



# Stressing the metabolic powers of fibroblast growth factor 21

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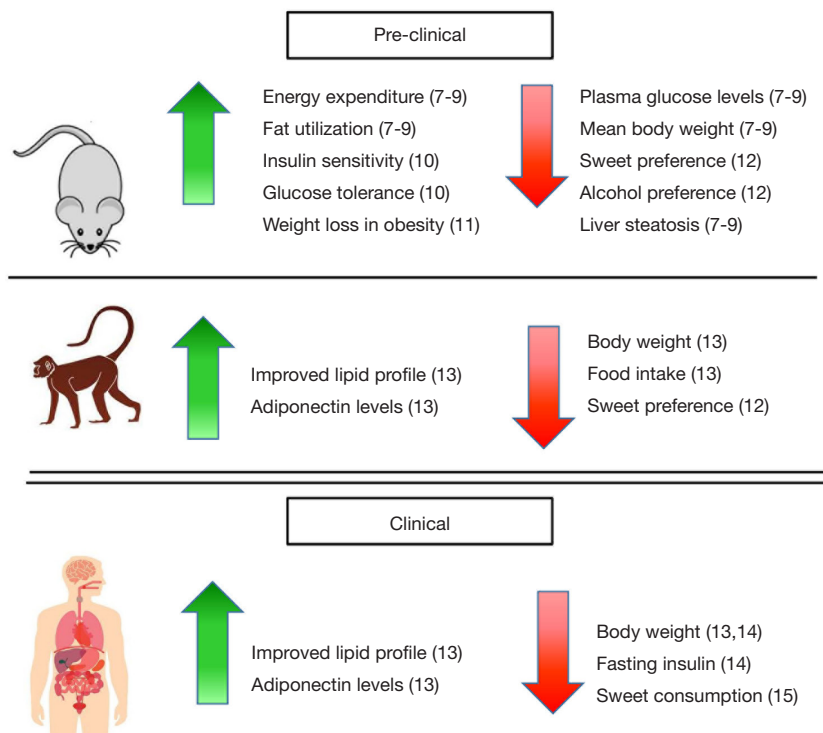
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The global prevalence of obesity has continued to rapidly increase over the last few decades. As well as the range of metabolic disorders associated with obesity, diseases like diabetes, which was the second leading cause of death related to obesity in 2015, all add to the global burden of disease (1). Currently, the options for treatment of obesity and its associated disorders are limited. Soluble pro-inflammatory cytokines including interleukin-6 and granulocyte macrophage colony stimulating factor have exhibited profound beneficial metabolic effects. These include promotion of fat oxidation and insulin sensitivity and reduction in food intake and body weight, respectively (2,3). The adipokines adiponectin and leptin may also have a positive effect on body weight regulation (4,5). Additionally, overexpression of interleukin-10 protected mice from diet induced inflammation and insulin resistance in skeletal muscle (6). Also emerging as an appropriate therapeutic agent is the soluble fibroblast growth factor 21 (FGF21) (*Figure 1*) (16). Early investigations in which FGF21 was administered to mouse models of obesity and diabetes revealed that it reduced plasma glucose levels, lowered mean body weight and reversed hepatic steatosis (7-9). In monkeys, it has been shown that FGF21 decreases sweet preference via dopamine signalling (12). Soberg *et al.* highlighted that in humans, the preferences for sweet tasting substances is determined by genetic variation of the *fgf21* gene (15). While native FGF21 has an unfavorable pharmacokinetic profile, a number of pharmaceutical companies have developed FGF21 analogues that appear more suitable for use in humans (16). Published data

from clinical trials administering two FGF21 mimetics, LY2405319 and PF-05231023, to patients with type 2 diabetes have demonstrated that this approach is safe (13,14). LY2405319 was found to significantly reduce body weight and fasting insulin levels of patients receiving the treatment compared to placebo after 28 days but only showed a dose dependent trend of lowering fasting glucose levels (14). After 25 days of treatment with PF-05231023, patients had a significant reduction in body weight compared to the placebo group. Patients receiving PF-05231023 also showed a trend towards lower glucose and insulin levels compared to subjects receiving placebo although they were not statistically different (13). Interestingly, in a larger clinical trial of PF-05231023 involving obese subjects with or without type 2 diabetes, no significant reduction in body weight could be observed (17). As clinical trials using these and other newly developed FGF21 mimetics continue, an alternative therapeutic approach aimed at augmenting endogenous FGF21 levels and/or activity is being pursued (16). Many studies have demonstrated that FGF21 expression can be upregulated in multiple tissues in response to stress (18). Sources of stress, including amino acid deprivation and endoplasmic reticulum stress, can initiate the integrated stress response (ISR) pathway by activating one of four kinases, PKR-like ER kinase (PERK), double-stranded RNA-dependent protein kinase (PKR), heme-regulated eIF2 $\alpha$  kinase (HRI) or general control non-depressible 2 (GCN2) (*Figure 2*). These kinases can then phosphorylate eIF2 $\alpha$ , a central component of ISR signalling, which leads to upregulated activity of the transcription factor activating



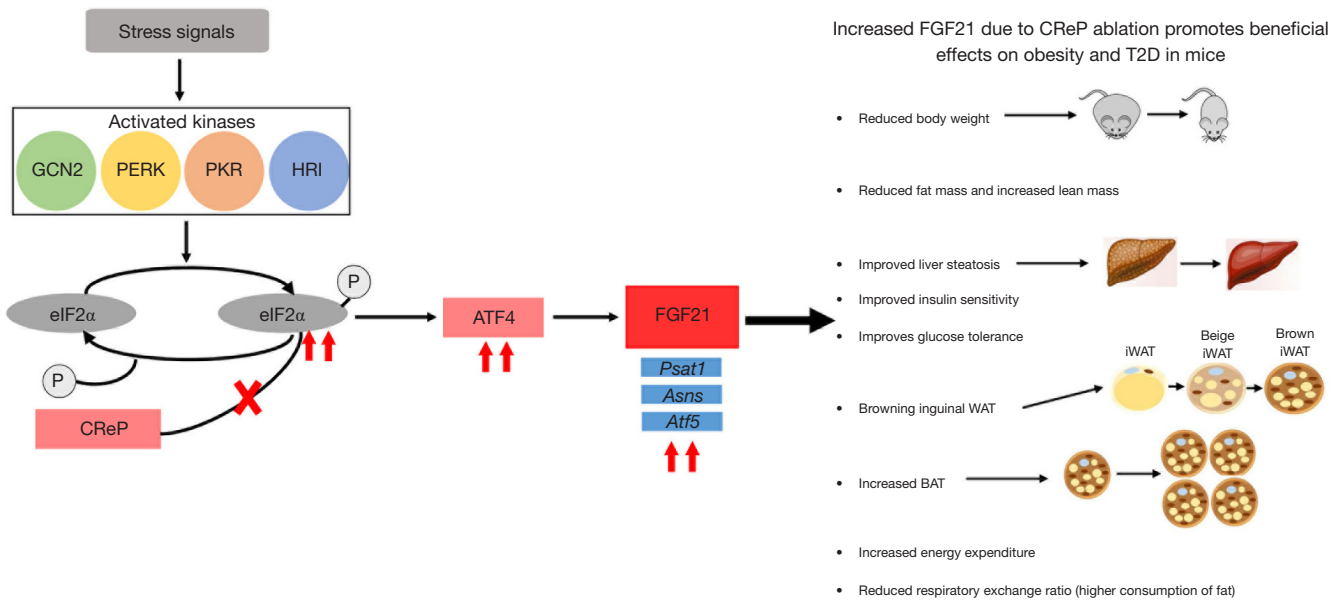
**Figure 1** The metabolic benefits of FGF21 and FGF21 mimetics in mice, non-human primates and humans. FGF21, fibroblast growth factor 21 (7-15).

transcription factor 4 (ATF4) and subsequent increased transcription of the *Fgf21* gene (18).

In a recent publication in *Hepatology*, Xu and colleagues generated a mouse model,  $\text{CReP}^{\text{LKO}}$ , in which constitutive repressor of  $\text{eIF2}\alpha$  phosphorylation (CReP) was specifically knocked out in the liver (19). CReP is a subunit of the  $\text{eIF2}\alpha$  phosphatase complex that is involved in the downregulation of  $\text{eIF2}\alpha$  phosphorylation. As expected, a hepatic knockout of CReP resulted in increased levels of phosphorylated  $\text{eIF2}\alpha$  in the liver of  $\text{CReP}^{\text{LKO}}$  mice compared to wild-type (WT) (Figure 2). The  $\text{CReP}^{\text{LKO}}$  mice were grossly normal in appearance with minimal impact on the protein synthetic and secretory function of the liver observed. Their liver to body weight ratio was 14% lower than WT. The  $\text{CReP}^{\text{LKO}}$  mice contained strikingly brown-coloured subcutaneous inguinal white adipose tissue (iWAT). Prior to being placed on a high-fat diet (HFD), the  $\text{CReP}^{\text{LKO}}$  mice had 17% lower body weight compared to WT, a difference that grew to 32% after both groups of mice were placed on an HFD for 12 weeks. After 12 weeks on an HFD, the  $\text{CReP}^{\text{LKO}}$  mice had a significantly lower fat mass percentage than the WT, paralleled by a higher lean mass percentage, which suggests a reduction of adiposity. In comparison to the WT mice, the

$\text{CReP}^{\text{LKO}}$  mice exhibited a higher energy expenditure but lower respiratory exchange ratio both before and after being on an HFD. This is indicative of a higher consumption of fat as an energy source in the  $\text{CReP}^{\text{LKO}}$  mice consistent with the brown-coloured iWAT found in them. Insulin sensitivity and glucose homeostasis was improved in the  $\text{CReP}^{\text{LKO}}$  mice both before and after being on an HFD. This allowed the mice to maintain normal glucose and serum insulin levels while on an HFD for 12 weeks. The observed increases in energy expenditure and insulin sensitivity with elevated FGF21 levels are consistent with those of previous studies (10,11). The livers of the  $\text{CReP}^{\text{LKO}}$  mice were in a noticeably healthier condition than the WT mice, which developed clear signs of hepatic steatosis, after 12 weeks on an HFD (Figure 2).

These phenotypical differences between the  $\text{CReP}^{\text{LKO}}$  and WT mice were underpinned by key differences at the molecular level between the two types of mice. With the higher level of  $\text{eIF2}\alpha$  phosphorylation in the livers of  $\text{CReP}^{\text{LKO}}$  mice, there was also an elevated level of the downstream ISR signalling proteins, ATF4 and ATF5. The team also detected higher mRNA levels of genes including *Atf5*, *Asns*, and *Psat1*, which are regulated by the transcription factors ATF4 and ATF5. The aforementioned



**Figure 2** ISR activation leading to eIF2 phosphorylation. ATF4 activation and FGF21 induction promotes metabolic benefits. GCN2, general control non-depressible 2; PERK, PKR-like ER kinase; PKR, double-stranded RNA-dependent protein kinase; HRI, heme-regulated eIF2α kinase; ISR, integrated stress response; FGF21, fibroblast growth factor 21; ATF4, activating transcription factor 4.

browning of iWAT in CReP<sup>LKO</sup> mice was accompanied expectantly with increased expression of brown adipocyte markers *Ucp1*, *Dio2* and *Elovl3* in WAT and brown adipose tissue (BAT). Without the degree of hepatic steatosis observed in the WT mice, there were lower levels of proteolytic activation of SREBP-1c and expression of its target lipogenic genes in the livers of CReP<sup>LKO</sup> mice as expected. The greater insulin sensitivity in the CReP<sup>LKO</sup> mice corresponded with higher insulin-mediated AKT phosphorylation in their livers and WAT compared to WT. Of most interest were the higher levels of *FGF21* mRNA in the liver and FGF21 protein in the serum of CReP<sup>LKO</sup> mice, which was maintained throughout HFD feeding. These levels were in fact similar to the levels seen in fasted WT mice. When CReP<sup>LKO</sup> mice were fasted, the FGF21 levels increased further from their already elevated level.

To further elaborate on their findings, Xu *et al.* used two further mouse models, protein phosphatase 1 regulatory subunit 15b<sup>loxP</sup> (*Ppp1r15b<sup>loxP</sup>*) and double knockout (DKO) mice. By injecting an AAV-Cre vector into *Ppp1r15b<sup>loxP</sup>* mice, Xu *et al.* were able to ablate CReP expression in the liver of adult mice. Following the injection of AAV-Cre, the adult *Ppp1r15b<sup>loxP</sup>* mice exhibited many features observed in the CReP<sup>LKO</sup> mice including, upregulation of *Fgf21*, *Asns* and *Psat1* in the liver, no elevation of ER stress markers and

increased expression of *Ucp1* and *Elovl3* in WAT and BAT. Body weight and blood glucose levels in *Ppp1r15b<sup>loxP</sup>* mice were reduced by 6% and 20%, respectively, within three days of the AAV-Cre injection. *Ppp1r15b<sup>loxP</sup>* mice that had been on an HFD also saw a reduction in their body weight and blood glucose levels by 20% and 45%, respectively, two weeks after being injected with AAV-Cre. Improvements in insulin sensitivity and liver steatosis, as well as a decrease in fat mass were also observed in *Ppp1r15b<sup>loxP</sup>* mice on an HFD following the AAV-Cre injection. The results observed in the AAV-Cre injected *Ppp1r15b<sup>loxP</sup>* mice suggest that activation of ISR in the liver is a viable target for the treatment of obesity and is capable of providing multiple beneficial metabolic effects.

The DKO mouse was generated by crossing CReP<sup>LKO</sup> mice with *Fgf21<sup>loxP</sup>* mice, which do not express FGF21. All the beneficial effects of CReP ablation seen in CReP<sup>LKO</sup> mice were negated by the loss of FGF21 in the DKO mice, which were more similar to WT mice. There was no browning of WAT seen in DKO mice with *Ucp1* and *Elovl3* mRNA levels similar to those in WT mice. The energy expenditure, respiratory exchange ratio, insulin sensitivity and glucose homeostasis in the DKO mice were also all similar to WT mice. The generation of the DKO mice by Xu *et al.* eloquently demonstrated that the all the benefits seen in the CReP<sup>LKO</sup> mice were dependent on presence of FGF21.

The protective benefits of ISR signalling and the central role that eIF2 $\alpha$  plays clearly shown by Xu *et al.* mirrors the findings of Scheuner and colleagues who studied a mouse model in which eIF2 $\alpha$  could not be phosphorylated (20,21). Eif2s1<sup>tm1Rjk</sup> mice were generated by using gene targeting to mutate the Ser51 phosphorylation site of eIF2 $\alpha$  to prevent its phosphorylation (20). Homozygous Eif2s1<sup>tm1Rjk</sup> mice died within 18 hours of birth due to hypoglycemia and were found to have only 20% of the serum insulin level of WT mice (20). Heterozygous Eif2s1<sup>+ / tm1Rjk</sup> mice on an HFD gained significantly more body weight, had reduced energy expenditure and a higher respiratory exchange ratio in comparison to WT mice on the same diet (21). After 5 weeks on an HFD, heterozygous Eif2s1<sup>+ / tm1Rjk</sup> mice developed severe glucose intolerance compared to WT mice on an HFD in which only a minor change in glucose intolerance was observed in the same timeframe (21). The work of Xu *et al.* and Scheuner *et al.* highlight the importance of the ISR pathway and provide a possible therapeutic target in eIF2 $\alpha$  to protect against the metabolic impact of obesity via the upregulation of endogenous FGF21. Drugs that inhibit eIF2 $\alpha$  phosphatases like CReP include salubrinal, sephrin1, guanabenz and nelfinavir. These drugs are available but it is important to note that targeting the ISR pathway does not have uniform consequences in all cell types and may result in unwanted side effects (22).

In addition to the ISR pathway there are alternative pathways that have been demonstrated to be involved in the upregulation of endogenous FGF21 expression. Metformin, for example, was found to increase the level of FGF21 expression in obese mice (23). Increased liver expression of FGF21 was observed in Sprague-Dawley rats that had received a combination of docosahexaenoic acid and triiodothyronine (24). Our own work has highlighted that when haematopoietic cells express TNFSF14, there is a concomitant increase in hepatic FGF21 levels (25). More importantly, the increased FGF21 expression aligned with an improved metabolic profile (25).

Gene therapy is also a novel avenue being pursued to elevate the endogenous level of FGF21 to treat obesity. Jimenez and colleagues used a variety of adeno-associated virus vectors to overexpress FGF21 in the liver of C57BL/6 mice on an HFD, the liver or epididymal WAT of ob/ob mice (a model of obesity) and skeletal muscle of WT mice (26). Gene therapy induced FGF21 overproduction in each mouse model and irrespective of the tissue source of increased FGF21, FGF21 prevented body weight gains that were observed in control mice treated with null vectors (26).

Overall, there have been exciting developments in the search for viable therapeutic options for the treatment of obesity and the metabolic disorders related to it. The work of Xu *et al.* clearly demonstrates the protective benefits of engaging the ISR pathway against the metabolic dysfunction of obesity. It represents another piece of evidence that highlights the beneficial role that FGF21 can play in the treatment of metabolic diseases. However, optimism about a treatment must be tempered with caution as there still remains much to be elucidated about how these pathways interact within the body and what impact long-term supra-physiological levels of FGF21 may have on the body.

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