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Stable hydrogen isotope measurements of songbird feathers: effects of intra-feather variability and sample processing

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Abstract

Deuterium composition of feathers has been widely applied to establish provenance of birds. Surprisingly, little attention has been paid to the potential sources of unwanted hydrogen isotopic variability that may affect the comparability and replicability of deuterium measurements. Feathers may exhibit hydrogen isotopic differences between distal and proximal sections, as well as between rachis and vane. For this reason, it is recommended to always analyse ground vane from the same section of the feather. However, this protocol may be seriously limiting when working with songbird feathers, which are usually small and light. In this paper, intra-feather hydrogen isotopic variability was studied in four species of songbirds by comparing four sections of the same feather. No difference in hydrogen isotopic composition was found between distal and proximal sections of the vane. This was expected as songbird feathers grow rapidly and individuals avoid geographical movements during moult, favouring a homogeneous source of hydrogen during keratin synthesis. Conversely, as previously demonstrated for other avian species, rachis showed systematically c. 11% δ^2 H more negative values than vane. This is probably a result of their biochemical and ultrastructural differences, which seem strongly conserved across taxa. Such differences imply that we need to describe which feather part was used in the methods of our reports to ensure full comparability and replicability among studies. Nevertheless, vane and rachis δ^2 H were strongly correlated, demonstrating that rachis can be also used in isotope analyses because its δ^2 H can be accurately rescaled. This is useful since rachis represents an important fraction of feather mass in songbirds. Values of δ^2 H from ground and non-ground samples from the same feather were also compared. There was no effect of the processing protocol, suggesting that grinding is unnecessary when working with passerine feathers.

Keywords Deuterium · Keratinous material · Methodology · Passerines · Protocol · Tissue

Zusammenfassung

Messungen stabiler Wasserstoffisotopen in Singvogel-Federn: Effekte der Intra-Federn-Variabilität und Probenverarbeitung

Die Deuteriumzusammensetzung von Federn wird üblicherweise eingesetzt, um die Herkunft von Vögeln festzustellen. Überraschenderweise wird dabei aber nur wenig auf potenzielle Quellen einer unerwünschten Wasserstoffisotopen-Variabilität geachtet, die die Vergleichbarkeit und Wiederholbarkeit der Deuteriummessungen beeinflussen können. Federn können zwischen ihren distalen und proximalen Abschnitten sowie zwischen Wirbelsäule und Flügeln Unterschiede in den Wasserstoffisotopen aufzeigen. Deshalb empfiehlt es sich, Analysen stets von den gleichen Federabschnitten zu machen. Diese Regel führt jedoch bei Singvogelfedern zu einer ernsthaften Einschränkung, weil diese normalerweise klein und leicht sind. In dieser Studie untersuchten wir bei vier Singvogelarten die Wasserstoffisotopen-Variabilität in den gleichen Federn, indem vier Abschnitte derselben Feder verglichen wurden. Zwischen dem distalen und proximalen Abschnitt desselben

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Flügels wurden keine Unterschiede in der Wasserstoffisotopen-Zusammensetzung gefunden. Das war erwartet worden, weil Singvogelfedern schnell wachsen und die einzelnen Tiere Ortsveränderungen während der Mauser vermeiden und somit während der Keratin-Synthese eine homogene Wasserstoffversorgung haben. Andererseits, wie schon früher für andere Vogelarten festgestellt, zeigt der Schaft systematisch ca. 11% δ^2 H mehr negative Werte als die Federfahne. Vermutlich liegt das an biochemischen und ultrastrukturellen Unterschieden, die sich durch alle Taxa hindurchziehen. Diese Unterschiede machen es empfehlenswert, immer anzugeben, welche Federabschnitte verwendet wurden, um eine volle Vergleichbarkeit und Wiederholbarkeit der Untersuchungen zu gewährleisten. Nichtsdestotrotz korrelierten die δ^2 H von Schaft und Fahne hoch miteinander und zeigten, dass auch der Schaft für die Isotopenanalyse benutzt werden kann, weil seine δ^2 H-Werte entsprechend korrekt skaliert werden können. Das ist deshalb sehr hilfreich, weil bei Singvögeln der Schaft einen wichtigen Anteil der gesamten Federmasse darstellt. Die δ^2 H-Werte von zermahlenen und nicht zerkleinerten Proben der gleichen Federn wurden ebenfalls untersucht und kein Unterschied festgestellt, was nahelegt, dass bei der Arbeit mit Singvogelfedern ein Zermahlen der Federn unnötig ist.

Introduction

The study of stable isotopes has become a popular tool in trophic and movement ecology research (Hobson 1999; Rubenstein and Hobson 2004; Bowen et al. 2005; Hobson 2011; Meier-Augenstein et al. 2013). In birds, stable isotope measurements have been successfully used to trace the origin of migrants (Dunn et al. 2006; Cárdenas-Ortiz et al. 2017), establish the connectivity between breeding and nonbreeding populations of migratory species (Chamberlain et al. 1996; Hobson and Wassenaar 1997; Marra et al. 1998), unravel the diet of species difficult to track (Procházka et al. 2010; Robillard et al. 2017), discover historical changes in food webs (Emslie and Patterson 2007; Blight et al. 2015), and determine nutrient allocation during reproduction (Klaassen et al. 2001; Jaatinen et al. 2016). Therefore, by selecting suitable element(s), tissue(s) and sampling period(s) within the life cycle, stable isotope research can solve a broad range of essential questions in bird ecology.

Feathers have been regularly used in stable isotope research due to a number of advantages (Smith et al. 2003; Bortolotti 2010). They are replaced regularly and moult strategies are usually known (e.g., Jenni and Winkler 2011), thus becoming an excellent tissue to have a snapshot of the isotope composition of the consumed diet, occupied habitat and geographical location. Furthermore, they are easy to collect and store (McDonald and Griffith 2011), and by being inert, they maintain the chemical signature of the environment where they grew (Wassenaar 2019). In addition, feathers can also be used as a source of DNA, hormones, and to quantify colouration (Katzner et al. 2012), providing further information of the studied individuals, thus improving our understanding of ecological and evolutionary processes of interest.

In spite of the widespread application of feather stable hydrogen isotope analyses, little research has been published on the methodological aspects of its measurement (Wassenaar and Hobson 2006; Smith et al. 2008, 2009; Paritte and Kelly 2009; Wiley et al. 2010; Bontempo et al. 2014; Soto

et al. 2017; Wassenaar 2019). This is surprising given the lack of a unified approach to collect and prepare feathers for posterior isotope-ratio mass spectrometry (but see Wassenaar 2019). This heterogeneity in methodologies may affect replicability and comparability among studies (Jardine and Cunjak 2005; Lott and Smith 2006; Wassenaar and Hobson 2006; Paritte and Kelly 2009; Qi and Coplen 2011; Meier-Augenstein et al. 2013; Bontempo et al. 2014).

Feathers are not a homogenous tissue and thus, some intra-feather variability in isotope composition should be expected. For this reason, the section of the feather selected for the analysis can affect the comparability of deuterium measurements between samples. Feathers are composed of a vane and rachis, which have different ultrastructure and density that can affect isotope deposition (Bortolotti 2010; Grecian et al. 2015). In fact, previous studies demonstrated that rachis normally is more depleted and variable in deuterium than vane (Wassenaar and Hobson 2006; Bontempo et al. 2014). For this reason, it is recommended to exclude rachis from analyses. In addition, some deuterium variability has been observed between the distal and proximal sections of feathers (Wassenaar and Hobson 2006; Smith et al. 2008, 2009; Wiley et al. 2010; Bontempo et al. 2014; Wassenaar 2019), suggesting changes in the source of hydrogen used to synthesize keratin during feather growth. To avoid this problem, one should always select the same section of the feather (Kelly et al. 2002; Smith and Dufty 2005; Pérez and Hobson 2006; Smith et al. 2008, 2009).

Elimination of the rachis and selection of a small vane section are feasible approaches in large birds, in which rachis and vane are patently different and a stamp-size piece of vane may provide several mg of material, which is enough for several analyses (e.g., several elements and/or replicates). However, in many songbirds, even full flight feathers may be really tiny and light (<3 mg). From such feathers, once the rachis is removed, we might barely obtain 1 mg of keratinous material. One mg is a quite limiting amount of sample, as each isotopic analysis needs around 0.3 mg. This challenge has not limited the application of hydrogen stable isotopes



research on songbirds (e.g., Chamberlain et al. 1996; Marra et al. 1998 and Pérez and Hobson 2006). Nevertheless, intrafeather isotope variability in this group should be properly described to elucidate whether or not protocol recommendations cited previously can be flexible and consequently, different sections of the same feather can be used without risk of getting biased results. This issue seems especially relevant, since the vast majority of studies to date about intra-feather isotope variability focused on flight feathers of non-passerines (Wassenaar and Hobson 2006; Smith et al. 2008, 2009; Wiley et al. 2010; Bontempo et al. 2014; Grecian et al. 2015).

Another step during feather processing can be to grind the samples to powder. Reference keratinous materials for the analysis are powdered, thus samples in the same state would ensure full comparability (principle of identical treatment; Chesson et al. 2009; Wassenaar 2019). In fact, tests with mammal hair have demonstrated some effects of particle size on the final measured deuterium abundance (Chesson et al. 2009; Qi and Coplen 2011). Apparently, powdered vs. nonpowdered materials would have different hygroscopic properties that affect their capacity to retain residual moisture content, thus having an effect on the exchangeable fraction of hydrogen (Chesson et al. 2009; Qi and Coplen 2011). Surprisingly, no similar tests have been carried out with feathers, therefore, it is unknown to what extent the findings with mammal hair are generalizable to other keratinous materials. To challenge the principle of identical treatment in this particular step of the protocol seems especially useful when working with songbird feathers, which are usually small. In this case, grinding can be a serious issue because a fraction of the sample is unavoidably lost during the process (e.g., small barbules adhered on the manipulation tools by static electricity). Therefore, it would be necessary to investigate whether or not entire (small) pieces of feather provide unbiased measurements of deuterium composition.

The aim of this study was to explore the effects of intrafeather hydrogen isotopic variability and sample processing on deuterium abundance measured in songbirds' feathers. The specific objectives were: (1) to study hydrogen isotopic gradients along the feather by comparing samples from the more distal and proximal parts of the same feather. I predicted there would be no deuterium differences, as songbird feathers grow quickly and individuals avoid large displacements during moulting. (2) To study the effect of feather ultrastructure on deuterium composition by comparing sections of rachis and vane from the same feather. In this case, I predicted subtle, but quantifiable deuterium differences between the sections. (3) To test the effect of two different protocols of feather processing by analysing ground and non-ground sections from the same feather. I expected an effect of grinding on deuterium measurements, since the exchangeable fraction of hydrogen is affected by particle size, as it has been demonstrated in hair.

Methods

The feathers (n=35) were collected from four species (Chaffinch, *Fringilla coelebs*, n=5; Grasshopper Warbler, *Locustella naevia*, n=4; Bluethroat, *Luscinia svecica*, n=4; Song Thrush, *Turdus philomelos*, n=22) trapped during the autumn migration of 2017 in the Manecorro ringing station located in the Doñana National Park (SW Spain; Arroyo 2004). The sixth secondary was plucked from the right wing of birds. Samples were stored in paper envelopes until their analysis.

Feather calamus was removed because this section of the shaft contains skin and blood residues, which may bias deuterium measurement of feather keratin (Hobson et al. 1999; Mazerolle and Hobson 2005; Wolf et al. 2012). Each feather was placed into a small tube and rinsed three times with absolute ethanol and a fourth time with distilled water. For each rinse, washing was enhanced by vortexing for 10 s. Wet samples were placed into a drying chamber at 50 °C during 48 h.

Each feather was divided into four subsamples (Fig. 1). First, the left section of the vane ('vane') was separated with surgical scissors and ground to a homogeneous powder. Therefore, 'vane' provided deuterium composition representative for the full feather's vane in powder form. From the remaining sample of the feather, the 'tip', 'bottom' and 'rachis' (Fig. 1) were cut with a scalpel. 'Tip' and 'bottom' aimed to represent the distal and proximal sections of the vane, respectively. Their comparison determined possible isotopic gradients along the feather (objective 1). Irrespective of the existence of such gradients, the 'vane' had to have an average composition of deuterium between 'tip' and 'bottom', otherwise any deviation would imply an effect of the processing protocol (objective 3). It is necessary to clarify that 'tip' included the upper section of the rachis. At the distal region of small feathers, these two structures showed similar thickness and consistency making difficult its separation. Finally, 'rachis' was the rachis section contiguous to 'bottom'. Here, rachis was thick and patently different from vane, even in the

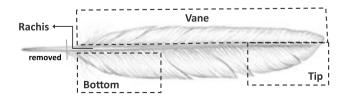


Fig. 1 The four subsamples obtained from each feather. See text for details on the procedures applied to each section



smallest feathers of the grasshopper warbler. The comparison between 'rachis' and 'bottom' tested the second objective.

An average of 0.303 mg (0.024 SD) of feather from each section was weighed with an electronic microbalance (Sartorius[®] Cubis Microbalance, ± 0.001 mg) and placed into 5.0×3.3 mm silver foil cups (Lüdiswiss, Switzerland) and folded. Encapsulated samples (n=140) were stored in 96-well plates during 1 month to ensure equilibrium of the exchangeable hydrogen fraction with ambient water vapour (Wassenaar and Hobson 2003). Samples were analysed in a random order.

Isotope measurements were performed on hydrogen derived from high-temperature flash pyrolysis at 1450 °C by means of Flash HT Plus elemental analyser coupled to a Delta-V Advantage isotope-ratio mass spectrometer via a CONFLO IV interface (Thermo Fisher Scientific, Bremen, Germany). Four keratin reference materials with a known deuterium isotopic composition of the non-exchangeable hydrogen component were used to calibrate results to the Vienna Standard Mean Ocean Water. The reference materials were: CBS (caribou hoof standard, -157.0%), KHS (kudu horn standard, -35.3%; both supplied by Environment Canada), USGS-42 (human hair from Tibet, -72.2\%; supplied by the Reston Stable Isotope Laboratory) and LIE-PA2 (an internal standard made from razorbill feather, +3.8%). Reference materials were encapsulated following the same protocol described previously and simultaneously to feather samples. Samples and reference materials were stored together for hydrogen equilibration (Wassenaar and Hobson 2003; Wassenaar 2019).

The isotopic composition is reported in the conventional delta (δ^2 H) per mil notation (‰). Replicate assays of laboratory standards routinely inserted within the sampling sequence indicated analytical measurement errors of $\pm 1.6\%$ (SD, n = 1023).

Feather processing and isotopic analyses were performed at the Stable Isotope Lab of the Estación Biológica de Doñana (Spain), which is certified to ISO9001:2015 and ISO14001:2015 quality and environmental management systems.

I used repeated measures ANOVA to test for differences between the four studied sections of the feather. The species was included as a categorical predictor. Greenhouse–Geisser $(\hat{\epsilon})$ and the Huynh–Feldt $(\tilde{\epsilon})$ estimations of sphericity of the variance–covariance matrix (Greenhouse and Geisser 1959; Huynh and Feldt 1976) were used to correct the degrees of freedom of the model. I applied Bonferroni post hoc tests to determine statistical differences between feather sections (Maxwell 1980). To obtain conversion formulae, I fitted linear regression models between pairs of feather sections that were statistically different in the previous post hoc test. All analyses were conducted with Statistica 7.0 (StatSoft, Inc).

Results

There were statistically significant differences among the $\delta^2 H$ in each section of the feather (Table 1). 'Rachis' had $\delta^2 H$ values around 11% lower than 'vane' and 'bottom', and around 8% lower than 'tip' (Fig. 2). Interestingly, 'tip' also differed significantly from 'vane' and 'bottom' (Bonferroni post hoc test p < 0.001) showing, in general, more negative $\delta^2 H$ values (Fig. 2). The $\delta^2 H$ values for all feather sections differed consistently in all species but the grasshopper warbler (Fig. 2; Table 1). Moreover, as 'vane' and 'bottom' had the same $\delta^2 H$, the hypothesis of deuterium gradients along the feather was rejected, as well as, any potential effects of vane grinding.

In spite of the differences, $\delta^2 H$ values between different sections of the feather were strongly correlated (Table 2). Thus, variability among individuals was captured in the same way by any feather section.

Discussion

The most important intra-feather variation in deuterium composition in passerine feathers was between rachis and vane, a difference already noticed in several non-passerine

Table 1 Results of the repeated measures ANOVA for the δ^2 H measured in each section of the feather

	SS	d.f.	MS	F	p	G–G adj. d.f.	G–G adj. p	H–F adj. d.f.	H.F. adj. <i>p</i>
Between species									
spp.	1406.5	3	468.8	0.388	0.763				
Error	37,499	31	1210						
Within species									
Feather section	1457.0	3	485.7	91.408	< 0.001	2.429	< 0.001	2.906	< 0.001
Feather section × spp.	91.8	9	10.2	1.920	0.058	7.288	0.075	8.718	0.061
Error	494.1	93	5.3			75.306		90.082	

G–G are the results obtained from the adjustment of the Greenhouse–Geisser $\hat{\epsilon}$ = 0.810, while H–F are those from the adjustment of Huynh–Feldt $\tilde{\epsilon}$. = 0.967



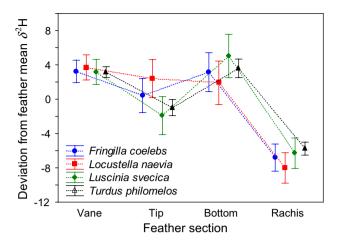


Fig. 2 δ^2 H for each section of the feather in each studied species. Due to the large differences in the δ^2 H among individuals, δ^2 H values have been rescaled using the mean value of each individual to improve visualization. Therefore, each point shows the average of individuals' deviations in δ^2 H. Bars denote the 95% confidence intervals

Table 2 Conversion equations to rescale $\delta^2 H$ values from different sections of a feather

Sections	r^2	p	Equation
Vane versus rachis	0.9807	< 0.001	Vane = 10.19687 + 1.01084 * rachis
Vane versus tip	0.9721	< 0.001	Vane = 0.23132 + 0.94339 * tip

species (Wassenaar and Hobson 2006; Wiley et al. 2010; Bontempo et al. 2014). Such difference between rachis and vane was expected because they have biochemical and structural differences (Harrap and Woods 1967; King and Murphy 1987), which may affect hydrogen isotope incorporation (Bortolotti 2010; Grecian et al. 2015) and exchange (Meier-Augenstein et al. 2013). The magnitude of the observed differences in δ^2 H (c. 11%) was similar to the 6.8% (2.6% SD) found previously for the Swainson's Thrush (*Catharus ustulatus*; Wassenaar and Hobson 2006). Although the number of analysed samples for the grasshopper warbler, the chaffinch and the bluethroat was small, this 11% difference between rachis and vane was surprisingly constant among individuals and species. Overall, this finding, as well as that by Wassenaar and Hobson (2006), suggests strong similarities among passerines in the biochemical processes controlling hydrogen isotope composition in feathers, regardless of their phenotypic characters, such as size, shape and colour (but see Michalik et al. 2010). This hypothesis concurs with the fact that differences in amino acid composition between rachis and vane are larger within than between species (Schroeder et al. 1955; Harrap and Woods 1967; King and Murphy 1987), suggesting a strong evolutionary conservation of feather keratins (Murphy et al. 1990). As amino acid composition of proteins affects the hydrogen exchangeable fraction (Meier-Augenstein et al. 2013), the slight differences in the amino acid composition between rachis and vane seem the most plausible underlying cause for their $\delta^2 H$ differences. Nevertheless, a more extensive sample of species would be necessary to confirm that hydrogen isotopic differences between rachis and vane are conserved across all passerine taxa. In the same way, a similar study for other elements used regularly in isotope research, such as C or N, would be advisable too (but see Bontempo et al. 2014; Grecian et al. 2015).

The observed δ^2 H differences between vane and rachis require special attention. In fact, both parts should be considered to some extent as different tissues, since they present a different factor of hydrogen isotope discrimination (Wolf et al. 2012; Grecian et al. 2015), making impossible a direct comparison of their results. For instance, if the tip of a small feather is clipped and it is necessary to use the whole piece in the isotope analysis, the δ^2 H will be around 3% lower than expected in a pure vane sample. In consequence, the measured δ^2 H will bias c. 350 km of latitude the assignment of the provenance of the bird. Therefore, it seems essential to state explicitly in the methodology of our studies, whether or not rachis was included. Surprisingly, this information is usually not reported, hindering the comparability and replicability of the studies (Meier-Augenstein et al. 2013). Since most ecologists rely on specialists to carry out their laboratory analyses, they can be underestimating or even ignoring these potential issues (Jardine and Cunjak 2005; Hawke et al. 2018).

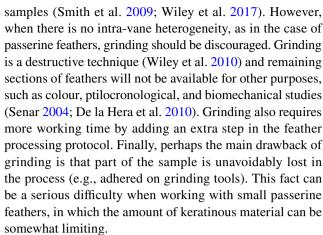
Due to the hydrogen isotopic differences between rachis and vane, some authors recommend to use only vane as a source of keratinous material for deuterium analyses (Wassenaar and Hobson 2006; Bontempo et al. 2014; but see Meehan et al. 2003). However, in some instances this is not possible. Samples may be limited to small feathers, such as contour feathers, since they might be the only ones that moulted during the life cycle period of our interest (Chamberlain et al. 1996; Kelly et al. 2002; Mazerolle and Hobson 2005; Hobson 2007; Langin et al. 2007; Rader et al. 2017) or the only ones allowed to be collected in museum specimens (Smith et al. 2009; Wiley et al. 2017). In other cases, because of ethical reasons feather samples can be limited to a clipped piece of few millimetres (Pérez and Hobson 2006; McDonald and Griffith 2011). Since passerine feathers are light structures, small samples, as those previously cited, can seriously restrict the possibilities to obtain deuterium measurements due to insufficient vane mass. Moreover, analyses for other isotopes or replicates would be unfeasible, limiting our ability to get the maximum information per sample. In these situations, a flexible protocol allowing including the rachis in the analysis can be helpful since the rachis



represents a relevant fraction of the total feather mass (King and Murphy 1987; Murphy et al. 1990). For instance, in this study, the feathers weighed between 3 and 5 mg and the rachis represented c. 50% of their weight. As demonstrated here, δ^2 H values from rachis were strongly correlated to those values from vane. This property grants the possibility to apply linear transformations to the results to get unbiased deuterium measurements in samples containing rachis.

Contrary to previous work (Wassenaar and Hobson 2006; Smith et al. 2008, 2009; Wiley et al. 2010; Bontempo et al. 2014; Grecian et al. 2015; Wassenaar 2019), feathers did not show isotopic gradients between bottom and tip. This opposite result can be due to the fact that I worked with passerines, while this type of intra-feather variation has been found in non-passerines. In species, such as raptors (Smith et al. 2008, 2009) or herons (Wassenaar and Hobson 2006), full replacement of a single flight feather takes between 7 and 14 weeks (Meehan et al. 2003; Zuberogoitia et al. 2018). During this period, individuals may move long distances (e.g. migration) and their feathers may incorporate along their length all this environmental variability. However, in passerines, single feathers are replaced in few days (Newton 1967; Bensch et al. 1991; De la Hera et al. 2011). Furthermore, passerines during moult do not move long distances because of the energy demands of the moulting process and their impaired flying abilities (Merilä 1997; Swaddle and Witter 1997). Thus, in passerines, a feather grows completely at the same place during a short time period. In consequence, feathers would use a homogeneous source of hydrogen, which would favour a homogeneous deuterium composition along their entire length. Therefore, by resampling different sections of the vane, e.g. to repeat analyses, δ^2 H measurements are not affected. This finding is also useful when the available section of the feather varies between individuals, as when the target feather is broken.

In opposition to mammal hair (Chesson et al. 2009; Qi and Coplen 2011), grinding did not have any effect on δ^2 H measures in feather samples. Probably, grinding does not have a critical effect on hydrogen exchange between feather materials and ambient humidity due to the natural structure of the vane. By grinding this complex structure of barbs and barbules up to small particles, one obtains a similar material than by cutting a small vane piece of a few millimetres. Therefore, in spite of the fact that we would be violating the principle of identical treatment (Wassenaar and Hobson 2003; Chesson et al. 2009; Wassenaar 2019), empirical evidence suggests that we may be somewhat flexible in the grinding step of the protocol. Grinding might be recommended when there is evidence of intra-feather heterogeneity (Wassenaar and Hobson 2006; Smith et al. 2008; Wiley et al. 2010). Under these circumstances, tissue homogenization can provide a more integrative value for the full feather enhancing any comparison between the



In conclusion, working with passerine feathers may be arduous because of the limited amount of sample that they provide. This study demonstrates that protocols can be flexible to some extent because: (1) any section of the vane can be used since there are no hydrogen isotopic differences along the feather; (2) the rachis can be included despite its lower δ^2 H, since conversion equations can be applied to get unbiased results; (3) grinding can be avoided, since particle size does not have any effects on the exchangeable hydrogen fraction in feather material. This flexibility in the processing protocol can help to maximize the information obtained from passerine feathers.

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