

# Changes to Metabolism and Cell Physiology that Enable Mammalian Hibernation

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**Abstract** Heterothermy is a widespread, adaptive strategy used by many species of bird and mammal to conserve energy during periods of energetic deficit, the expression of which varies greatly depending on the species and environment. A temporary, reversible reduction in metabolic rate and body temperature (i.e., torpor) is an adaptive response used by many species of birds and mammals to conserve energy during periods of resource scarcity. Long-term employment of torpor (i.e., hibernation) is a seasonally expressed phenotype, the genetic and regulated pathways of which can be found throughout all mammal lineages, including hibernators and nonhibernators alike. In mammals, adaptations that allow for hibernation can be classified as those involved in preparation for hibernation, metabolic reduction, continued cellular function and protection, and arousal. Key physiological changes involve seasonal regulation of metabolic hormones, a shift to largely using endogenous fuel sources (i.e., increased lipolysis), global down regulation of protein transcription by posttranslational modification and microRNA save for the increased production of a small number of protective proteins, shifts in membrane composition, and thermogenesis by brown adipose tissue. There is some evidence of cold acclimations in nonhibernators, such as during fetal development, but responses are limited and cursory, and

eventually cellular damage occurs. Therefore, it appears that a complete suite of adaptations to metabolism, vital physiological functions, and thermogenic mechanisms is required for the successful expression of the hibernation phenotype.

**Keywords** Cellular physiology · Gene expression · Hibernation · Hypothermia · Metabolism · Phenotypic plasticity · Torpor

## Abbreviations

ATP	Adenosine triphosphate
BAT	Brown adipose tissue
MR	Metabolic rate
PUFA	Polyunsaturated fatty acid
$T_a$	Ambient temperature
$T_b$	Body temperature
$T_{set}$	Hypothalamic body temperature setpoint
WAT	White adipose tissue

## Introduction

Endothermic animals rely on metabolically produced heat to maintain high body temperature ( $T_b$ ) independent of environmental conditions [104]. Most enzyme-mediated reactions occur optimally within a very narrow temperature range [92], and endothermic thermoregulation ensures biological function over a wide range of ambient temperatures ( $T_a$ ). However, there is considerable energetic demand for the maintenance of high metabolic rates. An adaptive thermoregulatory strategy known as torpor—the use of temporary, reversible reductions in metabolic rate (MR) and  $T_b$ —has evolved in some animals to deal with

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situations in which maintaining high  $T_b$  is energetically restrictive [77, 198]. Torpor is a physiological state of heterothermy in which MR and  $T_b$  are concomitantly and purposefully reduced to conserve energy, followed by an increase in metabolism and a rewarming of the body to a state of normal function using endogenous sources of heat [198].

Torpor can be defined more narrowly as being daily or seasonal (i.e., hibernation). Daily torpor is characterized by bouts lasting less than 24 h with  $T_b$  often remaining some degrees above  $T_a$  [176]. In contrast, hibernation is extended and often lasts for months with torpor bouts of multiple days to weeks interspersed with short periods of arousal and euthermy [124]. Hibernation can be further split into obligate hibernation, defined by predictably seasonal drops in MR,  $T_b$ , and activity regardless of external factors (e.g., [162]), and facultative hibernation when animals use torpor for extended (i.e., more than 24 h) periods outside the typical “hibernation season” in response to an ephemeral energetic challenge (e.g., [202]). During hibernation, body temperature setpoint ( $T_{set}$ ) is considerably reduced and  $T_b$  often remains at or very near (within 1 °C of)  $T_a$  [85]—in the range of 2–10 °C for most hibernators, but as low as –2.9 °C in extreme cases [11, 139]. Concomitantly, the rates of biological processes are slowed to a fraction of normal values. Torpid metabolic rate during hibernation can be less than 1 % of resting metabolic rate [74], and heart rate can decrease by more than 97 %; for example, mean heart rate in the hibernating Gould’s long-eared bats (*Nyctophilus gouldi*) drops to as low as 5 beats/min during hibernation compared to a mean of 228 beats/min in a normal resting state [49, 50]. Respiratory rate is also affected in torpor; it is not only reduced but also completely arrested for periods of time, with some hibernators experiencing prolonged periods of apnea [93].

Many small mammals and some birds use torpor. Examples of heterothermic species can be found throughout the classes Aves and Mammalia [77, 176, 198], including hummingbirds [127] and nightjars [27], marsupials [167], rodents [72, 150], bats [182], and primates [52]. The phylogenetically widespread distribution of heterothermic species (Fig. 1) suggests torpor is an ancestral trait [136]. Moreover, although there is considerable variation in the degree to which heterothermic species throughout the class Mammalia express torpor, species that are closely related tend to express similar torpor characteristics [176], which further suggests heterothermy is a plesiomorphic trait. With the exception of the common poorwill, *Phalaenoptilus nuttallii* [27], hibernation is restricted to mammals, mainly those species living in temperate regions that experience extended periods of food shortage (i.e., winter; [198]). The hibernation phenotype may have evolved in some lineages as an adaptation to further support maintenance and

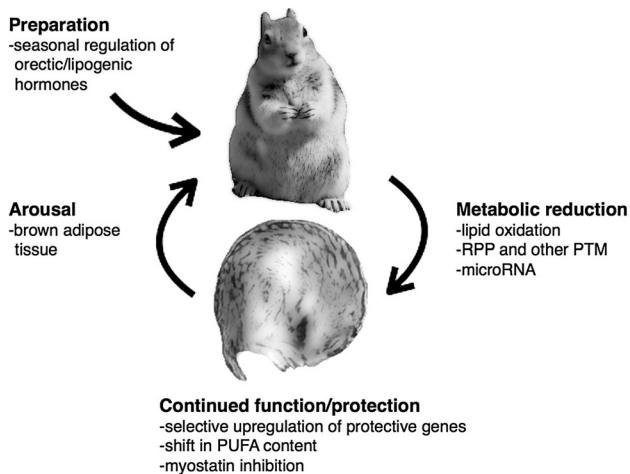
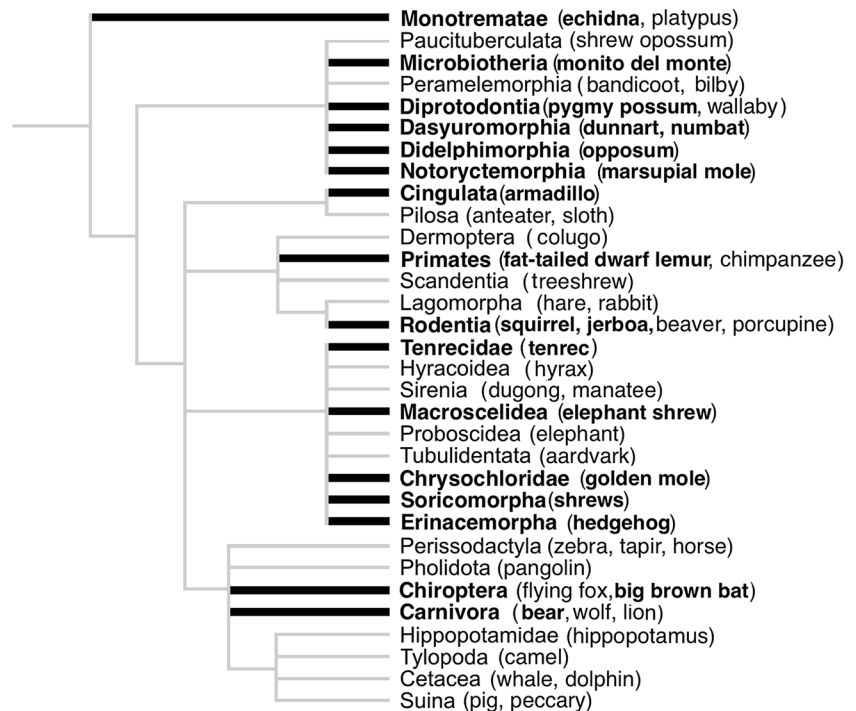
survival at lower  $T_b$  and slower MR during periods of energetic deficit [8, 112].

My review focuses on the cellular and physiological changes associated with hibernation, as well as the molecular underpinnings when appropriate. Adaptations associated with hibernation can be classified into four main categories (Fig. 2): (1) those that allow animals to prepare themselves for hibernation (i.e., put on fat); (2) those that result in state of metabolic reduction, and thus lower  $T_b$ ; (3) those that allow continued cellular function in the cold and prevent damage induced by reduced biological synthesis; and (4) adaptations allowing for a rapid return to normal function upon arousal, which allows the oft-overlooked characteristic of hibernation in that it is regulated and all associated physiological changes are non-permanent and reversible. Arousal by endogenous means is a key feature for distinguishing torpor from unregulated hypothermia (i.e., uncontrolled periods of lower than normal  $T_b$ ; [79]). The majority of our knowledge about hibernation physiology stems from research on rodents, particularly sciurid hibernators, such as ground squirrels and marmots. Seasonally obligate hibernators such as these represent the most extreme and most well-studied phenotype. Therefore, much of this review focuses on the hibernation biology of this group of mammals during the hibernation season. However, consistent patterns in metabolism and physiology can be found in all hibernators, including facultative hibernators that enter long-term torpor during other times of the year. In such cases, many changes are unnecessary or less pronounced, such as accumulation of large fat reserves, but the mechanisms of metabolic depression are likely the same as those used during seasonal hibernation.

## Hibernation Triggers and Control of Physiological Processes

There is considerable debate as to which specific system triggers seasonal torpor and influences patterns of physiological activity once in hibernation. Activity is purported to be largely under the control of a master endogenous clock located within the suprachiasmatic nuclei (SCN) of the hypothalamus [173]. This system is driven by photoperiod and results in circadian (i.e., daily) oscillations in metabolism, appetite, and activity [124, 173]. Patterns of daily torpor appear to fall under circadian control, which would allow animals to continue to arouse and function (e.g., forage and locomote) during normal periods of activity, but circadian rhythm appears to have little influence on hibernation. Seasonal changes in day length and levels of light-driven hormones, such as melatonin, likely drive circannual rhythms as well. For example, waning daylight

**Fig. 1** Dendrogram depicting major groups (Families and Orders) within the class Mammalia that are known to use torpor (*black lines and bold text*). Examples of species within each group that use torpor appear in *bold text*. The wide distribution of torpor suggests it is an ancestral trait. The dendrogram was prepared using data from Lovegrove [136], Ruf and Geiser [176], and the Interactive Tree of Life [132]



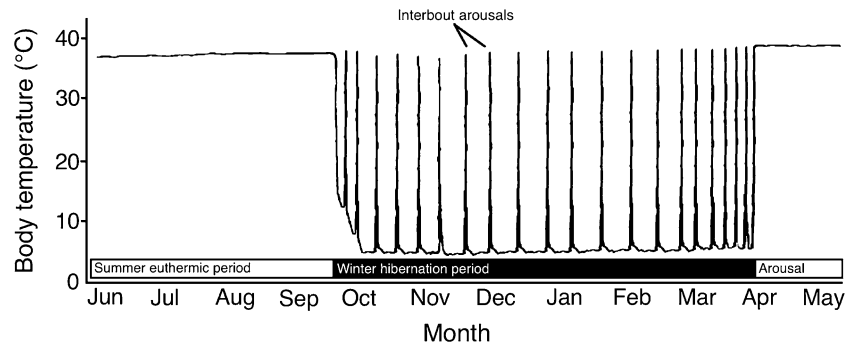
**Fig. 2** A schematic outlining the four categories of adaptations for hibernation: (1) those that allow animals to prepare themselves for hibernation; (2) those that result in state of metabolic reduction and thus lower  $T_b$ ; (3) those that allow continued cellular function in the cold and prevent damage incurred from reduced biological synthesis; and (4) adaptations allowing for a rapid return to normal function upon arousal

levels are purported to trigger seasonal pre-hibernation fattening [171, 172]. In addition, timing of seasonal hibernation is predictable (see Fig. 3) and in most cases occurs regardless of environmental conditions. Circannual rhythms of hibernation can persist even under constant photoperiod and temperatures (e.g., [53, 119]), which suggests intrinsic circannual control [124]. However,

energy availability also influences entrance into hibernation. Reductions in food availability lead to metabolic stress [76], signaled by increased ratios of [ADP]:[ATP] and [NAD<sup>+</sup>]:[NADH]. The cellular response to falling concentrations of energy substrates is a reduction in metabolic rate and an increased reliance on lipolysis for ATP production [76], two main characteristics of hibernation. Furthermore, phenology of hibernation is influenced by sex and body condition, and can vary within a species and even within a single population [201]. For example, male little brown bats (*Myotis lucifugus*) enter hibernation later than females and fatter females emerge earlier than other roostmates [157]. Thus, onset of hibernation appears to be an underlying response to photoperiod and influenced to some degree by other environmental factors that predictably wane prior to the hibernation season, such as the abundance of food on the landscape, as well as individual condition.

Photoperiod, however, is not likely to influence circadian rhythms of animals in concealed dens, roosts, or burrows, and food availability rarely fluctuates throughout the winter. This raises the question: what controls rhythms in activity and physiological processes during hibernation when these cycles are disrupted? Current opinion is split as to whether circadian control of activity during hibernation is modified to be photo-independent or completely abandoned. Transfer of control of physiological processes to an alternate, temperature-dependent circadian clock that is not affected by photoperiod would result in patterns of activity

**Fig. 3** Simplified schematic of  $T_b$  over time of a typical hibernator, showing euthermia in the summer, hibernation with frequent interbout arousals during the winter, and final arousal in the spring ( $T_b$  pattern adapted from Epperson and Martin [62])



that slow down with reduced  $T_b$  [141]. Under such control, maintenance of semi-regular patterns in arousal and activity (i.e., interbout arousals; see Fig. 3) during extended periods of darkness and quiescence would continue. The fact that duration of torpor bouts negatively correlates with  $T_b$  at temperatures above  $T_{set}$  during hibernation (e.g., [82]) seems consistent with this hypothesis. In addition, patterns of activity when arousals occur seem to remain under some circadian influence during part hibernation. For example, in hibernating *Myotis* bats (Natterer's bat, *M. nattereri*; Daubenton's bat, *M. daubentonii*; whiskered bat, *M. mystacinus*; and Brandt's bat, *M. brandtii*), timing of peak activity throughout winter remains linked to sunset and diurnal activity is uncommon [107]. However, there is evidence that the influence of a circadian rhythm is lost during hibernation. Clock genes *Period1* and *Period2* are expressed constantly during hibernation as opposed to a normal, rhythmic, diel pattern [165], and patterns of arousal during hibernation that contradict a diel rhythm have been reported. For example, arousals in some populations of little brown bats show no circadian pattern [51]. Furthermore, warmer ambient temperatures result in more frequent arousals by bats hibernating in buildings than in caves [89], which suggests roosting ecology (and therefore ambient conditions) affects arousal patterns. In general, currently available data are inconclusive as to whether a modified circadian clock persists or if another intrinsic timing mechanism is involved. Patterns of physiological activity are likely controlled by interactions between endogenous clocks (both circadian and circannual), photoperiod, temperature, resource availability, and metabolite imbalance.

The intrinsic mechanisms behind the onset of hibernation also remain poorly understood. Current opinion is that changes in MR and  $T_b$  during hibernation are controlled by way of a "sliding setpoint." Adaptations of the hypothalamus adjust the temperature around which  $T_b$  is metabolically defended [101]. Just as a furnace does not produce heat until  $T_a$  reaches the setpoint of the thermostat, metabolic heat production in the body is largely reduced until  $T_b$  reaches the newly established setpoint ( $T_{set}$ ).

During entrance into hibernation, decreased  $T_{set}$  and concomitant reduction of metabolic rate precede a more gradual decline in  $T_b$  [102]. The metabolic response of hibernating animals when  $T_b$  drops below  $T_{set}$  is also important and further supports the idea of setpoints rather than complete abandonment of thermoregulation. As  $T_b$  drops below  $T_{set}$ , metabolic activity increases proportionately and torpor bouts are shortened to bring  $T_b$  back up to  $T_{set}$  (e.g., [56, 78]). This further suggests thermoregulation during hibernation is not completely abandoned and hypothalamic control is maintained at low temperatures [101]. Ecologically, maintenance of  $T_b$  around a setpoint lends importance to hibernacula microclimate;  $T_a$  in hibernacula above or below the species-specific  $T_{set}$  increases metabolic activity and diminished energy savings.

Although the precise mechanisms involved in hypothalamic control of  $T_{set}$  and  $T_b$  remain largely unknown, recent evidence suggests involvement of central A1 adenosine receptors (A1AR) in the preoptic area of the hypothalamus (POAH) and the circadian-regulated nucleotide adenosine 5'-monophosphate (AMP). Neurons within the POAH monitor  $T_b$  and regulate appropriate responses [149]. Activation of A1AR appears necessary for torpor; blocking A1AR activity prevents and disrupts hibernation [113] and A1AR agonists can successfully induce a torpor-like state, even in nonhibernating lab rats (*Rattus norvegicus* [193]). In addition to activating A1AR in the POAH, AMP also reduces activity of neurons responsible for inducing a thermogenic response to cold, which ultimately lowers  $T_{set}$  [153]. Levels of AMP are elevated in the blood of hibernating mice kept in constant darkness, and AMP induces hypothermia in mice (*Mus* spp.) despite normal circadian light cycles [212]. Seasonality of hibernation also suggests a relationship between AMP and A1AR in inducing hibernation. Stimulation of A1AR induces metabolic suppression during but not outside of the hibernation season (i.e., winter) in Arctic ground squirrels (*Urocitellus parryii* [116]), likely due to seasonal adenosine sensitization (i.e., increased A1AR density) in the winter or relatively low levels of AMP during summer. These findings support the role of AMP and its effect on A1AR in the onset of torpor.

However, physiological parameters measured during induced hypothermic states are typically inconsistent with those of true hibernation [186, 187].

## Adaptations in Preparation of Hibernation

### Fat Storage

The most outwardly obvious change in preparation for hibernation is the accumulation of large fat stores (e.g., [17, 31, 128]). Food intake doubles or triples preceding hibernation, and body mass increases by as much as 62 % in free-ranging individuals [177]. In nonhibernators, satiety signals maintain body adiposity and energy balance. Numerous pathways and proteins activated by accumulation of surplus adipose tissue elicit a counter response. For example, adipocytes in white adipose tissue (WAT) release leptin when adiposity is high, which acts on the hypothalamus in long-term suppression of appetite and enhances lipid oxidation [35, 183]. Leptin also inhibits AMPK-activated protein kinase (AMPK), a metabolic “master switch” that stimulates appetite [123] and ATP production from energy stores [147]. High adiposity also reduces hunger by inhibiting the release of ghrelin, a hormone that stimulates appetite and food intake [48]. Physiological changes in hibernators allow them to temporarily override signals of adiposity to continue pre-hibernation fat storage.

Concentrations of leptin and ghrelin typically follow seasonal fluctuations (reviewed in [96]). Body mass and leptin levels continue to increase concomitantly during the pre-hibernation period in woodchucks (*Marmota monax* [45]), and leptin is positively correlated with increasing fat mass and fat cell size during pre-hibernation mass gain in yellow-bellied marmots (*Marmota flaviventris* [67]). An increase in leptin levels is expected with hyperphagia, but its anorexigenic (i.e., appetite suppressing) effects are inhibited, suggesting seasonal resistance. In little brown bats, a leptin increase occurs just prior to fattening but decreases during fattening, suggesting complete dissociation of leptin release from adiposity during pre-hibernation fattening [126]. Appetite can be suppressed with a leptin injection during pre-hibernation hyperphagia in Arctic ground squirrels, which also suggests sensitivity to the hormone is retained but dissociation occurs [160]. Regardless of mechanism, seasonal regulation of the anorexigenic effect of leptin allows hibernators to continue to eat and fatten during pre-hibernation. Leptin sensitivity is restored shortly after entering hibernation; high concentrations of leptin in hibernating Columbian ground squirrels (*Urocitellus columbianus*) continued to restrict food intake despite food being available [68]. Likewise, golden-man-

ted ground squirrels (*Callospermophilus lateralis*) experimentally induced to hibernate in warm temperatures ( $T_a = 22\text{ }^\circ\text{C}$ ) had higher metabolic rates, declining fat reserves, and an associated drop in leptin levels and return of appetite [68].

Ghrelin inhibition also appears to be seasonally regulated in hibernators. Despite increasing adiposity during the hyperphagic pre-hibernation period of golden-mantled ground squirrels, levels of ghrelin remain higher than any other time of the year [99], suggesting a seasonal upregulation of circulating levels of the hormone [192, 209] and possible dissociation from the adiposity feedback loop. In addition to directly stimulating appetite, ghrelin amplifies its hyperphagic effect by phosphorylating and activating AMPK [97, 98]. Although AMPK stimulates appetite, it also inhibits lipogenesis and other anabolic processes that are counterproductive during pre-hibernation fattening [90]. However, anabolic-inhibiting effects of AMPK are not observed in the fall. Seasonal dissociation of AMPK may mitigate this conflict, but a test of this idea is required.

Differential insulin sensitivity also plays an important role in pre-hibernation adiposity. Plasma insulin promotes uptake of blood glucose to be stored as glycogen in the liver and muscles, decreases lipolysis and increases lipid synthesis, and stimulates the production of triglycerides and adiposomes [208]. Effects of plasma insulin are therefore advantageous during periods of pre-hibernation fat accumulation. However, levels of plasma insulin are proportional to levels of cerebrospinal fluid (CSF) insulin, which decreases appetite and thus inhibits adiposity [207]. The responses of hibernating animals to insulin suggest a seasonal sensitivity. In both golden-mantled ground squirrels and yellow-bellied marmots (*Marmota flaviventris*), insulin level increases expectedly with adiposity during autumn fattening [21, 66] but does not signal satiety [66]. In grizzly bears (*Ursus arctos horribilis*), insulin sensitivity also follows seasonal patterns with adipocytes showing three distinct levels of responsiveness throughout the cycle [155]. During pre-hibernation fattening, adipocytes are hypersensitive to serum insulin only to become insulin resistant during the onset of hibernation and then resume normal levels of insulin sensitivity post-hibernation [155]. Selective insulin resistance appears similar to leptin resistance in that it is reversed shortly after the individual enters hibernation [65]. Insulin levels remain low throughout much of the hibernation period in most species [65] but gradually rise during the last half of the hibernation period in the little brown bat [14]. This may reflect the importance of a readily available energy pool immediately upon arousal and emergence to ensure the ability to fly.



Although the exact mechanism behind selective hormone resistance is unclear, a recent study hints at tissue-specific changes to hormone receptors as having a role. Regardless of age or sex, androgen receptors of pre-hibernating Arctic ground squirrels are selectively upregulated in skeletal and brain tissue [19]. This stimulates building of muscle tissue, which is necessarily catabolized along with lipids during hibernation in the harsh Arctic climate, without costs typically associated with high testosterone levels in other systems, such as immunosuppression [135] or increased reproductive effort [204]. Whether selective receptor regulation is responsible for the shifts in sensitivity to other hormones, such as insulin, during hibernation remains to be seen but is a promising place to start.

## Metabolic Adaptations for Hibernation

### Metabolic Fuel

The primary source of fuel for mitochondrial respiration shifts from carbohydrates to lipids during hibernation. Indirect evidence of reliance on lipids as a fuel source comes from the observation of a reduction of fat stores as winter progresses (e.g., [21, 31]). Measurable shifts in respiratory quotient (RQ;  $\text{CO}_2\text{:O}_2$  in exhaled breath) also indicate a switch from carbohydrates to fats during hibernation. A RQ value close to 1.0 indicates carbohydrates are the main metabolic substrate, whereas values of 0.7 suggest lipid oxidation [138]. During hibernation, RQ in European hedgehogs (*Erinaceus europaeus*) is 0.7, indicating lipid metabolism [188]. Likewise, in Arctic ground squirrels, RQ averages 0.70 during steady-state torpor at  $T_a$  between 4 and 8 °C, indicating exclusive use of lipids for energy production [32]. However, outside this range of temperatures at -16 and 20 °C, RQ rises to values >0.85, suggesting a return to primarily carbohydrate-fueled metabolism during arousal [32].

Some mechanisms behind the switch in metabolic fuel source simultaneously inhibit the use of carbohydrates and increase the availability of fatty acids. Upregulation of genes for pyruvate dehydrogenase kinase isoenzyme 4 (PDK4) and pancreatic triacylglycerol lipase (PTL) occurs in the heart, skeletal muscle, and WAT of thirteen-lined ground squirrels (*Ictidomys tridecemlineatus*) during hibernation [7]. Induction of PDK4 inhibits activity of pyruvate dehydrogenase (PDH) by phosphorylation. Conversion of pyruvate to acetyl-CoA is dependent on PDH, and inhibition of this pathway halts entry of glycolytic intermediates into the TCA cycle [6]. In conjunction with PDK4 inhibition of glycolysis, PTL stimulates lipolysis by breaking down adiposomes (i.e., “fat droplets”) to release

fatty acids for use in the  $\beta$ -oxidation pathway. High lipolytic activity at low temperature is inherent to PTL and conserved across all mammal lineages, but its upregulation and consequential surge during lipolysis appears to be unique to the hibernating state [5]. Other proteins that are elevated during hibernation include liver fatty acid-binding protein, fatty acid transporter, and 3-hydroxy-3-methylglutaryl-CoA synthase, all of which support the switch to lipid metabolism [63].

### Metabolic Rate

In mammals, 90 % of oxygen consumption occurs in the mitochondria, of which 80 % is involved in ATP synthesis and 20 % in maintaining ion concentration gradients [169]. Given that mitochondrial respiration is a substantial site of energy turnover and heat production, changes within are key to metabolic reduction and  $T_b$  reduction. Rates of enzymatic reactions decrease as temperature drops [92]. The temperature coefficient of metabolic activity ( $Q_{10}$ ) is the change in rate of oxygen consumption over a 10 °C temperature range [91]. Most biological reactions have a  $Q_{10}$  of 2–3, but values >3 are typical during hibernation, suggesting additional physiological changes to inhibit metabolic activity beyond that expected based on decreases in  $T_b$  alone [74]. As much as 76 % of metabolic depression in hibernating edible dormice (*Glis glis*) is purportedly due to physiological inhibition as opposed to thermodynamic effects [100].

Decreases in metabolic rate are correlated with reductions in enzyme activity and available substrate [169]. For example, activity of glycogen phosphorylase, which converts stored glycogen into glucose, decreased by as much as 75 % of euthermic levels in liver and kidney of golden-mantled ground squirrels [29]. Reversible protein phosphorylation (RPP) has emerged as a crucial mechanism in the shift between active and inactive states in both metabolic enzymes and substrates [184]. Substantial reduction in metabolic activity occurs when metabolic enzymes and substrates are inhibited by covalent addition of phosphate groups, a potentially widespread process carried out by protein kinases and quickly reversed by protein phosphatases [184]. In golden-mantled ground squirrels, the proportion of active pyruvate dehydrogenase (PDH), which converts pyruvate into carbohydrates that enter the tricarboxylic acid (TCA) cycle, drops from 60 to 80 % in heart and kidney of euthermic animals to as little as 4 % during hibernation, the rest inactivated by RPP [29]. Phosphorylation of skeletal muscle hexokinase (HK), an enzyme that catalyzes the conversion of hexoses into hexophosphate, also occurs during hibernation [2]. In Richardson’s ground squirrels, activity of phosphorylating HK is reduced by 33 % compared to euthermic values [2].

This is particularly important for the reduction of glycolysis because HK converts glucose to glucose-6-phosphate (G-6-P), which then enters cells and is used in glycolysis. Deactivating enzymes involved in carbohydrate catabolism by RPP facilitates the switch from carbohydrates to lipids as the main metabolic fuel source and is rapidly reversible [184].

Most cellular processes that consume energy are directly or indirectly associated with membranes and the maintenance of concentration gradients [169]. Thus, activity of membrane-associated proteins has profound effects on metabolic rate [61, 110] and regulated suppression of ion channels by RPP is a key factor in reducing ATP turnover and metabolic reduction. For example, phosphorylation of  $\text{Na}^+/\text{K}^+$ -ATPase (i.e., sodium–potassium pump) in golden-mantled ground squirrels results in reduction of that pump's activity by up to 60 % [185]. Likewise, RPP of sarcoplasmic reticulum  $\text{Ca}^{2+}$ -ATPase in hibernating long-tailed ground squirrels (*Urocyon undulates*) reduces activity of the pump by 50 % compared to active animals [142].

Protein synthesis is responsible for up to 10 % of resting metabolic rate in mammals [169] and is largely globally downregulated during hibernation [122]. For example, in hibernating thirteen-lined ground squirrels, the rate of protein synthesis is reduced to 34 % of the euthermic value in brain tissue [73] and 15 % in kidney tissue [106]. Likewise, in torpid golden-mantled ground squirrels, transcription is reduced to 50 % of the value in euthermic animals between bouts of torpor [195]. Observed decreases in activity have  $Q_{10}$  values between 2 and 3, which are to be expected based on temperature effects alone [195]. However, at least some suppression of gene expression is purportedly due to posttranslational modification of proteins [185]. Small ubiquitin-related modifiers (SUMOs) have been shown to target and turn off transcription factors [86]. The brains, liver, and kidney of hibernating thirteen-lined ground squirrels showed elevated SUMOylation of proteins [131]. The effect of SUMOylation is reproducible in human-derived research cells, suggesting the underlying mechanism is not unique to hibernators [131]. However, the large-scale use of such modifications in metabolic reduction is not evident outside mammals with the hibernation phenotype.

Interestingly, a number of key metabolic proteins seem largely absent in metabolic rate depression during hibernation. In particular, AMPK has a central role in regulating energy balance in euthermic states and could be expected to have an essential role in global metabolic depression. AMPK is activated by a rise in cellular ADP:ATP ratio and stimulates catabolic ATP-generating processes, including mitochondrial biogenesis [166]. However, most studies report only tissue-specific changes in AMPK levels during

hibernation. In thirteen-lined ground squirrels, relative abundance of AMPK was three times higher in WAT of winter animals compared with that of summer animals, but there were no differences in other tissues [108]. Similarly in golden-mantled ground squirrels, only WAT showed higher total AMPK abundance in hibernation compared to euthermia, with no differences detected in brown adipose, muscle, or liver tissue [98]. The fact that changes in AMPK expression are not global and occur only in WAT suggests AMPK does not play a key role in metabolic depression during hibernation. Additional regulation of AMPK may not be necessary given that disruption of ADP:ATP ratio might occur during entry into torpor [118] but is largely restored once ATP-genesis and ATP-turnover balance in steady-state hibernation. Metabolic fuel source shifts from carbohydrates to lipids during hibernation, and higher AMPK levels in WAT may conserve metabolic ability of these tissues despite low  $T_b$ .

## Key Adaptations to Cold and Quiescence

### Gene Expression

Steady-state levels of most mRNAs and proteins do not decrease considerably during hibernation, despite reduced synthesis [39, 62, 180, 181]. Enzyme kinetics predicts protein degradation to slow down as  $T_b$  drops, but protein loss and degradation of mRNA proceed even slower than expected given the length of time that hibernators remain dormant. Mechanisms exist to conserve genetic precursors during hibernation, such as a seasonal increase in the robustness of the poly(A) tail of mRNA [122]. Residues along the poly(A) tail of mRNA isolated from hibernating Arctic ground squirrels showed increased binding sites for stabilizing proteins, an alteration not seen in summer-active ground squirrels [122]. Expression of proteins involved in folding, stabilizing, and transporting other proteins is also increased [63]. Nevertheless, protein loss does occur and may be a reason for interbout arousals. Protein synthesis during arousals is rapidly increased and often exceeds the rate during the active season [63].

A small number of some genes are upregulated during hibernation, aside from those associated with metabolism already mentioned (e.g., PDK4 and PTL). Levels of some proteins produced in the liver and released into the blood are increased to preserve cellular function at low temperatures, such as  $\alpha$ 2-macroglobulin ( $\alpha$ 2m; [62, 180]), which prevents clotting and aids in blood perfusion at low temperatures [181], and apolipoprotein A1 (apoA1), which transfers long-chain fatty acids to target tissues [62]. Sarcoplasmic reticulum  $\text{Ca}^{2+}$ -ATPase (SERCA2a) is upregulated in the heart of hibernating woodchucks relative

to active conspecifics [25]. Increased activity of SERCA not only preserves cardiac function at low  $T_b$  by maintaining  $Ca^{2+}$  balance [87], but also provides localized metabolic heat within the membrane of cardiomyocytes [54]. Selective regulation of protein synthesis allows hibernators to decrease energy consumption while simultaneously maintaining levels of those proteins important for cellular function.

MicroRNA (miRNA), small non-coding transcripts that bind to mRNAs and target them for degradation or inhibit their translation [13], has recently been described as having a pivotal role in controlling global gene expression during hibernation. Studies investigating miRNA expression during hibernation are limited but promising. Selective and tissue-specific regulation of miRNAs may preserve critical organ function and contribute to metabolic change during hibernation. In thirteen-lined ground squirrels, levels of miRNA-1 and miRNA-21 in kidneys are 100 and 30 % higher in hibernation compared to euthermia [148]. Increased cell proliferation is associated with miRNA-1 expression [43], and miRNA-21 inhibits cellular apoptosis [41]. Maintenance of kidney structure and function during hibernation is likely aided by upregulation of these miRNAs. Other miRNAs are downregulated during hibernation to reduce ATP turnover and promote lypolysis. Levels of miRNA-24 and miRNA-122a transcripts are reduced in heart and skeletal muscle of torpid animals [148]. Reduction in miRNA-24 activity is implicated in the suppression of cell growth and differentiation [129], and inhibition of miRNA-122a reduces lipogenesis and promotes fatty acid oxidation [64]. The role of miRNAs has only recently been suggested in hypometabolism, and empirical data are sparse. Given that microRNAs are evolutionarily conserved in all animals and regulate 30–50 % of all protein-coding genes [13], their involvement in hypometabolism is highly likely and further research is warranted.

### Membrane Function

Lipid membranes are selectively permeable boundaries vital to almost all cellular processes. Membrane fluidity and activity of inter-membrane pumps are reduced as temperature drops [4, 42]. Cold-induced leakage and disruption of ion gradients typically causes catastrophic loss of cellular and organ function [33, 199]. Unlike the widespread polyunsaturation observed in cold-adapted ectotherms [46, 95], a phenomenon known as “homeoviscous adaptation” [178], mammalian hibernators maintain membrane function at cold temperatures without consistent patterns of polyunsaturation, suggesting different mechanisms are involved [4].

Most membranes show some change in unsaturation index during hibernation, and there are measurable effects

on hibernation. Increased consumption of dietary PUFAs in the fall facilitates entrance into torpor and lengthens and deepens torpor bouts during hibernation (e.g., [69, 75, 81]). However, increased membrane PUFA composition is a double-edged sword; a global increase in total membrane PUFA concentration would result in a concomitant and counterproductive rise in membrane-associated metabolic activity [109]. PUFAs are also more prone to react with reactive oxygen species (ROS), which causes peroxidative damage [70], making high concentrations unfavorable. Animals experimentally fed diets higher in PUFA than would be normally ingested hibernate at higher  $T_b$  and for shorter bouts than those fed moderate levels of PUFA (e.g., [70]). Rather than an increase in total membrane PUFA content, a shift in the ratio of n-3:n-6 class PUFAs is central in maintaining membrane function at low temperature. A higher proportion of n-6 fatty acids (e.g., linoleic acid) to n-3 fatty acids increases membrane-bound pump activity [175] without increasing the negative effects associated with PUFA. Hibernators also employ tissue-specific shifts in membrane PUFA composition, and increased total PUFA levels mainly occur in adipose tissue [151]. Oxidative stress caused by increased PUFA levels may additionally be offset by the inherently lower production of peroxides during ATP synthesis in some hibernators. For example, the mitochondria of little brown bats produce as little as one-third the amount of hydrogen peroxide per unit oxygen consumed during respiration than those of short-tailed shrews (*Blarina brevicauda*) and white-footed mice (*Peromyscus leucopus*; [30]).

Cholesterol also increases membrane fluidity at low temperature [179] but there is a distinct lack of evidence compared to studies on PUFA to suggest an active increase in cholesterol uptake by free-ranging hibernators. There is an increase in total cholesterol (plasma, HDL, and LDL) during hibernation, but this appears to be related to increased lipid metabolism in a closed system rather than functional membrane change [161]. However, increased dietary cholesterol has been shown to influence torpor patterns in experimentally hibernated animals, resulting in longer torpor bouts at lower  $T_b$  [83]. Increased cholesterol concentration would allow for increased fluidity and function of cell membranes at lower temperatures. It could be that tissue-specific shifts in cholesterol concentration occur during hibernation as is seen with PUFA, but to my knowledge this has not been observed.

There are also inherent differences in pump substrate affinity and configuration that enables hibernators to maintain optimal rates of ion turnover at lower temperatures than nonhibernators. For example, activity of erythrocyte  $Na^+/K^+$  pumps (i.e.,  $Na^+/K^+$ -ATPase) in Columbian ground squirrels and guinea pigs (*Cavia porcellus*) is similar at high temperatures (37 °C), but turnover



rates are 2–3 times higher in ground squirrels at 5 °C [60]. Affinity of  $\text{Na}^+/\text{K}^+$ -ATPase for ATP is reduced at low temperatures in cold-sensitive cells (e.g., those of guinea pigs), whereas cells of hibernators show an increase in ATP affinity of  $\text{Na}^+/\text{K}^+$ -ATPase when cooled [143]. Cells of hibernators also maintain calcium balance better than nonhibernators at low temperatures. Rats exhibit an increase in the influx of  $\text{Ca}^{2+}$  into the cytoplasm as temperature drops, while the opposite is true in Columbian ground squirrels because fewer calcium channels are present and open [134]. Likewise, Richardson's ground squirrels have fewer  $\text{Ca}^{2+}$ -ATPase and higher threshold activation levels in smooth muscle compared to guinea pigs [205]. Having fewer, less responsive ion channels allows for greater control over ion flux at cold temperatures, but there are associated costs. Stronger stimuli are likely needed for muscle contraction; thus, performance may suffer even at euthermic temperatures. Such trade-offs are consistent with the hypothesis that hibernators are thermal generalists and perform well over a wide range of temperatures but are not thermal specialists (i.e., obligate homeotherms) within their optimum temperature range (i.e., at euthermic temperatures; [8]).

### Neural Function

Continued function of the nervous system is critical during hibernation. Cardiovascular performance needs to be regulated, and animals need to remain responsive to external stimuli. Although  $T_{\text{set}}$  varies depending on species and environment, hibernators typically avoid reaching potentially damaging  $T_b$  much below freezing and actively increase metabolic rate and the use of non-lipid fuels to do so [32]. Bats arouse even with non-tactile stimuli [190], suggesting hibernating animals remain attuned and responsive to their environment. Furthermore, the continued ability to regulate vital processes while in hibernation is not the only neural function conserved. Although previous studies on the effects of hibernation on memory retention have shown conflicting results (e.g., [144–146]), more recent studies tend to show no evidence of adverse effects. For example, Alpine marmots (*Marmota marmot*) trained to perform tasks before hibernating showed no decrease in their ability to complete these tasks after a hibernation period of six months [44], and greater mouse-eared bats (*Myotis myotis*) trained to find food in a maze performed at the same level pre- and post-hibernation [174]. An inherent benefit of quiescence may be that the lack of stimuli reduces the formation of new neural networks and junctions and increases the ability to maintain current pathways and synaptic associations. However, without mechanisms to protect and preserve these pre-hibernation memories, they

would be expected to diminish to some degree during hibernation.

Bouts of cerebral ischemia result in a permanent loss of dendritic branching and synaptic connections, as well as permanent neuronal damage or death [24]. Neural function does decrease during long bouts of hibernation, but neurons of hibernating animals show reduced temperature-dependent loss of synaptic associations and remarkable plasticity. Hypothermia contributes inherently to decreased damage from ischemia. Ischemia-induced cell death primarily occurs by excitotoxicity, which is damage to neurons from massive efflux of neurotransmitters and excessive excitation, ultimately caused by inability to maintain cellular concentration gradients [211]. Cold temperatures inhibit the release of neurotransmitters, such as glutamate, thereby decreasing the risk of excitotoxicity [10]. However, there is a further and prolonged decrease in the activity of ion channels during hibernation than would be expected based on temperature alone.

Channel arrest and downregulation of neuroreceptor activity further protects neurons during hibernation. Actively downregulating *N*-methyl-D-aspartate receptors (NMDAR; a predominant glutamate receptor) by dephosphorylation during hibernation decreases the risk of excitotoxicity [170]. Hibernating Arctic ground squirrels show decreased phosphorylation and function of NMDAR compared to conspecifics during arousals and rats [213]. Reduction of nerve branching does persist during hibernation but may be an adaptation to further lessen the risk of excitotoxicity. Limiting the connectivity between neurons reduces the potential for a glutamatergic surge upon arousal [140].

However, mechanisms to protect neural function in hibernators are not perfect. Although slowed, decreases in neuronal function do occur during long, deep bouts of hibernation as a result of decreased electrical activity. Sleep, particularly rapid eye movement (REM) sleep, is necessary for refreshing somatic connections and maintenance of memory [20]. These restorative functions of sleep do not occur at  $T_b$  below 25 °C during hibernation [125, 196], and the neural effects of torpor are similar to those seen after periods of sleep deprivation [55]. Hibernating animals may therefore arouse periodically to “sleep” and restore neural function. Dendrites of ground squirrels reduced during the course of hibernation are rapidly regenerated upon arousal with no permanent damage [163]. European hamster (*Cricetus cricetus*) also show decreases in the length and complexity of branching of neurons in the intrahippocampal region during hibernation, but there is a remarkable and total reversal to pre-hibernation profiles within 2 h following arousal [140].

## Cardiovascular Function

Cold-induced influx of  $\text{Ca}^{2+}$  into the cytosol causes arrhythmias in cold-sensitive cells [33, 199]. Hibernators maintain calcium balance much better than nonhibernators. European hedgehogs and Richardson's ground squirrels display a distinct lack of intracellular build-up of  $\text{Ca}^{2+}$  with falling temperature, while guinea pigs show a predictable linear increase in cytosolic  $\text{Ca}^{2+}$  concentration [88]. This is partly due to fewer, less responsive membrane-bound  $\text{Ca}^{2+}$  channels in hibernators but tissue-specific membrane polyunsaturation also plays a role. The lack of unsaturation in plasma membranes during hibernation is not global. Higher levels of PUFA occur in the sarcoplasmic reticulum of cardiomyocytes and increase SERCA2a activity [87]. Sequestration of  $\text{Ca}^{2+}$  in the SR by PUFA-facilitated SERCA2a activity and a concomitant shift toward n6-class PUFA in the cell membrane (as seen in other tissues) result in less cytoplasmic  $\text{Ca}^{2+}$  and proper functioning of the heart even at low  $T_b$  [200].

## Skeletal Muscle Function

Maintenance of muscle tissue during hibernation is crucial for shivering thermogenesis and locomotion during arousals and upon emergence. In most mammals, periods of inactivity and disuse lead to breakdown of muscle vasculature [194] and muscle atrophy [168, 191]. However, atrophy is minimal, often nonexistent, in the muscles of hibernators [47, 130, 210]. For example, the muscle fiber properties of flight muscles in hibernating little brown bats and big brown bats (*Eptesicus fuscus*) under natural conditions are the same pre- and post-hibernation [26, 103]. This is unexpected for such long periods of dormancy, especially when compared to rapid and considerable muscle loss from starvation [15, 16], disuse [191], or unloading [152, 168].

Maintenance of muscle tissue during hibernation is due to a combination of adaptations that inhibit muscle protein loss. Inactivity and starvation typically stimulate expression of myostatin, a protein that inhibits growth and differentiation of muscle tissue, causing muscle atrophy [115]. However, assays of muscle tissue sampled from thirteen-lined ground squirrels during different periods throughout hibernation showed no increase of myostatin while entering or during hibernation [28]. The mechanism behind the downregulation of myostatin is unclear, but that myostatin levels increase markedly early during arousals suggests physiological inhibition. The protein p38 mitogen-activated protein kinase (p38<sup>MAPK</sup>) may also play a role in retention of muscle composition. In hibernating little brown bats, p38<sup>MAPK</sup> and several downstream proteins are phosphorylated (i.e., activated) and in higher

abundance compared to euthermic conspecifics [59]. Effects of activating the p38<sup>MAPK</sup> pathway include enhancement of lipid catabolism and prevention of muscle atrophy [59]. Interestingly, muscle mass may even increase despite continued inactivity during hibernation in some species. In thirteen-lined ground squirrels (*Ictidomys tridecemlineatus*), protein synthesis in skeletal muscle drops 66 % from summer levels and muscle volume declines proportionally to other tissues with declining body mass during early hibernation (October through February). However, the ratio of muscle volume to body mass shows a disproportionate increase in the months prior to emergence in May [105]. The fact that no other tissues examined (heart, liver, and intestine) showed such increases suggests muscle tissue is not only maintained but also prioritized and actively synthesized mid-hibernation [105]. The mechanism behind this shift is unclear but could involve seasonal, tissue-specific androgen sensitivity, which has only recently been documented in Arctic ground squirrels [19].

RPP plays a role in maintaining muscle function in cold, anaerobic states. The enzyme creatine kinase (CK) uses ATP to catalyze the conversion of creatine (Cr) into phosphocreatine (PCr), which acts as a readily available source of phosphate for the later conversion of ADP to ATP under anaerobic conditions in muscle tissue [197]. In hibernating Richardson's ground, CK mRNA is reduced by 70 % compared to euthermic values, but CK activity and protein are reduced by only 20 % [1]. The balance occurs as a pool of available but phosphorylated (i.e., inactive) CK [1]. Phosphorylation of CK is quickly reversed, and the conversion of Cr into PCr to power muscle activity is rapidly resumed during arousal [1]. Similarly, hexokinase (HK) is differentially phosphorylated between hibernation and euthermy [2]. Hexokinase facilitates glycolysis by phosphorylation of glucose into G-6-P, the main sugar used for glycolysis [2]. In Richardson's ground squirrels, HK persists in low and high phosphate forms during hibernation, but is quickly and completely phosphorylated (i.e., activated) by AMPK during arousal [2]. Thus, in hibernators, muscle metabolism and activity is mediated through regulated and reversible phosphorylation of key enzymes. The lag between ramping up protein and energy substrate production during arousal is mitigated by dephosphorylating an existing pool of enzymes. The benefit is an immediate source of energy for shivering thermogenesis and movement upon arousal.

## Other Adaptations

Immunosuppression is a consequence of low  $T_b$  [22, 34]. Entrance into hibernation is associated with a temperature-dependent reduction of plasma leukocytes, with as many as

90 % sequestered in lymphatic organs [23]. Reduction in immune capacity leaves hibernating individuals susceptible to parasites and disease, but it also prevents the wide-scale recruitment of cytokines and consequent inflammatory response that typically follows cerebral ischemia [3, 159]. Immune responses, such as fever, are absent during hibernation but common immediately or shortly after individuals arouse [164].

Appetite is suppressed during hibernation, and in the absence of eating, the gut undergoes some reduction in mass. For example, in Alpine marmots, stomach and small intestine mass is 105 and 259 % greater during mid-summer than the end of hibernation, respectively [111]. In addition, no net flux of nutrients occurs in mucosal tissue during hibernation [37, 40]. However, despite mucosal atrophy and reduced mass of gastrointestinal organs during hibernation, absorptive capacity is retained. Intestinal tissue of hibernating thirteen-lined ground squirrels showed no net nutrient absorption at hibernation  $T_b$  (7 °C), but when warmed to euthermic  $T_b$  (37 °C), absorption was highest in tissue of fall and winter animals [37]. High capacity for absorption is expected during fall hyperphagia but retention throughout winter is not expected given the cost of the required protein synthesis. By sustaining high absorption capacity in less tissue, hibernators save energy on tissue maintenance and conserve the ability to resume digestive function when critical, such as during arousals (if the hibernator caches food and eats periodically during winter) and at the end of hibernation when body mass may be low and body condition affects other life history traits, such as reproduction (perhaps especially important for insectivorous hibernators that rely completely on fat reserves). Retention of absorptive capacity may be due to persistent expression of moesin in hibernating animals. Moesin is associated with formation of microvilli in the intestinal brush-border membranes of hibernating mammals, but is not found in euthermic animals or in adult nonhibernating mammals [38].

### Requirements for Rewarming and Arousal

Rewarming using endogenously generated heat is characteristic of torpor [114], and all small mammals, aside from tenrecs (*Tenrec ecaudatus* [137]) periodically arouse throughout hibernation [71, 203]. Restoration of neural function [163], mating [12], maintenance of muscle tissue [210], and access to food and water [112] have all been suggested as reasons for arousal. For whatever the reason, arousals play an essential role in hibernation biology and use a disproportionately large amount of energy [9, 56]. Over 80 % of energy reserves can be spent warming up and maintaining high  $T_b$  during arousals and euthermic bouts

[76]. Thus, it is essential that hibernators have mechanisms to make this process as efficient as possible. One such adaptation is the extensive use of brown adipose tissue (BAT) for rewarming.

### Non-shivering Thermogenesis

Non-shivering thermogenesis is largely dependent on BAT, which provides the predominant source of energy and heat during arousal from torpor [154]. Shivering is the most common method of producing heat in low temperatures but is not used in deep torpor because mechanisms that increase muscle function at low temperatures (e.g., a pool of inactive and available CK and HK) likely remain ineffective given the slowed rates of enzyme activity. A massive source of endogenous heat is needed to facilitate the shift in physiology and metabolic activity. Unlike the process of shivering, the energy released by oxidizing BAT is entirely heat [36]. Thus, the exclusively thermogenic result of oxidizing BAT is the key benefit of its use in arousal from hibernation.

Brown adipose tissue is under adrenergic control, and the  $\beta$ -adrenergic pathway in particular stimulates thermogenesis [36]. For reasons unknown, the signaling pathway that stimulates BAT thermogenesis in hibernators is more responsive as temperature drops. In Syrian hamsters (*Mesocricetus auratus*), stimulation of the  $\beta$ -adrenergic pathway at 36 °C produces no change in the metabolic activity of BAT in hibernating, nonhibernating, or cold-acclimated individuals, but stimulation at 12 °C results in an almost 200 % increase in hibernating animals and no change in the other groups [120]. Brown adipose tissue has a higher density of mitochondria than any other tissue and contains several smaller lipid droplets compared to a single large droplet as seen in WAT [36]. Peroxisome proliferator-activated receptor gamma coactivator 1-alpha (PGC-1 $\alpha$ ) is a co-regulator of metabolism that stimulates mitochondrial biogenesis and may play a key role in the number of mitochondria in bats. In little brown bats, PGC-1 $\alpha$  is upregulated during hibernation to a higher degree in BAT than all other tissues [58]. In addition to occurring in higher density, mitochondria in BAT are specialized for heat production by the uncoupling of ATP production. Thermogenins (UCP-1) are found in high density in the mitochondria of BAT and allow electrons to bypass ATPase while re-entering the mitochondrial matrix [121]. The electron transport chain found in the mitochondria of BAT is also adapted for increased activity. For example, in hibernating little brown bats, expression of genes for some components of the electron transport chain in the membranes of BAT mitochondria is upregulated by as much as 400 % compared to other tissues during hibernation and all tissues in

euthermic conspecifics [57]. Thus, the capacity for oxidation and heat production is increased on the scale of each individual mitochondrion as well as BAT tissue in general. Taken together, BAT provides most hibernators with a mechanism for increased heat production heat at relatively low  $T_b$  compared to other tissues, including WAT and muscle.

A lack of BAT does not necessarily preclude the use of hibernation. Echidnas (*Tachyglossus aculeatus*; Order: Monotremata) and four species of pygmy possum (family Burramyidae; Infraorder: Marsupialia) are known to hibernate (reviewed in [84]); yet all lack functional BAT [94]. The patterns of arousals during hibernation in these Australasian species are similar to those of other hibernators and frequent arousals are observed [84], yet it appears they use WAT to fuel arousals, the mechanisms of which remain unknown. The small difference between active and hibernating MR and  $T_b$  in echidnas [156] might make it possible to rewarm using only WAT and shivering thermogenesis. In addition, the range of these hibernators is restricted to Australasia, and roosting or nesting ecology may partly explain persistence of arousals during hibernation despite lack of BAT, as is the case with the sole avian hibernator. The common poorwill roosts in areas exposed to sunlight and uses radiant solar heating to passively rewarm during arousals [206], which may negate a reliance on an endogenous heat source. Considerable energy can be saved by passively rewarming [80], and a minimum  $T_b$  threshold might be reached in doing so that would allow mechanisms for shivering thermogenesis to become functional, thus negating the reliance on BAT for rewarming. Common tenrecs (*Tenrec ecaudatus*), which are protoendothermic eutherians that have BAT, are known to go months in hibernation without arousing [137]. Given that  $T_a$  did not fall below 22 °C during this study, the lack of arousals suggests the physiological costs of hibernating (e.g., neurological degeneration or accumulation of metabolites) at such temperatures may not be substantial enough to require frequent arousals in this species. Interestingly, BAT found in the lesser hedgehog tenrec (*Echinops telfairi*) is not concentrated in the intrascapular region, as is typical in mammalian hibernators, but rather is found in the abdominal region and ostensibly used to maintain high  $T_b$  during reproduction [158]. The focused distribution of BAT in tenrecs may explain their inability to endogenously produce heat for arousal during hibernation. Although there are other adaptations that allow few hibernating animals to avoid the need for BAT, the predominant hibernation phenotype is characterized by long-term, deep, seasonal hibernation interspersed with frequent arousals, which are largely fueled by BAT.

## Conclusion

Phenotypic plasticity is common, and hibernation is arguably one of the most profound examples among mammals. Adaptations associated with hibernation are multifaceted and allow for increased energy storage, pronounced metabolic depression and reduced  $T_b$ , selective maintenance of vital biological processes and cellular protection in an ischemic and hypothermic state, and the means for complete reversal to a euthermic state, all with virtually no pathological consequence. Hibernators become hyperphagic and considerably increase energy intake during the pre-hibernation period by way of seasonal resistance to hormones that typically inhibit appetite, dissociation of key hormones from providing adiposity feedback, and concomitant stimulation of lipid synthesis pathways. The switch in metabolic fuel during hibernation is facilitated by a decreased availability of carbohydrates and increased amounts of available lipids. Key enzymes that halt the flow of carbohydrates into metabolic pathways and stimulate the release of fatty acids for oxidation are upregulated during hibernation. Entrance into hibernation is influenced by photoperiod, temperature, and energy availability, and adaptations to the circadian system maintain semi-regular arousal patterns. However, the underlying intrinsic mechanism remains poorly understood.

The main purpose and consequence of hibernation is energy savings. Temperature-dependent effects on enzyme activity inherently decrease the rate, and therefore energy demand, of biological processes with falling  $T_b$ , but ATP turnover is further reduced physiologically. RPP plays a key role in activating and deactivating metabolic enzymes and substrates, such as PDH. Phosphorylation of enzymes involved in carbohydrate catabolism reduces the amount of substrate available for oxidation and consequently the rate of mitochondrial respiration and energy consumption. A large proportion of ATP turnover is also associated with maintenance of ion gradients across cellular membranes, and RPP of membrane proteins reduces activity and energy consumption, further facilitating metabolic reduction. The process of RPP is reversible given that it is a temporary modification and does not permanently change the substrate. Therefore, the pool of enzymes and proteins remains readily available and functional, even at euthermic temperatures. Although protein synthesis and biological processes are largely diminished and some even stopped during hibernation, tissue-specific changes maintain vital functions and protect tissues during this period of ischemia and hypothermia. For example, increased expression of SERCA2a in cardiac muscle facilitates cardiac function at lower temperatures [87], and mRNA is structurally changed to increase its stability during hibernation [122].



Membrane permeability is also maintained at a minimally functional level by shifts in the type of membrane fatty acids, an adaptation that avoids a counterproductive PUFA-associated increase in energy consumption. In addition, processes that normally occur during long periods of inactivity and lead to tissue damage are not evident during hibernation (e.g., lack of myostatin in skeletal muscle). Through a combination of protective adaptations and the ability to mute the body's normal, damaging responses, hibernators are able to mitigate the costs of low  $T_b$ , ischemia, and inactivity to realize the energetic benefits of quiescence and metabolic reduction.

The most intriguing aspect of hibernation is that adaptations for it involve modifications of existing physiological mechanisms that likely occur in all mammal lineages. Genes differentially expressed during hibernation that underlie the changes involved in hibernation physiology occur in all mammal lineages. Therefore, hibernation can be considered a seasonally expressed phenotype involving adaptive regulation of a number of pathways and processes that are evolutionarily conserved across all mammalian taxa. In fact, nonhibernators show similar responses to hypothermia and ischemia, as well as seasonal fluctuations in metabolism. WAT in some nonhibernators adapts to the cold by activating "brite" adipocytes, which express UCP-1 and increase thermogenic capacity [133]. Likewise, hypometabolism becomes evident and insulin resistance increases in some human populations in response to prolonged exposure to cold during winter [117]. However, hypothermic responses of nonhibernators are acute and typically differ from those of true hibernators, and are often followed by pathological consequences. Nonhibernators incur damage during periods of ischemia and hypothermia likely because they do not possess the entire suite of adaptations to cold and metabolic reduction as in hibernators. Physiological reductions to metabolism, maintenance of vital functions and protection of tissues in cold temperatures, and a means to warm up endogenously to resume function at normal temperatures are all necessary for the hibernation phenotype. However, some studies suggest an embryonic state of gene expression is achieved during hibernation. For example, moesin is only expressed in the enterocytes during fetal and hibernation states [38], and pancreatic lipase is found only in the pancreas of nonhibernators but throughout other tissues as well in fetuses of hibernators [7, 189]. Although heterothermy likely evolved separately in birds and mammals [77], there are examples of bird species using torpor early in development as altricial young but not in adulthood (e.g., [18]) and, with further study, we may find the same pattern in nonhibernating mammals. Perhaps many more mammals than we know are born with the ability to hibernate and to lose it with maturity. With a better understanding of the

mechanisms involved in hibernation, at least some of the benefits seen during torpor, such as resistance to metabolic disease despite adiposity (e.g., [155]), may be achievable in nonhibernators (e.g., humans), a prospect that adds urgency and excitement to further studies in hibernation physiology.

## Glossary

Daily torpor	Short-term employment of torpor characterized by bouts typically lasting less than 24 h with body temperature ( $T_b$ ) often remaining some degrees above ambient temperature ( $T_a$ ).
Endothermy	Regulation of $T_b$ using a high rate of metabolism to produce endogenous heat.
Euthermy	Maintenance of $T_b$ at a relatively high temperature conducive to normal biological function.
Heterothermy	The diel or seasonal pattern of temperature regulation where $T_b$ varies outside the normal euthermic range
Hibernation	Extended employment of torpor characterized by bouts typically lasting days to weeks with $T_b$ dropping to near $T_a$ . Obligate hibernation lasts for months during predictable periods of energy imbalance (the "hibernation season") and consists of multiple bouts interspersed with regular arousals. Conversely, facultative hibernation typically occurs during ephemeral energy crises outside of any predictable season
Hypothermia	An uncontrolled state of reduced $T_b$ (i.e., heat loss exceeds capacity for heat production)
Torpor	A temporary and controlled reduction in metabolic rate (MR) and body temperature ( $T_b$ ) followed by a return to euthermy using an endogenous source of heat. Torpor is often employed as an adaptive response to conserve energy during periods of resource scarcity

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