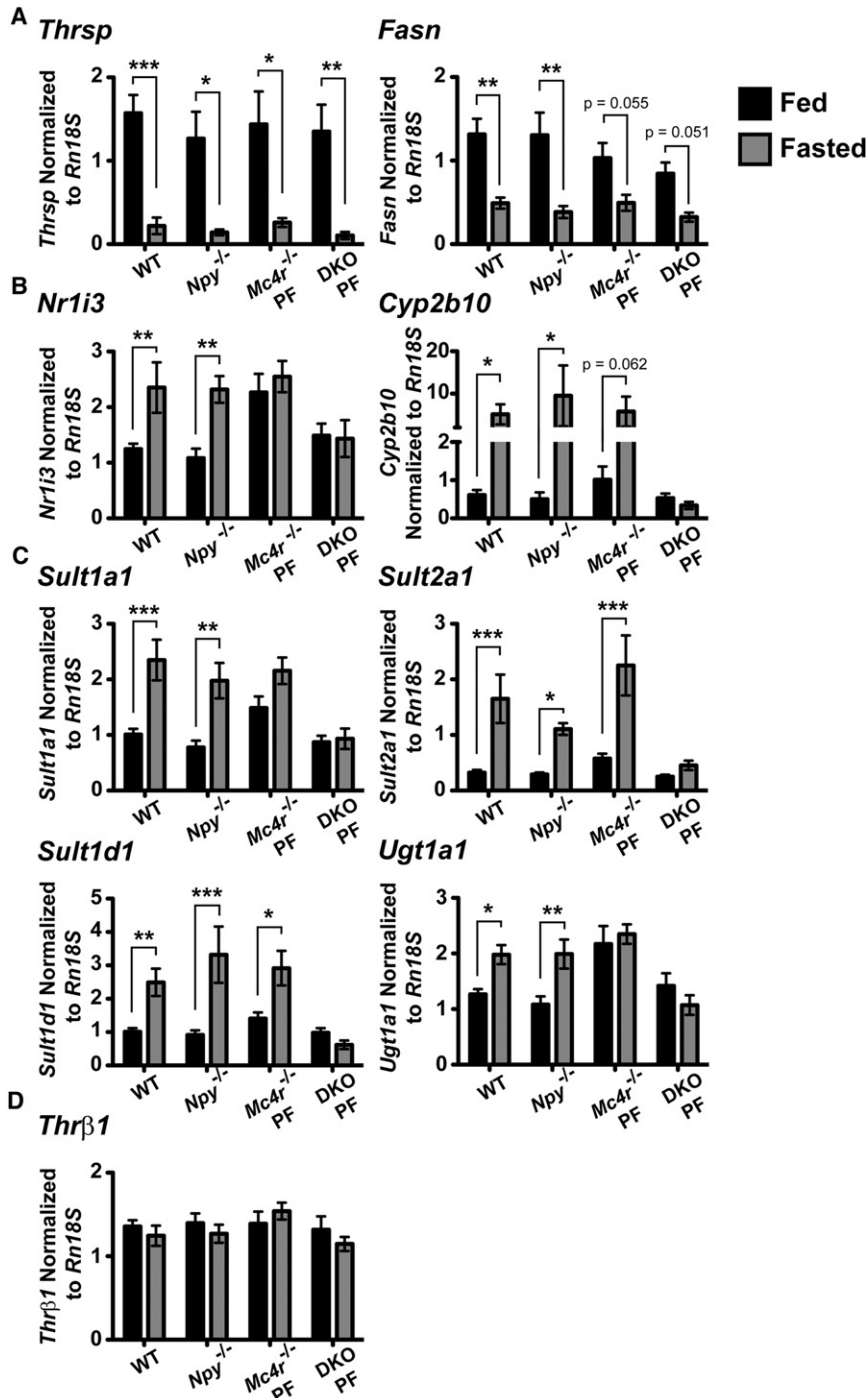


Figure 4. MC4R and NPY Are Necessary for Upregulation of *Nr1i3* and Its Target Genes during Fasting

Expression of various genes in the liver was measured via qPCR and normalized to *Rn18S* expression in WT (n = 17, n = 10), *Npy*^{-/-} (n = 10, n = 7), and pair-fed *Mc4r*^{-/-} PF (n = 7, n = 7) and DKO PF (n = 9, n = 7) mice. (A) Expression of *Thrsp* and *Fasn*. (B) Expression of *Nr1i3* and its classical target gene, *Cyp2b10*. (C) Expression of the *Nr1i3*-target genes involved in hepatic TH metabolism: *Sult1a1*, *Sult2a1*, *Sult1d1*, and *Ugt1a1*. (D) Expression of *Thrβ1*. In all panels, data are presented as mean ± SEM. Significance was measured with two-way ANOVA with Bonferroni post hoc test. * = p < 0.05; ** = p < 0.01; *** = p < 0.001.

tane receptor (CAR), or nuclear receptor subfamily 1, group I, member 3 (*Nr1i3*), and its target genes, sulfotransferases (Sults) and UDP-glucuronosyltransferases (Ugts), are pivotal to hepatic TH metabolism as they accelerate deiodination and clearance of TH (Maglich et al., 2004; Mol and Visser, 1985; Qatanani et al., 2005; Vansell and Klaassen, 2002; Visser, 1996). Furthermore, *Nr1i3* and several of its target genes are upregulated in the liver during fasting to increase TH metabolism (Maglich et al., 2004). Indeed, fasting-induced repression of TH levels is impaired in *CAR*^{-/-} mice (Maglich et al., 2004).

First, to establish the sensitivity of the livers of *Npy*^{-/-} and pair-fed *Mc4r*^{-/-} and DKO mice to fasting signals, we measured expression of lipogenic genes that decrease during fasting via qPCR (Carr et al., 1984; Clarke et al., 1990). Thyroid hormone responsive SPOT14 (*Thrsp*) mRNA was significantly repressed with fasting in all genotypes (Figure 4A) while fatty acid synthase (*Fasn*) mRNA levels also decreased with fasting in WT (2.6-fold) and *Npy*^{-/-} (3.9-fold) mice and trended toward reduced levels in pair-fed *Mc4r*^{-/-} (2.1-fold, p = 0.055) and DKO (2.6-fold, p = 0.051) mice.

We next assessed the roles of MC4R and NPY signaling pathways in the regulation of *Nr1i3* and its target genes in the

liver. *Nr1i3* expression was upregulated during fasting (1.8-fold, Figure 4B) in WT mice, which has been demonstrated previously (Maglich et al., 2004). Additionally, *Nr1i3* expression increased with fasting in *Npy*^{-/-} mice (2.1-fold), however expression of *Nr1i3* remained unchanged in pair-fed *Mc4r*^{-/-} and DKO mice. To understand the impact of impaired *Nr1i3* regulation during fasting, we measured the expression of the classic *Nr1i3*-target gene, cytochrome p450, family 2, subfamily b, polypeptide 10 (*Cyp2b10*) (Honkakoski et al., 1998).

Hepatic Metabolism of Thyroid Hormone during Fasting Requires the MC4R and NPY

To explain the adaptive response of TH levels in *Npy*^{-/-} mice during a fast despite the absence of *Thr* repression, we hypothesized that fasting-induced peripheral hepatic metabolism of TH was operative in *Npy*^{-/-} mice, but not in pair-fed DKO mice (de Herder et al., 1988). This circuit would allow TH levels to fall, independent of TRH, and would require the MC4R. Previous studies have shown that the orphan nuclear receptor constitutive andros-

Expression of *Cyp2b10* has been shown to increase with fasting, and our results agree, as WT (8.4-fold) and *Npy*^{-/-} (18.8-fold) mice both upregulate hepatic *Cyp2b10* expression following a 36 hr fast (Figure 4B) (Brown et al., 1995). *Cyp2b10* expression in pair-fed DKO mice showed no response to fasting, whereas pair-fed *Mc4r*^{-/-} mice trended toward an upregulation (5.7-fold, *p* = 0.062). These results implicate both MC4R and NPY in Nr1i3-mediated regulation of hepatic TH metabolism, as *Cyp2b10* may respond to fasting in pair-fed *Mc4r*^{-/-} mice, but not in DKO mice.

As failure of hepatic *Nr1i3* to respond to fasting signals in *Mc4r*^{-/-} and DKO mice may explain the inability of DKO mice to suppress TH levels during a fast, we next measured the Nr1i3 targets responsible for hepatic thyroid hormone metabolism. Previous studies have demonstrated that Phase II sulfotransferases mediate hepatic metabolism of TH by sulfating the phenol group and marking TH for inactivation, which occurs via the type 1 deiodinase (Mol and Visser, 1985). Sulfotransferase 1A1 (*Sult1a1*), which sulfates TH in vitro and has a high affinity for T₄ and T₃, is upregulated during fasting (Kester et al., 1999; Li et al., 2001; Maglich et al., 2004). *Sult1a1* expression was significantly increased with fasting in WT (2.3-fold) and *Npy*^{-/-} (2.5-fold) mice compared to their fed counterparts (Figure 4C). Both pair-fed *Mc4r*^{-/-} and DKO mice failed to upregulate *Sult1a1* with fasting. Additionally, we examined two other sulfotransferases: *Sult2a1* preferentially sulfates T₃, and *Sult1d1* has a very similar protein structure to *Sult1a1*, although its role in the liver is still unknown (Li and Anderson, 1999; Sakakibara et al., 1995). Both genes were significantly upregulated in WT (5.1-fold and 2.5-fold, respectively), *Npy*^{-/-} (3.7-fold and 3.6-fold, respectively) and pair-fed *Mc4r*^{-/-} (3.9-fold and 2.1-fold, respectively) mice, whereas pair-fed DKO mice did not regulate these genes during fasting (Figure 4C). Taken together, these data confirm that pair-fed DKO mice have impaired Nr1i3-mediated hepatic TH metabolism. To a lesser degree, pair-fed *Mc4r*^{-/-} mice are also defective in Nr1i3 signaling, as they fail to upregulate both *Nr1i3* and *Sult1a1* with fasting, but *Cyp2b10*, *Sult2A1*, and *Sult1d1* are responsive. Removal of NPY signaling in addition to MC4R signaling ablates the liver's ability to metabolize TH via sulfation.

Nr1i3 also regulates another pathway of phase II TH metabolism, glucuronidation. Conjugation of TH by Ugts increases the water solubility of TH, promoting biliary excretion (Burchell and Coughtrie, 1989). We measured the expression of the phenol/bilirubin UGT 1 family, polypeptide A1 (*Ugt1a1*) responsible for the glucuronidation of T₄ (Beetstra et al., 1991; Findlay et al., 2000). Following a 36 hr fast, *Ugt1a1* expression was upregulated in both WT (3.8-fold) and *Npy*^{-/-} (3.5-fold) mouse livers compared to fed counterparts (Figure 4C) (Maglich et al., 2004). No change in *Ugt1a1* expression was detected in pair-fed *Mc4r*^{-/-} and DKO mice. These data further emphasize the role of MC4R and NPY in Nr1i3-mediated hepatic metabolism as both sulfation and glucuronidation of TH are impaired in pair-fed DKO mice. Furthermore, these results underline the mechanism by which *Npy*^{-/-} mice and not DKO mice are able to repress TH levels during fasting despite impaired central HPT axis regulation. To ensure TH signaling was not impacted in fasting mouse livers, we measured the expression of thyroid hormone receptor β1 (*Thrβ1*) mRNA levels. *Thrβ1* levels were

not affected by fasting or by deletion of NPY, MC4R, or both (Figure 4D).

Taken together, these data demonstrate that signaling through NPY and the central melanocortin system regulates peripheral thyroid hormone metabolism in the liver and provides evidence of an additional role for this pathway in the regulation of energy expenditure.

Sulfotransferases Become Nr1i3-Resistant with HFD

To determine the impact of obesity and high leptin levels on thyroid hormone metabolism in the liver, we examined the expression of *Nr1i3* mRNA and its target genes in the livers of mice on a chow diet or HFD. As before, we first measured *Thrsp* and *Fasn* expression to determine the sensitivity of livers to fasting in mice on a HFD. Both *Thrsp* and *Fasn* expression decreased with fasting in mice on a chow diet (3.8-fold and 1.4-fold, respectively) and HFD (3.6-fold and 2.7-fold, respectively, Figure 5A). Next, we measured *Nr1i3* expression during fasting in HFD-fed mice. *Nr1i3* levels were upregulated during fasting in mice on a HFD (3.9-fold) compared to their fed controls, which was similar to mice on a chow diet (3.2-fold, Figure 5B). The classic Nr1i3 target, *Cyp2b10*, was increased in response to fasting in mice on a chow diet (6.0-fold, Figure 5B). However, there was no change in *Cyp2b10* levels in fasted mice on a HFD. Additionally, HFD-fed mice failed to upregulate the expression of *Sult1a1*, *Sult2a1*, and *Sult1d1* during fasting, whereas *Ugt1a1* levels increased with fasting (2.3-fold, Figure 5C). *Thrβ1* levels were unaffected by HFD (Figure 5D). Although *Nr1i3* expression is responsive to fasting in mice on HFD, these data suggest that the expression of *Cyp2b10* and sulfotransferases are resistant to Nr1i3-mediated induction during fasting. Resistance to Nr1i3 on a HFD does not permeate to all target genes since *Ugt1a1* still increases during fasting in HFD-fed mice.

DISCUSSION

Nutritional regulation of the HPT axis provides an important adaptive response to caloric restriction. Indeed, humans who undergo weight loss or controlled caloric restriction experience both reduced TSH and thyroid hormone levels (Rosenbaum et al., 2005; Rosenbaum et al., 2002). Previous studies in both humans and rodents have demonstrated that the fall in leptin levels during caloric restriction controls thyroid hormone levels (Ahima et al., 1996; Rosenbaum et al., 2002; Sanchez et al., 2004). Prevailing hypotheses suggest that leptin's role, which can be either direct through leptin receptors in the PVN or indirect via POMC and AgRP/NPY neurons in the arcuate nucleus, is mediated by its ability to regulate *Trh* expression in hypophysiotropic neurons in the PVN (Fekete et al., 2001; Fekete et al., 2000a; Fekete et al., 2002b; Huo et al., 2004; Kim et al., 2000; Légrádi et al., 1997; Legradi et al., 1998). While showing the importance of POMC and AgRP/NPY signaling in the regulation of *Trh* expression, we also demonstrate their effect on a key peripheral hepatic circuit that controls thyroid hormone levels in the setting of nutritional stress. Thus, leptin engages the HPT axis on multiple levels to control energy expenditure during nutritional stress.

Prior to this report, the specific roles of arcuate nucleus neuropeptides downstream of leptin in the regulation of *Trh* gene

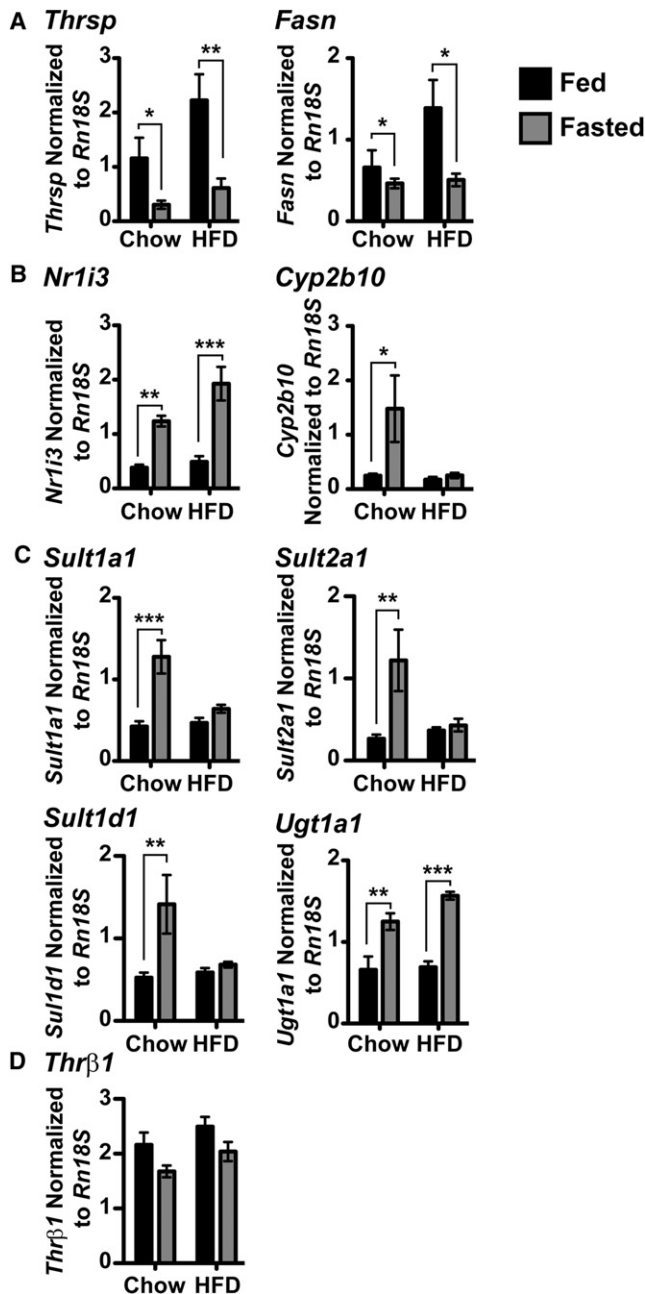


Figure 5. Genes Enhancing Hepatic Thyroid Hormone Metabolism are Nr1i3-Resistant in HFD-Fed Mice

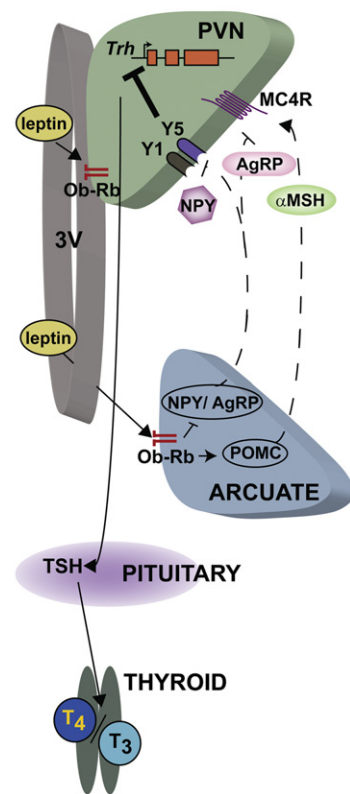
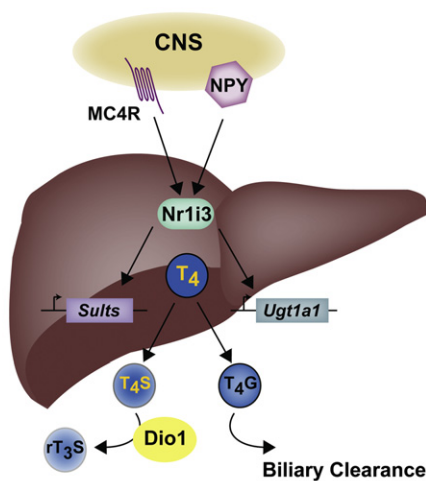
Expression of various genes in the liver was measured via qPCR and normalized to *Rn18S* expression in fed and fasted C57BL/6 mice on chow (n = 4, n = 4) or HFD (n = 4, n = 4). (A) Expression of *Thrsp* and *Fasn*. (B) Expression of *Nr1i3* and *Cyp2b10*. (C) Expression of the Nr1i3-target genes: *Sult1a1*, *Sult2a1*, *Sult1d1*, and *Ugt1a1*. (D) Expression of *Thrβ1*. In all panels, data are presented as mean ± SEM. Significance was measured with two-way ANOVA with Bonferroni post hoc test. * = p < 0.05; ** = p < 0.01; *** = p < 0.001.

expression had not been determined. To assess these roles, we created mice that lack either MC4R signaling or NPY signaling or both. Although we have deleted MC4R and NPY signaling globally, reports to date implicate the arcuate nucleus as the source

of NPY, AgRP, and α -MSH to the PVN (Fekete et al., 2001; Fekete et al., 2000a; Legradi et al., 1998). Previous studies that modified NPY signaling in different areas of the hypothalamus did not report abnormalities in thyroid hormone levels (Chao et al., 2011; Gardiner et al., 2005; Tiesjema et al., 2007; Tiesjema et al., 2009; Yang et al., 2009). Importantly, we had to control for the obese phenotype in *Mc4r*^{-/-} and DKO mice in order to use these models as the high leptin levels present in obesity preclude a fall in thyroid hormone levels during a fast. As suggested from earlier studies, input from the arcuate nucleus is not required for basal function of the HPT axis, as *Mc4r*^{-/-}, *Npy*^{-/-}, and DKO mice all have normal TH, serum TSH, and *Tshβ* levels at baseline (Erickson et al., 1997; Fekete et al., 2004). In contrast, mice with leptin receptor mutations that impair signal transducer and activator of transcription 3 (STAT3) signaling have central hypothyroidism (Bates et al., 2004). Taken together, these contrasting data suggest that leptin acts independently of α -MSH, AgRP, and NPY to establish the set point of the HPT axis in the basal fed state (Perello et al., 2010). Although this does suggest a direct action of leptin on TRH neurons, it remains possible that cocaine and amphetamine-regulated transcript (CART) production in the arcuate nucleus could play a role initiating the set point of the HPT axis (Broberger, 1999; Fekete et al., 2000b; Kádár et al., 2010; Raptis et al., 2004).

Unlike the basal state, the indirect pathway plays a critical role in fasting. The data presented herein demonstrate that NPY signaling is required for suppression of *Trh* in hypophysiotropic TRH neurons and that both the MC4R and NPY signaling pathways are required for the full suppression in TH levels seen during a fast (Figure 6). It is likely that NPY targets TRH neurons via the NPY Y1 or Y5 receptors and represses *Trh* expression by downregulation of the cAMP-PKA pathway, a known activator of *Trh* expression (Fekete et al., 2002a; Herzog et al., 1992; Sarkar and Lechan, 2003). The failure of *Npy*^{-/-} mice to repress *Tshβ* during a fast suggests that TRH peptide levels in these animals are also unchanged and further establishes the role of NPY in controlling *Trh* expression in the PVN.

Despite the clear role of NPY in controlling fasting-induced suppression of *Trh* expression, TT_4 and TT_3 levels were still partially decreased in *Npy*^{-/-} mice during a fast, which indicates that another pathway controlling TH levels must be operative. Strikingly, the HPT axis was entirely resistant to fasting in pair-fed DKO animals, demonstrating that the separate pathway controlling TH levels may require the MC4R. While it is possible that changes in TSH bioactivity contribute to the fasting response of the HPT axis in general, this is unlikely in DKO mice. Here, serum TSH and TT_4 levels are the same between fed and fasted mice, which is indicative of similar TSH bioactivity (Nikrodhanond et al., 2006). Likewise, serum TSH and TT_4 levels are similar across all genotypes in the fed state, suggesting that TSH bioactivity is not altered by MC4R and/or NPY deletion. Certainly, when *Trh* falls during a fast in WT and pair-fed *Mc4r*^{-/-} mice, both total and biologically active serum TSH may fall. Given that the MC4R activates key hepatic pathways that control metabolism, we focused on the hepatic metabolism of TH due to its important role in determining TH levels during acute illness (Nogueiras et al., 2007; Wiersinga, 2005). Notably, we have demonstrated that the MC4R and NPY are required for the fasting-induced upregulation of the nuclear receptor *Nr1i3*, which

A Central T₄ RegulationB Hepatic T₄ MetabolismFigure 6. Model of NPY and MC4R Signaling in T₄ Regulation during Fasting

(A) During fasting, reduced leptin levels invoke an increase in *Npy* expression in the arcuate nucleus. NPY signals through Y1 and Y5 receptors in the PVN to repress *Trh* expression. Central repression of *Trh* leads to a decrease in *Tshβ* mRNA levels in the pituitary and, thus, reduced T₄ and T₃ levels from the thyroid. *Trh* and serum TSH levels remain unchanged in fasting *Npy*^{-/-} mice, yet T₄ and T₃ levels are decreased. Adapted from Vella and Hollenberg (2009) with permission of The Endocrine Society.

(B) *Npy*^{-/-} mice are able to suppress TH levels during fasting through Nr13-mediated hepatic metabolism of TH, which requires both MC4R and NPY. The MC4R and NPY may act on the liver directly, through sympathetic outputs or a combination of both. The Nr13-targets SULT1A1, SULT2A1, SULT1D1, and UGT1A1 inactivate TH through sulfation (T₄S) or glucuronidation (T₄G). These genes are regulated during fasting in *Npy*^{-/-} mice, but not in DKO mice. T₄S is converted to reverse T₃ sulfate (rT₃S) by the type 1 deiodinase (Dio1), and T₄G is cleared through biliary excretion. 3V = third ventricle. AgRP = agouti-related peptide. α-MSH = α-melanocyte-stimulating hormone. Nr13 = constitutive androstane receptor. CNS = central nervous system. Dio1 = type 1 deiodinase. MC4R = melanocortin 4 receptor. Ob-Rb = leptin receptor, long form. NPY = neuropeptide Y. POMC = pro-opiomelanocortin. PVN = paraventricular nucleus. *Sults* = sulfotransferases. T₃ = triiodothyronine. rT₃S = reverse triiodothyronine sulfate. T₄ = thyroxine. T₄G = glucuronidated thyroxine. T₄S = thyroxine sulfate. TSH = thyroid-stimulating hormone. *Trh* = thyrotropin-releasing hormone. *Ugt1a1* = UDP-glucuronosyltransferase 1A1. Y1, Y5 = NPY receptors.

then controls the expression of key sulfotransferases and *Ugt1a1* that are necessary for the metabolism and excretion of T₄ and T₃ (Maglich et al., 2004). Whereas pair-fed *Mc4r*^{-/-} mice are partially deficient in the activation of this Nr13-mediated pathway, DKO mice are completely defective, further supporting the concept of a compound effect by NPY and MC4R signaling on TH levels (Lechan and Fekete, 2006). Further work will be required to delineate how the hypothalamic neurocircuitry, sympathetic output, and/or direct effects controlled by MC4R and NPY signaling regulate Nr13-driven metabolism of TH in the liver. Importantly, we have seen upregulation of *Nr13* and its targets at least as early as 16 hr of fasting (data not shown), consistent with both the measured 13–18 hr half-life of T₄ in mice and the greater than 50% drop in TT₄ levels during a 36 hr fast, which supports the relevance of this pathway in vivo (van Buul-Offers et al., 1983).

In summary, the development of mice that lack key arcuate nucleus inputs to hypothalamic TRH neurons in the PVN has allowed us to establish the mechanism by which leptin regulates the HPT axis during fasting. While we demonstrate the unique requirement of NPY signaling for suppression of the central axis, we identify a fasting-induced hepatic pathway necessary for full suppression of TH levels that requires both MC4R and NPY signaling and is independent of TRH expression (Figure 6). These data suggest that the fall in TH levels seen during caloric restriction in humans could be targeted either centrally or peripherally and that the fasting-induced activation of the

MC4R and NPY controls not only food intake but also energy expenditure through the regulation of TH levels.

EXPERIMENTAL PROCEDURES

Animals

All experiments were approved by the Beth Israel Deaconess Medical Center Institutional Animal Care and Use Committee. For HFD experiments, male C57BL/6 mice were obtained from The Jackson Laboratory (Bar Harbor, ME). NPY/MC4R mice were generated by crossing the previously described mice heterozygous for the loxTB *Mc4r* allele (*Mc4r*^{loxTB}) with mice heterozygous for a functional *Npy* allele (*Npy*^{+/+}) (Balthasar et al., 2005; Erickson et al., 1996). Details of the cohorts in these studies are provided in the Supplemental Experimental Procedures. Collections of serum, flash frozen tissue, and formalin-fixed brains and storage have been previously described (Astopova et al., 2011; Sugrue et al., 2010). All mice were given free access to water and the appropriate diet. In fasting experiments, mice were fasted for 36 hr. In pair-feeding experiments, obesity-prone *Mc4r*^{-/-} and DKO mice were fed the same amount of food consumed by WT mice on the previous day, from weaning to the end of the experiment. Pair-fed *Mc4r*^{-/-} and DKO mice with a body fat percentage of greater than 15% were excluded from the study.

CLAMS

NPY/MC4R mice from cohort 1 were placed in a Comprehensive Lab Animal Monitoring System apparatus (Columbus Instruments, Columbus OH) at 8–12 weeks of age and monitored for oxygen consumption, using a previously described protocol (Segal-Lieberman et al., 2003a; Segal-Lieberman et al., 2003b). Mice were acclimated in CLAMS cages for 48 hr prior to measurement of metabolic parameters.

DEXA Analysis

Body mass composition of male and female mice from NPY/MC4R cohort 1 was measured at 18–20 weeks of age by dual energy X-ray absorptiometry using a densitometer for mice (Piximus Lunar PIXI 51048).

EchoMRI

At 9 weeks of age, NPY/MC4R mice from cohorts 2 and 3 were subjected to magnetic resonance imaging (MRI) using EchoMRI (Echo Medical Systems, Houston, TX) to determine the body composition.

Serum Analysis

Total serum T₄ (TT₄) and T₃ (TT₃) were measured by solid-phase RIA (Coat-a-Count; Diagnostic Products Corp., Los Angeles, CA) in 25 and 50 μ l of serum, respectively. Serum leptin concentration was measured via ELISA (Quantikine M Mouse Leptin Immunoassay, R & D Systems, Minneapolis, MN). Serum TSH was measured via IRMA (TSH IRMA, ALPCO, Salem, NH) in 100 μ l of serum.

Real-Time Quantitative PCR

Total RNA was extracted from frozen tissues with STAT-60 reagent (Tel-Test, Friendswood, TX). Then 0.5 μ g of total RNA was reverse transcribed using Advantage RT-for-PCR kit (CLONTECH, Mountain View, CA) with random hexamer primers. TaqMan gene expression assays for all mRNAs were purchased from Applied Biosystems (Carlsbad, CA). Quantitative PCR was performed in triplicate using the 800HT thermal cycler (Applied Biosystems, Foster City, CA). Relative mRNA levels were calculated using the standard curve method and normalized to peptidylprolyl isomerase A (cyclophilin, *Ppia*) (pituitary) or 18S ribosomal RNA (*Rn18S*, liver).

In Situ Hybridization

Trh expression was detected by ISH using ³⁵S-radiolabeled antisense riboprobe on formalin-fixed brain slices as described previously (Sugrue et al., 2010). Dark-field digital images of the paraventricular nucleus (PVN), the anterior hypothalamic area (AHA), dorsomedial hypothalamus (DMH), and lateral hypothalamic area (LHA) were acquired with the same exposure time, brightness, and contrast on Zeiss Axioimager Z1 with Axiovision 4.5 software (Oberkochen, Germany). The 10 \times magnification images were quantified using ImageJ (Sugrue et al., 2010). To ensure that hypophysiotropic neurons in the PVN were the focus of *Trh* mRNA measurements, quantification of brain sections was limited to the medial portion of the PVN, which has been described in detail recently (Kádár et al., 2010). Positive pixels per unit area (pixel density) was calculated on both sides of the PVN by highlighting the medial portion of the PVN excluding the periventricular zone and subtracting background from areas not expressing *Trh*. Pixel density per mouse was averaged from a minimum of three sections of the medial portion of the PVN.

Statistical Analyses

Data are presented as sample means \pm SEM. Sample numbers range from n = 3 to n = 40 and are stated specifically within the text or figure legends. The text or figure legends highlight how differences between groups were measured, which includes unpaired t test, one-way ANOVA with Tukey's multiple comparison post hoc test, repeated-measures two-way ANOVA with Bonferroni post hoc test, and two-way ANOVA with Bonferroni post hoc test.

SUPPLEMENTAL INFORMATION

Supplemental Information includes four figures, Supplemental Experimental Procedures, and Supplemental References and can be found with this article online at doi:10.1016/j.cmet.2011.10.009.

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