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**RESEARCH PAPER** 

# Hibernation strategy – related profound differences in the whole-body fat composition of bats

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**Abstract.** Bats can use a wide range of roosts as hibernacula, resulting in diverse hibernation strategies. The ecological needs of a species during hibernation translate into particular torpor-arousal patterns and physiological demands. For mammalian hibernators, the oxidation of fatty acids from triacylglycerols stored in white and brown adipocytes provides the main energy to fuel hibernation. The relative content of saturated, monounsaturated, and polyunsaturated fatty acids in body fat brings multifarious costs and benefits, and their importance during hibernation is likely changing. While considering the level of fatty acid saturation and their properties, we hypothesised that whole-body fat composition varies between bat species (*Nyctalus noctula, Myotis myotis*) that employ different hibernation strategies. Therefore, the focus of this study was to determine the relative fatty acid composition of the whole-body fat of these species. We found evidence that the body fat of *N. noctula* has a higher relative content of MUFAs than *M. myotis*, which, on the other hand, has high SFAs and PUFAs. Such profound differences in fatty acid profiles suggest that the studied species' distinct hibernation strategies and torpor-arousal patterns are reflected in functional differences.

Key words: energy reserves, fatty acids, PUFA, MUFA, SFA

# Introduction

Torpor and hibernation represent powerful strategies enabling animals to cope with periods of low food availability and unsuitable environmental conditions (e.g. short photoperiod or challenging weather). For seasonal hibernators, entrance into hibernation is anticipated several weeks in advance by changes in behaviour and physiology that lead to the accumulation of energy stores (Kunz et al. 1998, Speakman & Rowland 1999). These changes are reflected in an adjustment of the thermoneutral zone and a decrease in basal metabolic rate (BMR) (Ruf & Geiser 2015). During hibernation, individuals achieve a minimum torpid metabolic rate of 4% of BMR and a variable reduction of their body temperature ( $T_b$ ), ranging on average for most species between 0 and 10 °C (Ruf & Geiser 2015). Hibernation corresponds to multiple and successive torpor bouts lasting for days to a few weeks, during which animals rely entirely on fuel stores, such as body fat and/or food caches (Humphries et al. 2003, Dark 2005). Torpor bouts are periodically interrupted by brief arousals to a euthermic state, which consumes up to 85% of winter energy stores (Thomas et al. 1990). As torpor entails several potential costs (Humphries et al. 2003,

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\* Corresponding Author Downloaded From: https://bioone.org/journals/Journal-of-Vertebrate-Biology on 17 Nov 2024 Terms of Use: https://bioone.org/terms-of-use Boyles et al. 2020), torpor expression and quantitative energetics are influenced by species, environmental conditions, and individual states, including available energy reserves (Boyles et al. 2007, Czenze et al. 2017, Blažek et al. 2019, Bachorec et al. 2021, E. Bachorec, unpublished data). The main energy source during periods of food scarcity is stored in the white and brown adipose tissue as triacylglycerols (TGA). In addition to an increased mass of both types of adipose tissue (Feist et al. 1986, Kunz et al. 1998, McGuire et al. 2009), heterothermic mammals also increase the proportion of polyunsaturated fatty acids (PUFA) in their body fats prior to entering hibernation (Schalk & Brigham 1995). The seasonal increase in PUFAs, which have a characteristic of low melting points, is important in maintaining the fluidity of depot fats for fuelling metabolism during torpor, as well as the fluidity of membrane phospholipids at low body temperatures (Irving et al. 1957, Mead 1986, Aloia 1988). Besides the effect of the total amount of fat reserves on torpor expression in terms of optimal hibernation theory (Boyles et al. 2020), the fatty acid composition of body fat also influences the utilisation of torpor and metabolic rate (MR) of animals. Previous studies concluded that an increased relative proportion of PUFAs in fat reserves of hibernators resulted in decreased  $\mathrm{T}_{\mathrm{b}}$  and prolonged torpor bouts and ultimately increased survival during the hibernation season (Geiser et al. 1990, 1992, Frank 1992, Florant et al. 1993). Some evidence also indicates that monounsaturated fatty acids (MUFA) may positively affect torpor, i.e. longer torpor bouts, lower T<sub>b</sub> and MR (Frank & Storey 1996, Geiser et al. 1994). On the other hand, depot fat, rich in saturated fatty acids (SFA), increases metabolic rate and  $T_{\mu}$  which may explain differences in MR among torpid animals (Geiser 1993). Additionally, fatty acids are not uniformly mobilised from adipocytes; some are preferentially mobilised while others are preferentially retained (Raclot & Groscolas 1993, Price et al. 2013). Several studies have observed that short-chain SFAs and fatty acids with one or more double bonds are mobilised and metabolised quicker in fish, birds, and mammals (Raclot & Groscolas 1995, Sidell et al. 1995, Price et al. 2008, 2013). For mammalian hibernators, the synthesis of MUFAs and the specific retention of PUFAs acquired through feeding might be the only way to increase the proportion of unsaturated fatty acids in their fat depot. Insectivorous bats, which have access to only low levels of PUFA, should select their insect prey to maximise PUFA intake (Schalk & Brigham 1995). Voigt et al. (2019) argue that the diet of most insectivorous bats is low in PUFA content. Thus consumption of a diet enriched in MUFA and

synthesis of MUFA are mechanisms to incorporate unsaturated fatty acids in depot fat, as suggested for hibernating echidnas (Falkenstein et al. 2001).

Hibernation, however, can be employed in various forms. Bats can use a wide range of natural and artificial structures as roosts. While caves provide a stable thermal microclimate for hibernating bats such as greater mouse-eared bats Myotis myotis (Zukal et al. 2005, 2017), tree roosts and crevices in buildings used by common noctule bats Nyctalus noctula provide much less insulation from the external daily T<sub>a</sub> cycle (Sluiter et al. 1973, Turbill 2006). Nyctalus noctula have been observed to hibernate in road bridge crevices at T<sub>a</sub> of -13 °C (Cel'uch & Ševčík 2008). Situations like this often force bats to arouse from torpor and move to a proper shelter. As such, noctules are well adapted for hibernation in harsh microclimatic conditions. Additionally, in contrast to *M. myotis*, *N. noctula* is noticeable by flight and foraging activity during winter (also at  $T_2 < 0$  °C) (Gaisler et al. 1979, Avery 1986, Cel'uch & Kaňuch 2005, Kaňuch et al. 2005). Therefore, comparing the relative body fat composition of two species with different ecology and roost requirements seems to be a good model for testing proximate mechanisms of thermal response and hibernation patterns. We hypothesise that the two species will differ in the whole-body fatty acid composition. Regarding the character of hibernation strategy, we predict that N. noctula would benefit from high content of easily and rapidly metabolizable fatty acids (MUFA, PUFA), while M. myotis would rely on energy-dense fatty acids (SFA).

## **Material and Methods**

## Sample preparation and analysis

Whole body fatty acid composition was compared across 18 common noctule bats and 14 greater mouseeared bats (Table S1). All bats died during the second part of deep hibernation season (late February-mid March). *Nyctalus noctula* cadavers were collected within a few days after disposing of them from their shelter in a public building. *Myotis myotis* cadavers were collected within a 3-week period during regular data collection visits (Bachorec et al. 2021). Before that, bats were naturally hibernating and fasting. As we only received the cadavers, we could not influence the sampling design and used the material available.

Sample preparation and analysis took place during the summer of 2022. Cadavers without heads and wings were homogenised, and 16 g samples were prepared from each. Total fat content was determined gravimetrically, where samples (3-5 g) were acid



Fig. 1. Principal component analysis based on the whole-body fatty acid proportions of two bat species grouped by the level of saturation.

hydrolysed via 50 ml of boiling diluted hydrochloric acid (4 mol/l) for lipids to be released. Afterwards, the hydrolysed sample was filtered and dried for 1 hour at 103 °C. The fat extraction procedure was performed using the Soxtec 2050 (FOSS, Denmark) petroleum ether extractor. For the determination of fatty acids, extracted fat samples were esterified to fatty acid methylesters via transmethylation using methanolic solution of potassium hydroxide (2 mol/l). Extracted fatty acid methylesters were separated, identified, and quantified using high-performance capillary gas chromatography with FID detector (HP5890A, Germany). The fat and fatty acid analyses were performed by an external company (State Veterinary Institute, Czech Republic) following ISO standards (ISO-1443; ISO-12966-2; ISO-12966-4).

#### **Statistical analyses**

Fatty acids that showed no variability and/or had a low proportion relative to the total fat profile were excluded from the comparison. These included caproic (0.01%), capric (0.02%), caprylic (0.01%), nervonic (0.04%), and undecanoic acid (0.01%). Principal components analysis (PCA) via '*prcomp*' function was employed to summarise the variation and detect distinction in the fatty acid profiles of whole-body fat between

N. noctula and M. myotis. PCA was performed on all fatty acids pooled into three categories based on the saturation level (SFA, MUFA, PUFA) and the specific fatty acids in these categories alone. As our data are in the form of proportions relative to the total fatty acid profile, a non-parametric test was used to test the differences between the two species. Therefore, we used npMANOVA using the 'adonis2' function from the 'vegan' package (Oksanen et al. 2013), which is a permutational analysis using distance matrices and is appropriate for comparing relative proportions. To assess differences between species, a comparison was carried out on all fatty acids pooled together and with fatty acids grouped into categories based on the saturation level. Analyses were performed in R Studio (R Core Team 2022).

# Results

Oleic acid had the highest proportion to the total fatty acid profile, ranging from 49.7-71.3% in *N. noctula* and 32.4-59.8% in *M. myotis*. On average, a high proportion was also observed in linoleic acid (13.9 and 12.6%), palmitic acid (9.4 and 12.7%) and stearic acid (4.3 and 9.4%) in *N. noctula* and *M. myotis*, respectively. The mean values of the relative fatty

**Table 1.** Relative fatty acid content (mean % to total fat) and PCA scores for fatty acids grouped by the level of saturation.  $N_{N.noct} = 18$ ;  $N_{M.mvo} = 14$ .

|        | FA                          |          | $\bar{X}N$ . noctula | $\bar{X}M$ . myotis | PC1    | PC2    |
|--------|-----------------------------|----------|----------------------|---------------------|--------|--------|
|        |                             |          | n = 18               | n = 14              |        |        |
| SFA    | lauric                      | C12.0    | 0.112                | 0.074               | -0.258 | -0.490 |
|        | tridecanoic                 | C13.0    | 0.124                | 0.024               | -0.267 | -0.321 |
|        | myristic                    | C14.0    | 0.919                | 0.799               | -0.095 | -0.673 |
|        | pentadecanoic               | C15.0    | 0.054                | 0.236               | 0.197  | 0.213  |
|        | palmitic                    | C16.0    | 9.419                | 12.744              | 0.249  | -0.001 |
|        | heptadecanoic               | C17.0    | 0.132                | 0.696               | 0.357  | -0.084 |
|        | stearic                     | C18.0    | 4.294                | 9.440               | 0.328  | -0.050 |
|        | arachidic                   | C20.0    | 0.247                | 1.176               | 0.351  | -0.199 |
|        | heneicosanoic               | C21.0    | 0.013                | 0.354               | 0.332  | -0.198 |
|        | behenic                     | C22.0    | 0.056                | 0.776               | 0.341  | -0.197 |
|        | tricosanoic                 | C23.0    | 0.927                | 2.656               | 0.281  | -0.085 |
|        | lignoceric                  | C24.0    | 0.160                | 1.502               | 0.297  | -0.152 |
| MUFA   | myristoleic                 | C14.1_5  | 0.167                | 1.872               | -0.043 | 0.774  |
|        | palmitoleic                 | C16.1_7  | 4.937                | 2.176               | 0.395  | 0.479  |
|        | cis-10-heptadecenoic        | cC17.1_9 | 0.111                | 0.201               | -0.162 | -0.097 |
|        | oleic                       | C18.1_9  | 61.942               | 46.658              | 0.486  | -0.155 |
|        | elaidic                     | tC18.1_9 | 0.083                | 0.224               | -0.407 | 0.225  |
|        | gondoic                     | C20.1_9  | 0.345                | 1.441               | -0.526 | 0.175  |
|        | erucic                      | C22.1_9  | 0.044                | 0.295               | -0.369 | -0.236 |
| PUFA   | $\alpha$ -linolenic         | C18.3_3  | 0.968                | 0.991               | 0.295  | 0.013  |
|        | γ-linolenic                 | C18.3_6  | 0.023                | 0.214               | 0.308  | -0.030 |
|        | linoleic                    | C18.2_6  | 13.881               | 12.262              | -0.035 | 0.157  |
|        | linolelaidic                | tC18.2_6 | 0.054                | 0.117               | 0.321  | -0.405 |
|        | cis-11,14,17-eicosatrienoic | cC20.3_3 | 0.018                | 0.064               | 0.350  | 0.389  |
|        | cis-11,14-eicosadienoic     | cC20.2_6 | 0.131                | 0.783               | 0.315  | 0.464  |
|        | dihomo-γ-linolenic          | C20.3_6  | 0.071                | 0.143               | 0.293  | 0.463  |
|        | arachidonic                 | C20.4_6  | 0.013                | 0.216               | 0.341  | -0.324 |
|        | EPA                         | C20.5_3  | 0.101                | 0.154               | 0.312  | -0.049 |
|        | cis-13,16-docosadienoic     | C22.2_6  | 0.012                | 0.061               | 0.315  | -0.229 |
|        | DHA                         | C22.6 3  | 0.556                | 0.944               | 0.304  | -0.263 |
| pooled | SFA                         | —        | 16.496               | 31.017              | 0.634  | -0.423 |
|        | MUFA                        |          | 67.664               | 51.245              | -0.694 | 0.056  |
|        | PUFA                        |          | 14.973               | 17.731              | 0.340  | 0.904  |

acid content and PCA scores are shown in Table 1. The PCA based on fatty acid composition showed separation of two species regarding the proportion of SFA and MUFA with PC1 and PC2 explaining 98.9% of variability (Fig. 1). Clear separation of two species where with PC1 and PC1 explained 68.3% of variability was also observed in proportions of the specific SFAs, with the short chain SFAs (C12-C14) representing *N. noctula* and the long chain SFAs

(C15-C24) representing *M. myotis* (Fig. S1). The two bat species were clearly separated regarding MUFAs, with 71.8 % variability explained by the first two PCs (Fig. S2). A positive correlation of oleic and palmitoleic acid with PC1 characterises *N. noctula*, while the negative correlation of elaidic, erucic, gondoic and cis-10-heptadecenoic acid represents *M. myotis*. The principal component analysis of PUFAs composition, which explained 61% of variability on PC1 and PC2,



**Fig. 2.** Proportions of the whole-body fatty acids as percentage relative to the total fat grouped into categories by the level of saturation.

shows a positive correlation of all PUFA types with *M. myotis* and practically all *N. noctula* are centralised within negative PC values (Fig. S3). All PCA based on fatty acid composition explained more than 60.0% of the total variability.

The npMANOVA supported these distinctions by providing evidence of profound differences in the whole-body fatty acid composition between the two bat species under study. Species showed a significant effect on the fatty acid composition when proportions of all fatty acids were pooled ( $F_{1,28}$  = 19.416; partial R<sub>2</sub> = 0.409; *P* < 0.001). When fatty acids were divided into categories (SFA, MUFA, PUFA, Fig. 2), the npMANOVA showed strong evidence that the proportion of SFAs ( $F_{1,28}$  = 20.567; partial  $R_2$  = 0.423; P < 0.001) was higher in *M. myotis* and that the proportion of MUFAs ( $F_{1,28}$  = 23.549; partial  $R_2$  = 0.456; P < 0.001) was higher in *N. noctula*. This difference is present, however weaker, also in the case of PUFAs  $(F_{1.28} = 4.76; partial R_2 = 0.145; P < 0.05)$ , with M. myotis showing a slightly increased proportion of PUFAs.

## Discussion

We compared the whole-body fatty acid composition between two temperate bat species that use different hibernation strategies. Consistent with our prediction, the results show profound differences in the fatty acid composition of these two species. The fatty acid composition of body fat has been shown to reflect that of diet (Florant 1998, Falkenstein et al. 2001), although processes such as post-absorptive modification and *de-novo* synthesis cannot be ignored (Price 2010). It is known that *N. noctula* feeds mainly on Diptera, which are low in PUFAs, although Coleoptera and Lepidoptera are also important components of its diet (Schalk & Brigham 1995, Vaughan 1997), while M. myotis feeds primarily on large Coleoptera species (mainly Carabidae and Scarabaeidae) richer in PUFA (Arlettaz & Perrin 1995, Beck 1995, Schalk & Brigham 1995, Andreas 2002). Foraging habits and prey selection, especially during the pre-hibernation season, might result in divergent body fat composition to meet the specific needs of animals during hibernation (Schalk & Brigham 1995, Kunz et al. 1998).

Differences demonstrated by our results raise questions about differential energy availability and utilisation rate between the studied species during winter. Analysis of the whole-body fatty acid composition revealed that the proportion of SFAs to the total body fat was higher in *M. myotis*. For hibernating animals, SFAs present the main energy source (Florant et al. 1990, Geiser 1991, Price et al. 2013). Long-chain saturated fatty acids are energetically dense. However, they are mobilised and metabolised slower than unsaturated and shorter fatty acids (Price & Gugliemo 2009, Price 2010). The difference in SFA content likely reflects the energetic needs for a different length of the hibernation period which is shorter for N. noctula (mid-December-March) (Cel'uch & Kaňuch 2005, Zahn & Kriner 2016) compared to M. myotis (mid-November-mid-April) (Zukal et al. 2005, 2017). The higher content of energetically dense saturates in the body fat of *M*. myotis, conspicuous by deep, prolonged hibernation (Zukal et al. 2017), might thus represent a longterm energy store to fuel the overwinter demands. However, because body fats must remain fluid to be metabolised at low  $T_{b'}$  some level of unsaturation is needed, which is attained via specific retention of fatty acids with one or more double bonds (Irving et al. 1957, Mead 1986, Aloia 1988). We observed a high proportion of MUFAs (especially oleic acid) in both species. A high proportion of oleic acid in N. noctula was also observed by Voight et al. (2019), and the same was reported for several other bat species (Ewing et al. 1970, Levin et al. 2013, McGuire et al. 2013) as well as echidnas (Tachyglossus aculeatus) by Falkenstein et al. (2001). As one double bond has the most profound effect on fatty acid fluidity (Cossins & Lee 1985), it appears that high content of MUFAs can compensate for low dietary PUFA. It is the most likely way to cope for insectivores, which typically have a low PUFA content in their diet compared to herbivorous hibernators (Falkenstein et al. 2001). Moreover, high PUFA and MUFA content in body fat has positive effects on torpor expression (Geiser et al. 1990, 1992, 1994, Frank 1992, Florant et al. 1993, Frank & Storey 1996, Falkenstein et al. 2001) and thus improve energy conservation. Evidence of higher MUFA content at the expense of SFAs revealed by PCA and supported by MANOVA possibly accounts for the specific needs of *N. noctula*. This species hibernates in harsh, unstable conditions that may compel them to arouse and fall into torpor more frequently and quickly (Sluiter et al. 1973, Cel'uch & Kaňuch 2005, Kaňuch et al. 2005). Selective mobilisation of fatty acids could notably affect the supply of tissues and organs with specific fatty acids in situations of negative energy balance (Raclot & Groscolas 1993). We suggest that a higher content of MUFAs provides a rapid energy source for thermogenesis during short and mild arousals of N. noctula. Furthermore, hibernation shelters used by N. noctula often allow them to benefit from passive rewarming. By hibernating in a roost exposed to the daily T<sub>2</sub> cycle, they can rewarm more effectively and

save considerable energy during arousals that occur during the increased T<sub>a</sub> (Turbill & Geiser 2008). On the other hand, the slightly higher content of PUFAs in *M. myotis* is likely to fuel body tissues during long torpor bouts. Although BMR does not differ much between the studied species, N. noctula has a higher  $Q_{10}$  (3.4) compared to *M. myotis* (3.0) (Geiser 2004). This difference suggests that metabolic processes, including warming up during arousal, should be faster than in M. myotis. As reported by Rosner & Voight (2018), N. noctula rewarms in 40 min, with a peak MR after 25-28 min, faster than M. myotis, which rewarms after 60 min with peak MR at 35 min (E. Bachorec, unpublished data). However, a bigger differential between the species might be expected.

Altogether, a high MUFA content relative to the total fatty acid profile appears to be a general trait of hibernating insectivores. It provides hibernators with a constitution of fat that has suitable physical properties for deep torpor and periodic arousals. However, as our results showed, the fatty acid profile might vary with different hibernation strategies. The endogenous synthesis of MUFAs is independent of diet and seems to be an efficient adaptive mechanism that provides hibernators with a low PUFA diet to survive the dramatic changes in physiology during overwintering without additional fat restructuration. Even though our sampling was not standardised and might have accounted for some variance, the results presented in this study contribute to the knowledge about the fatty acid composition of hibernating bats and provide more detailed insight and improvement in modelling energetic costs during hibernation.

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# **Author Contributions**

E. Bachorec, J. Pikula and J. Zukal conceived and designed the study; J. Zukal and K. Zukalová collected material, with support from J. Pikula; K. Zukalová and V. Seidlová performed the laboratory analyses; E. Bachorec analysed the data; E. Bachorec and J. Zukal drafted the manuscript, to which all authors contributed with critical comments.

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# **Supplementary online material**

**Table S1.** Dataset supporting the study's findings (proportions of fatty acids relative to whole body fat) (https://www.ivb.cz/wp-content/uploads/JVB-vol.-72-2023-Bachorec-et-al.-Table-S1.xlsx).

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Principal component analysis based on the whole-body fatty acid proportions of two bat species: **Fig. S1**. Saturated fatty acids, **Fig. S2**. Monounsaturated fatty acids, **Fig. S3**. Polyunsaturated fatty acids (https://www.ivb.cz/wp-content/uploads/JVB-vol.-72-2023-Bachorec-et-al.-Fig.-S1-S2-S3.pdf).