

Targeting energy expenditure via fuel switching and beyond

J. G. Geisler

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Abstract Since over-nutrition accelerates the development of obesity, progression to type 2 diabetes, and the associated co-morbidity and mortality, there has been a keen interest in therapeutic interventions targeting mechanisms that may curb appetite, increase energy expenditure or at least attenuate insulin resistance. Over the past decade, numerous peri-mitochondrial targets in the de novo lipid synthesis pathway have been linked to an increase in energy expenditure and the drug development industry has pursued the gene products involved as candidates to develop drugs against. The basis of this link, and specifically the premise that lowering tissue and cellular malonyl-CoA can increase energy expenditure, is scrutinised here. The argument presented is that fuel switching as effected by changes in cellular malonyl-CoA concentrations will not trigger the mitochondria to increase energy expenditure because: (1) an increase in beta-oxidation by lowering respiratory exchange ratio (indicative of the metabolic fuel consumed) does not equal an increase in energy expenditure (how rapidly fuel is consumed); (2) the ATP:oxygen ratios (i.e. ATP energy made:oxygen required for the reaction) are similar when metabolising lipids (2.8) vs glucose (3.0); (3) substrate availability (NEFA) does not drive energy expenditure in vivo; and (4) the availability of ADP in the mitochondrial matrix determines the rate of energy expenditure, not the availability of fuel to enter the mitochondrial matrix. To increase mitochondrial energy expenditure, work must be done (exercise) and/or the mitochondrial proton leak must be enhanced, both of which

increase availability of ADP. In fact, despite the historic taboo of chemical uncoupling, this mechanism validated in humans is closest on task to increasing whole-body energy expenditure. Chemical uncoupling mimics the naturally occurring phenomenon of proton leak, accelerating the metabolism of glucose and lipids. However, it is completely non-genomic (i.e. the target is a location, not a gene product) and is not associated with addiction or mood alterations common to satiety agents. A significant hurdle for drug development is to discover a safe mitochondrial uncoupler and to formulate it potentially as a pro-drug and/or oral pump, to avoid the issue of overdosing experienced in the 1930s. The potential therapeutic impact of such a compound for an over-nutritioned patient population could be profound. If effective, the mitochondrial uncoupler mechanism could resolve many of the associated diseases such as type 2 diabetes, hypertension, obesity, depression, sleep apnoea, non-alcoholic steatohepatitis, insulin resistance and hyperlipidaemia, therefore becoming a ‘disease-modifying therapy’.

Keywords β -Oxidation · Brown adipose tissue · Chemical uncoupling · Disease-modifying therapy · Energy expenditure · Insulin resistance · Malonyl-CoA · Mitochondria · Obesity · Type 2 diabetes

Abbreviations

ACC	Acetyl-coenzyme A carboxylase
BAT	Brown adipose tissue
BMR	Basal metabolic rate
CPT1A	Carnitine palmitoyltransferase-1A
KO	Knockout
P:O	ATP:oxygen
RER	Respiratory exchange ratio
UCP1	Uncoupling protein 1
$\dot{V}O_2$	Volume of oxygen consumed

J. G. Geisler (✉)
Johnson & Johnson Pharmaceutical Research and Development,
L.L.C., Metabolic Disease, Drug Discovery,
Welsh & McKean Roads,
Spring House, PA 19477, USA
e-mail: Jgeisle1@its.jnj.com

The premise

In the last decade, seminal papers have been published in which gene deletion of targets peripheral to the mitochondria resulted in increased beta-oxidation and weight loss. For example, stearoyl-coenzyme A desaturase 1 null mutation was reported to protect against adiposity and increased fatty acid oxidation [1]. Subsequently, it was understood that this gene also had an essential role within the dermis and the animals were hypothermic at room temperature [2]. Similarly, diacylglycerol *O*-acyltransferase-2 has a role in providing a skin barrier [3]. These enzymes, as well as acetyl-coenzyme A carboxylase (ACC) 1 and 2 [4], fatty acid synthetase [5] and mitochondrial glycerol-3-phosphate acyltransferase-1 [6], are key enzymes in the de novo synthesis of triacylglycerol. Of central interest are ACC1 and 2, since they directly generate malonyl-CoA, which plays a dual role in the liver [4]. Malonyl-CoA donates the first two carbons for de novo synthesis of palmitate and binds to carnitine palmitoyltransferase-1a (CPT1A) as an allosteric inhibitor of fatty acid transport into the mitochondrial matrix. This second role interested pharmacologists because it was believed that lowering cellular malonyl-CoA would result in increased beta-oxidation of lipids and increased energy expenditure in the mitochondria. Why did we believe this hypothesis? First, malonyl-CoA allosterically inhibits CPT1A. Second, malonyl-CoA levels drop in the fasted state and subsequently beta-oxidation of fatty acids begins. Third, the knockout (KO) of *Acc2* (also known as *Acacb*) resulted in increased energy expenditure as measured by the volume of oxygen consumed $\dot{V}O_2$ [7, 8]. It has also been shown that inhibition of the downstream genes *Scd1* and *Dgat2* also lowers *Acc1* (also known as *Acaca*) and *Acc2* expression by a feedback mechanism [9–11]. These results supported drug research targeted approaches to lower malonyl-CoA. However, there are several problems with this premise, including the unusual phenotype of *Acc2* KO mice. First, it is unlikely that lowering malonyl-CoA's allosteric inhibition of CPT1A in the fed state will result in enhanced oxidation of lipids, with concomitant weight reductions and improved insulin sensitivity. Second, the first *Acc2* KO animals did not show a switching of fuels (i.e. change in respiratory exchange ratio [RER]), but rather an increase in $\dot{V}O_2$, which was not predicted by bioenergetics or evident with a dual *Acc1* and *Acc2* inhibitor in rats [12]. Recently, the *Acc2* KO model was independently generated and resulted in an opposite phenotype [13]. The regenerated *Acc2* KO animals had lower malonyl-CoA levels, lower RER and increased fatty acid oxidation, but there was no change in $\dot{V}O_2$, adiposity and body weight, or more importantly no improvements in glycaemic control during a glucose tolerance test on a low- or high-fat diet [13]. The

premise that purely switching metabolism from glucose to lipid oxidation by lowering malonyl-CoA will result in increased energy expenditure and be useful in treatment of metabolic disease is clearly unsupported. In fact, it may exacerbate the phenotype.

Further down the de novo lipid synthesis pathway, interesting observations have been made in the phenotype of mitochondrial *Gpat1* (also known as *Gpam*) KO mice on a high-fat diet [14]. Glycerol-3-phosphate acyltransferase-1 catalyses the initial and rate-limiting step for addition of the first long-chain acyl-CoA on to the glycerol-3-phosphate backbone to partition lipids towards synthesis of triacylglycerol and away from degradative pathways. The long-term consequence of placing the KO phenotype on a high-fat diet was increased insulin resistance. *Gpat1* KO mice had increased hepatic acyl-carnitines, a measurement of the abundance of partially processed fatty acids leaving the mitochondria and diffusing into the plasma compartment. The conclusion drawn from the *Gpat1* KO studies was that 'the amount of acyl-CoA exceeded the capacity of the mitochondrial oxidation pathway' [14]. The effect may be real, but the conclusion is inaccurate. Instead, it is possible that lowering malonyl-CoA without an increase in energy expenditure was the cause of increased acyl-CoA levels. It is unlikely that the mitochondrial machinery was overwhelmed by an excess of lipids to metabolise. It is, rather, more likely that the mitochondria could not metabolise more lipids because ATP levels were sufficient and ADP levels insufficient. Indeed, if these mice had been exercising or administered a mitochondrial uncoupler such as 2,4-dinitrophenol, it is possible that the mitochondria would have metabolised the back-log of acyl-CoAs, thus restoring sufficient levels of ADP.

Energy expenditure is dependent on ADP availability and not substrate availability

A widely held but false presumption that seems to act as a dividing line between the mitochondrial bioenergetic and the diabetes/obesity communities is that increasing substrate into the mitochondria will increase energy expenditure. In contrast to experiments on isolated mitochondria or in vitro studies [15], the infusion of lipids in humans did not change energy expenditure [16], but did increase insulin resistance [17]. If the allosteric block (malonyl-CoA) is removed from CPT1A, fatty acids will enter the mitochondria, but do not become oxidised until energy is needed. The ability to get into the mitochondria is a necessary condition, but not in itself sufficient for the oxidation of lipids. The metabolism of lipids is dependent on the availability of ADP [18], not on whether fatty acids can get into the mitochondria. If most of the ADP is locked up

as ATP, the condition typically found in the fed state, then intra-mitochondrial lipids will not be used. Mitochondrial toxicity or insulin resistance may arise if all normal functions (e.g. oxidative phosphorylation, electron transport, citric acid cycle) are to continue against a background of lipid overabundance [14]. Although oxidation of NEFA can be induced reproducibly in vitro [15], there is no evidence that increasing substrate in vivo will increase energy expenditure. Increasing substrate in humans has been shown to increase insulin resistance and fat storage [17, 19]. The function of CPT1A is not to block or modulate the rate of fatty acid oxidation, but rather to regulate lipid substrate availability into the inner mitochondrial matrix, thus governing against an overabundance of intra-mitochondrial fatty acids. The use of an in vitro beta-oxidation assay to select targets and compounds in order to suggest that the same effect occurs in vivo can potentially lead to incorrect conclusions. Evidence of increased energy expenditure must be demonstrated experimentally (e.g. $\dot{V}O_2$, tracers) in an intact animal.

The phenomenon of adaptive increases in energy expenditure in response to energy intake and weight gain is evident in rodents [20], but perhaps to a considerable less degree in humans. It has been clearly demonstrated in a head-to-head comparison of the *Ucp1* KO mouse with a wild-type mouse during which both were fed a high-fat diet and housed at thermoneutrality, i.e. $\sim 30^\circ\text{C}$ (82°F), when non-shivering thermogenesis is turned off. Both animals gained considerable weight relative to when housed at room temperature (25°C) [21], but at $\sim 30^\circ\text{C}$ the wild-type mouse gained far less weight than its *Ucp1* KO counterpart [22]. At this temperature, the sympathetic nervous system should not stimulate proton leak via uncoupling protein 1 (UCP1) in a fully ‘clothed mouse’ [23] and yet a compensatory effect of resistance to weight gain due to the high energy diet was still observed [24]. It is unlikely that such a mechanism will provide a meaningful impact in humans consuming high-fat diets (i.e. a diet-induced increase in activity and/or thermogenesis), since (1) the mechanism appears to be mediated by UCP1, (2) brown adipose tissue (BAT) mass is considerably lower and (3), more importantly, we have a pandemic of obesity.

Fuel switching does not increase energy expenditure

A second possible misconception is that switching fuels will increase oxidation and this in turn will increase energy expenditure. When malonyl-CoA is reduced, the NADH/FADH₂ needed to drive the electron transport chain and subsequently oxidative phosphorylation is forced to be derived from NEFA by beta-oxidation and from the remaining acetyl-CoAs by the citric acid cycle vs metabolism of glucose (Fig. 1). However, regardless of whether the source of

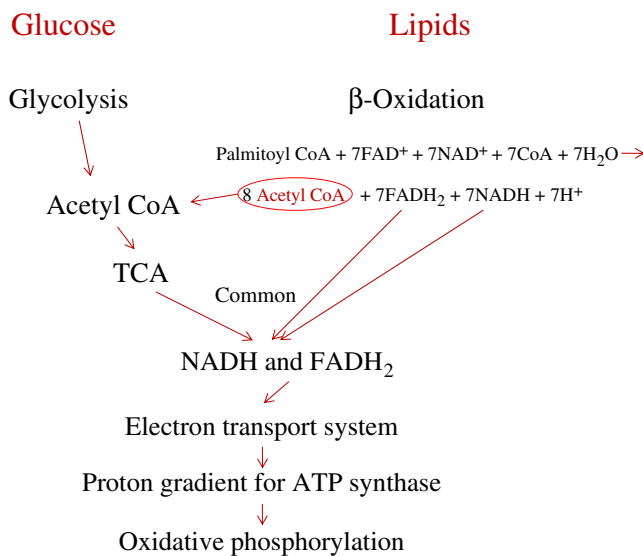


Fig. 1 Metabolism of glucose and lipids is the source of the proton gradient ($\Delta\tilde{u}_{H^+}$). NADH and FADH₂ derived from the metabolism of glucose and lipids provide the protons used to maintain the proton gradient across the mitochondrial matrix

NADH/FADH₂ is from glucose or lipid, the ATP:oxygen (P:O) ratio (ratio of ATP produced to the amount of oxygen required and consumed for the reaction) remains roughly equivalent (Fig. 2). The metabolism of 1 mole of glucose consumes 12 moles of oxygen and yields 36 moles of ATP. Therefore, the P:O ratio for glucose is 36/12=3. Metabolism of 1 mole of palmitate requires 45.7 moles of O₂ ($16/0.7 \times 2$) and yields 129 moles of ATP. Therefore, the P:O ratio is 129/45.7=2.8 [25]. This implies that no more energy is gained or lost simply by switching fuel, and therefore the mitochondria’s energy balance remains equal until the rate of oxygen consumption changes (energy expenditure). This was clearly demonstrated in rats administered a dual ACC1 and ACC2 inhibitor or in the recent *Acc2* KO model, where RER was constitutively lowered (as expected), but $\dot{V}O_2$ consumed remained unchanged [12, 13]. The reason why $\dot{V}O_2$ was unaltered is simply that switching fuels does not motivate the mitochondria to increase energy expenditure. The P:O ratios are equivalent, and therefore ATP levels or mitochondrial

Glucose	Lipids (palmitate)
$C_6H_{12}O_6 + 6O_2 \rightarrow 6CO_2 + 6H_2O$	$C_{16}H_{32}O_2 + 23O_2 \rightarrow 16CO_2 + 16H_2O$
Yield: 36 ATP	Yield: 129 ATP
P/O	P/O
$\frac{36 \text{ ATP}}{12 \text{ O}} = 3.0$	$\frac{129 \text{ ATP}}{46 \text{ O}} = 2.8$
← Fuel switch →	

Fig. 2 ATP:oxygen ratios for different fuels. The ratio of energy yield (ATP) to oxygen required for glucose vs lipid metabolism are roughly equal. Palmitate generates more ATP per molecule, but requires more oxygen to metabolise

membrane potential remain in balance. It should be noted that although there is a 6.67% difference between glucose and palmitate P:O ratio, it appeared insignificant with regard to providing an advantage in the constitutive fatty acid oxidation state for body weight, glycaemic control or adiposity in *Acc2* KO animals under high-fat diet conditions.

Animal models with dermis issues

The heat generated from brown fat (via UCP1) in rodents may also be linked to the possible misinterpretation of the metabolic phenotypes of mouse mutations and compounds that have been reported to reduce body weight in rodents. In rodents, BAT has the capacity to increase energy expenditure 60% over basal metabolic rate (BMR) through non-shivering thermogenesis [26]. If the function or heat retention capacity of mouse skin is compromised by changing the lipid content, non-shivering thermogenesis is engaged to maintain core body temperature via the sympathetic nervous system [2]. This creates the illusion, on translation to humans when characterising a new target, that the KO phenotype resulted in increased whole body energy expenditure and that the protein may be a suitable target for the development of drugs to treat metabolic disease. Before claiming increased energy expenditure, it should be tested whether the KO animals or drug candidates can reduce body weight at 30°C (thermoneutrality) or when animal models are crossed on to a *Ucp1* KO background to eliminate weight reductions due solely to non-shivering thermogenesis [22]. Ideally, compounds should be tested in *Ucp1* KO animals at 30°C to simultaneously eliminate shivering, non-shivering and diet-induced thermogenesis [27]. If greater energy expenditure and weight reductions are still present in the absence of UCP1, then the target or compound tested may well translate into weight loss in humans.

Breaking the taboo of chemical uncoupling

A significant research effort was undertaken in the 1990s after publication of a seminal paper connecting the phenomenon of non-shivering thermogenesis in BAT to UCP1 [28]. There was a keen interest in developing UCP1 drugs as agonists capable of harnessing the potential of UCP1 to achieve weight loss via energy expenditure mechanisms. However, it was quickly realised that UCP1 is restricted to BAT and adult humans appeared to no longer have this tissue. Only UCP1 has uncoupling properties [29]. These findings led to a steep decline in research on drugs targeting energy expenditure mechanisms and efforts were channelled towards anorectic mechanisms. Recently, BAT was found in humans along the vertebra and shown to

be induced upon exposure to cold, presumably also by the sympathetic nervous system [30, 31]. Expectations of significant weight loss mediated by human BAT should be treated with some caution if the intention is to use the existing tissue or to expand upon the mass of this tissue. One reason for such caution is that humans have proportionally less BAT per body weight than rodents. Current positron emission tomography-computed tomography (PET-CT) scans estimate BAT mass in humans to total ~13 g (0.02% of body weight for a person weighing 70 kg) [32], whereas mouse BAT mass is ~400 mg for 40 g body weight (1%) (J.G. Geisler, unpublished results). Therefore, humans have approximately 50-fold less BAT than mice. More importantly, since humans are thermoneutral [22], it is unlikely that UCP1 could be triggered to uncouple and provide meaningful weight loss at room temperature.

Although there are some 30,000 genes in the genome [33], the most effective means of increasing energy expenditure in humans may be non-genomic (i.e. independent of a gene or protein target), namely chemical uncoupling (i.e. location of the mitochondrial matrix would be the target). There has been renewed interest in chemical uncoupling due to a better understanding of mitochondrial function [24, 34] and the challenges of anorectic agents that target reward pathways [35, 36]. Chemical uncouplers mimic the function of UCP1 and the naturally occurring proton leak, which accounts for ~20 to 25% of the energy lost in liver and ~50% of that lost in muscle, i.e. roughly 25% of total BMR [37, 38]. Instead of a protein channel for the proton to enter through (e.g. UCP1), the molecule acts as its own proton transporter into the matrix from the cytosol. As a mechanism for effecting weight loss, chemical uncouplers are precisely on task for causing the mitochondria to increase energy expenditure and achieve this by metabolising glucose and NEFA in the process (Fig. 1). The mechanism works as follows. A weakly acidic uncoupler holds a proton that dissociates upon entering the basic environment within the mitochondrial matrix. As a proton has been lost, the molecule leaves the matrix as an anion and, upon returning to the acidic cytosolic environment, becomes reprotonated to a cation, upon which it again returns to the matrix to drop off another proton (Fig. 3). The cumulative effect of such a cycle is a reduction of the mitochondrial membrane potential and a lowering of the proton-motive-force (Δp) that would be used to allow a proton to travel through ATP synthase, causing a rotation and subsequent phosphorylation of ADP [39]. Because of this, ATP levels drop and the mitochondria instantly responds to re-establish the gradient by accelerating utilisation of NADH and FADH₂. Oxidation is sustained in the presence of an uncoupler because the mitochondrial membrane potential cannot be restored until the drug is cleared/metabolised. These electron transport system sub-

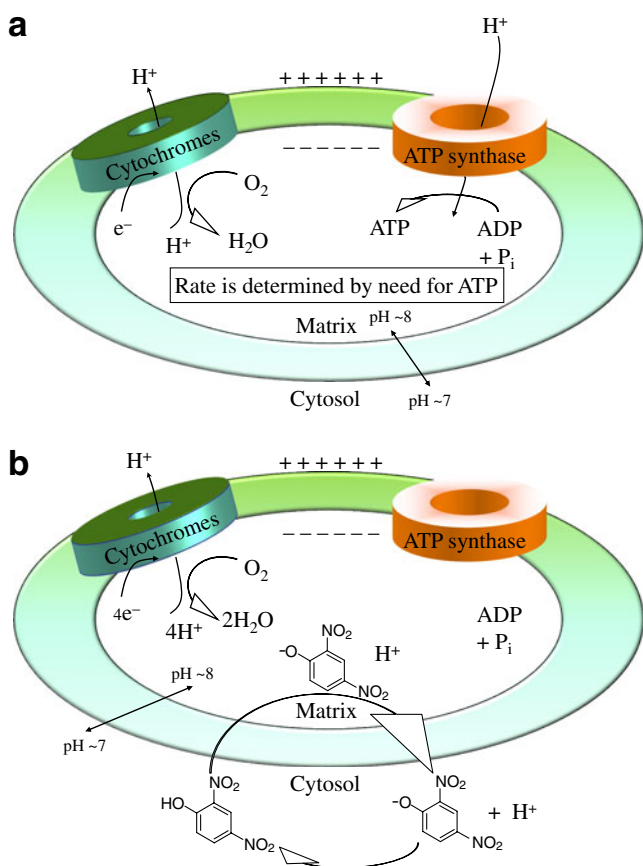


Fig. 3 Coupling vs uncoupling. Maintaining the proton gradient across the mitochondrial matrix and cytosol involves the pumping of protons out of the matrix via cytochromes I, III and IV. **a** The coupling of a proton transfer to the synthesis of ATP is a result of a proton returning through ATP synthase, causing a rotation and subsequent phosphorylation of ADP, thereby yielding an ATP molecule. **b** This mechanism is circumvented in the case of chemical uncoupling (i.e. entry of protons without phosphorylation to produce ATP). The proton transfer into the matrix is on a carrier, a weak acid molecule (e.g. 2,4-dinitrophenol or carbonyl cyanide *p*-trifluoromethoxyphenylhydrazone). When the weak acid (cation) enters the basic environment of the matrix, a proton is dissociated and the molecule, now an anion, returns to the cytosol to become reprotonated into a cation and start the cycle over again. All mitochondrial systems remain functional, but are accelerated

strates come from two sources, the metabolism of glucose and that of lipids. Energy expenditure is increased as monitored by an increase in oxygen consumed; however, the RER is roughly unchanged and therefore a mixed fuel is consumed. This mechanism motivates the mitochondria towards anti-obesity and anti-diabetes because the proton electrochemical gradient ($\Delta\tilde{\mu}_{H^+}$) is out of balance. This is fundamentally different from malonyl-CoA-lowering mechanisms (Fig. 4). It is important to note that for this mechanism to be efficient, it is critical that all components of the electron transport system, including ATP synthase, remain completely functional. A pure chemical uncoupler is not an inhibitor, but an enhancer of respiration.

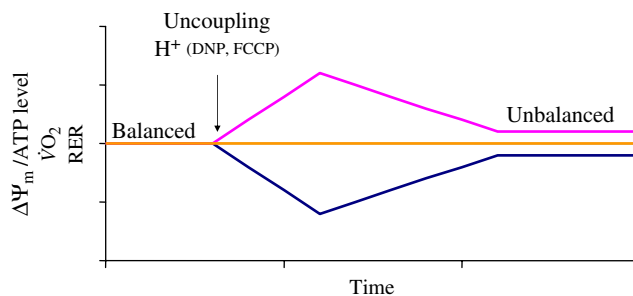


Fig. 4 Dynamics of energy compensation with chemical uncoupling in the fed state. Administration of a chemical uncoupler causes a drop in the mitochondrial membrane potential ($\Delta\Psi_m$) and ATP levels (blue line). The cell responds by increasing the rate of oxygen consumption ($\dot{V}O_2$ [pink line] or energy expenditure) to generate more NADH and FADH₂ as a source of protons (when oxidised back to NAD⁺ and FAD) as a mechanism to re-establish the electrochemical gradient ($\Delta\tilde{\mu}_{H^+}$). Since NADH and FADH₂ come from the metabolism of glucose and lipids, the RER (orange line) is primarily unchanged (high doses shift RER towards lipids). In the presence of a chemical uncoupler, energy expenditure is constitutively enhanced since the $\Delta\Psi_m$ is out of balance

Drug safety is a primary concern for the development of chemical uncoupling as a therapeutic approach for metabolic disease. Historical data from the 1930s demonstrate the abuse potential for uncoupler-based weight-loss agents, although 2,4-dinitrophenol was not addictive [40]. Beyond the benefits of weight loss ‘predominantly from the hips and abdomen’ [40], uncouplers reduce reactive oxygen species [41], a phenomenon that has been associated with improved infarct volume during ischaemia [42], longevity and ageing [43]. Although chemical uncoupling was proven to achieve weight loss in humans during the 1930s, the science of pharmacology and the challenges of drug development have changed dramatically over the past 80 years. It would need to be demonstrated that all vital functions are uncompromised with chemical uncoupling, in particular the heart in a setting of increased BMR. Another potential risk of chemical uncoupling is increases in core body temperature at high doses, although at lower doses the heat is usually dissipated with no change in temperature [44, 45]. Cataracts have been reported in some patients treated with 2,4-dinitrophenol, but this was not replicated in more recent animal models [46] and may be a property of 2,4-dinitrophenol vs general uncouplers. A key question is to what degree does energy expenditure have to be increased to achieve meaningful weight loss? According to calculations from a study conducted in 1935, a 10% increase in BMR resulted in weight loss of ~0.3 to 0.45 kg/week [44]. If the calculations are accurate and a less aggressive approach was taken today, then a 5% increase in BMR should result in half that weight loss, but still provide annual weight reductions of 10% or ~12 kg (26 lb) for patients with BMI over 30 kg/m². Interestingly, there were,

to the author's knowledge, no reports of a compensatory increase in food consumption due to the increase in energy expenditure in patients treated with 2,4-dinitrophenol in the 1930s [40, 44, 45]. In a recent study, 2,4-dinitrophenol was provided to mice for their entire adult lifespan and daily food consumption remained unchanged. However, a significant reduction in body weight was observed, as well as reduced serum glucose, insulin and triacylglycerol, relative to the untreated group [43]. Over the past 10 years, numerous patents have been published by Novo Nordisk indicating that the company has investigated chemical uncoupling [47, 48]. To make such an approach pharmaceutically possible, the therapeutic index has to be considerably improved over 2,4-dinitrophenol. Drugs that have the potential to make patients look or feel good are susceptible to abuse. 2,4-Dinitrophenol lacked weight loss-diminishing effects at higher doses, and patients tended to overdose. A modern chemical uncoupler may therefore need to be formulated with a delivery method (e.g. oral pump, pro-drug) that curtails overdosing. The positive and negative long-term consequences of increasing energy expenditure are unknown, but this could be addressed as a component of the drug approval process.

Wasting energy: a unifying theory to tackle over-nutritional phenotypes

The term 'over-nutritional phenotype' refers to the expressed or displayed effect when energy intake exceeds the tolerated threshold of an individual's combined genetic composition and level of physical activity, shown as an increased incidence of insulin resistance, obesity, type 2 diabetes, cancer, sleep apnoea, depression, inflammation, cardiovascular disease, hypertension, non-alcoholic fatty liver disease, etc. [49–60]. It is critical to understand that even with the most unfortunate genetic composition that predisposes an individual to metabolic disease [61], such as the well documented US Pima Indians with the world's highest prevalence of type 2 diabetes (38%), that physical activity (energy expenditure) and diet can have a profound effect on upon prevention. As an example, the lesser known Pima Indian tribe in Mexico has a 7% prevalence of type 2 diabetes, fivefold lower than the US tribe. However, the Mexican Pima Indians are much leaner (~30 kg lower body weight), chiefly due to greater physical activity (even in leisure time) and a low-fat diet including complex carbohydrates [62]. Comparative polymorphic analysis suggests that although the US and Mexican tribes are no longer identical, they share a closely held gene pool and that the difference in their phenotype is unlikely to be accounted for by a substantial genetic drift. For the remaining population that find it difficult or impossible to modify diet and

increase physical activity, wasting energy (the loss of potential substrates for 'useful work') may be a fundamental mechanism to target over-nutritional phenotypes (Fig. 5) and should be fully explored. Consider, for example, the energy that is lost by heat via chemical uncoupling vs making ATP or the loss of potential energy through urinary glucose excretion effected by sodium glucose transporter 2 inhibition [63]. These mechanisms create an unbalanced equation in favour of weight loss, without directly targeting pathways controlling mood and satiety. What other 'futile' cycles can be used to change the energy balance? One possibility, perhaps, is the reduction of fatty acid absorption in the intestine or the increase of fatty acid excretion in the urine, making fatty acids thus unavailable as an energy source or as signalling molecules. These ideas, centred on the notion of 'wasting energy', as well as a paradigm shift from the historic focus on searching for risk alleles towards a considerably increased emphasis on searching for protective alleles [64], could provide new therapeutic opportunities to treat the over-nutritional.

Satiety and energy expenditure hold the key to reversing the pandemic of over-nutritional phenotypes as a fundamental cure of symptoms manifested as obesity, diabetes, hypertension, cardiovascular disease, non-alcoholic steatohepatitis, cancer, depression, etc. Given the potential issues with drugs that result in satiety through attenuation of 'reward' pathways, e.g. increased incidence of depression and suicide with rimonabant, targeting energy expenditure may take centre-stage for the treatment of over-nutritional phenotypes. Importantly, for individuals beyond prevention, even moderate weight loss of ~8 kg (17 lbs) had a profound impact on type 2 diabetics [65], yet most patients find sustaining weight reduction difficult. In animal models

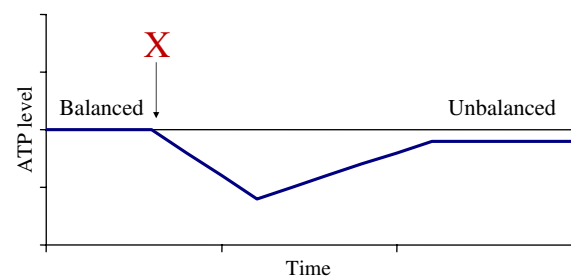


Fig. 5 Unifying theory: waste energy. There is no easy way to resolve an over-nutritional phenotype. However, it is clear that either energy-in must be reduced or energy-out increased to reduce body weight. From an energy-out perspective, wasting energy by some mechanism is essential to take the body metabolically out of balance. A compound that wastes energy (X) could, for example, do this by increasing urinary excretion of fuels (glucose, NEFA), releasing energy as heat, blocking intestinal absorption of potential fuels or by some as yet unknown mechanism. Whole-body energy-out has to be greater than energy-in and sustainable. The goal is to open the box for all possible disease-modifying therapies to realistically resolve the pandemic of over-nutritional phenotypes. Blue line, energy (ATP level)

in which $\dot{V}O_2$ was shown to be increased this effect was repeatedly found to be ‘disease modifying’ [43, 66–69]. Insulin resistance, increased body weight, adiposity, steatosis, and elevated plasma lipids and cholesterol can be resolved by chronic increases in $\dot{V}O_2$, suggesting a positive effect upon increasing energy expenditure. The significant hurdle in the search for novel drugs in this field is the identification of a mitochondrial uncoupler that is safe. However, given the potential benefits to patients and considering the pandemic of the over-nutritioned population, the endeavour is worth the effort.

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