



# Cetacean Skeletons Demonstrate Ecologically Relevant Variation in Intraskkeletal Stable Isotopic Values

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Conservation science requires quickly acquiring information and taking action in order to protect species at risk of extinction. Stable isotope measurements are one way to rapidly gather data regarding species' foraging ecology and habitat use, and passively collected samples limit additional stress to at-risk species. For these samples to be useful, however, we must know how representative they are of the stable isotope ratios of the entire organism. Bone tissue, often stored in museum collections or research centers, may be the most readily available tissue from rare, endangered, or extinct vertebrates, but using bone requires practitioners to understand intraskkeletal stable isotope variation. We sampled the same eight skeletal elements from 72 cetacean skeletons from 14 species to evaluate intraskkeletal variation in carbon and nitrogen isotope values. We found considerably more variation than anticipated. Carbon intraskkeletal ranges varied from 0.4 to 7.6‰, with 84.7% ( $n = 61$ ) of skeletons having a range  $> 1‰$ , and 55.5% ( $n = 40$ ) exhibiting a range  $> 2‰$ . Similarly, nitrogen intraskkeletal ranges varied from 0.4 to 5.2‰, with 59.7% ( $n = 43$ ) of skeletons exhibiting a range  $> 1‰$ , and 15.3% ( $n = 11$ ) with a range  $> 2‰$ . There were differences in which bones contributed most to intraskkeletal variation; however, we advise against using humeri and mandibles as these bones presented the most consistent trends in deviation from the intraskkeletal means for both isotopes. The large intraskkeletal variation we observed is likely due to changes in foraging behavior or habitat use being reflected differently in bone isotope ratios due to differences in bone turnover rates. We suggest that for cetaceans, intraskkeletal carbon isotope ranges  $> 1‰$  and nitrogen ranges  $> 2‰$  are ecologically relevant, and that using different bones from animals of the same population may produce false positive differences in foraging behavior or habitat within the population if intraskkeletal variation is not considered. Future studies should use the same bones from each animal and conduct species-specific analyses of intraskkeletal variation, if possible, when using specimens of opportunity. Failure to consider this variation could lead to erroneous conclusions regarding a species range or key habitats, jeopardizing conservation efforts.

**Keywords:** bone, carbon, cetacean, conservation, intraskkeletal variation, museum collections, nitrogen, stable isotopes

## INTRODUCTION

Conservation science is a crisis discipline because conservation action typically must be taken for species at risk of extinction before practitioners are confident in the sufficiency of their data (Soulé, 1985). One fundamental difficulty in wildlife conservation is rapidly understanding how species interact with, and utilize, their habitat (Aberg et al., 2000; Cristescu and Boyce, 2013). A variety of tools and research methods have been developed and employed to gain insight into habitat use. Typically, these methods require directly interacting with the animal in some manner, such as radio telemetry, capture or sedation to collect biological samples, or long-term observation; all of which can alter animal behavior (Brigham, 1989; Pietz et al., 1993; Guthery and Lusk, 2004; Brooks et al., 2008; Rachlow et al., 2014). Less invasive methods, such as camera traps and drones, still may alter animal behavior, as many animals identify the device in their habitat and interact with it (Meek et al., 2014, 2016; Mulero-Pazmany et al., 2017). Methods with no animal interactions, such as shore-based marine mammal observational studies, require thousands of observer hours, can be implemented over only limited spatial areas, are restricted in insight regarding surface-based activities, and are susceptible to observer error (Rugh et al., 1990; Aragones et al., 1997).

Stable isotope analysis (SIA) is an innovative technique for investigating wildlife habitat use, such as providing insight into foraging behavior, niche segregation, individual-level resource utilization, and diet shifts (Hobson, 1999; West et al., 2006; Newsome et al., 2007). Studies incorporating SIA can employ a variety of tissue types, each providing unique temporal snapshots of ecological or dietary conditions reflecting the timeframe when the tissue was generated. Although many SIA studies use actively collected samples that require direct human – animal interaction, such as biopsy plugs or blood samples, one of the most powerful aspects of this technique is the ability to gain insight from passively collected samples, such as molted feathers or fur (McKechnie, 2004; Thompson et al., 2005). Feathers, for example, often incorporate the isotopic value of the water and food resources in the region where they are grown, and are excellent samples for identifying migratory or breeding grounds without requiring human – bird interaction (Chamberlain et al., 1997; Hobson et al., 2001; Guillemain et al., 2019). Thus, SIA of specimens of opportunity provides a powerful monitoring system that minimizes invasive activities, limits impacts on animal behavior, and can be rapidly completed. Opportunistically collected samples provide a valuable alternative to capturing or harassing wildlife, but present new challenges. For rare or difficult to locate animals, small sample size or less than ideal samples can complicate analyses (Ben-David and Flaherty, 2012; Hopkins and Ferguson, 2012). If researchers are attempting to gain insight into longer-term behavior from passively collected samples, such as feathers, bone remains from archeological sites, or more recent skeletal remains, they must determine if the sample evaluated accurately reflects the tissue's value for the animal as a whole.

In rare, endangered, or extinct vertebrate species, bone is often the only tissue available for SIA, and is routinely stored in museum and research collections. Despite the ubiquitous

availability of bone tissue from vertebrates, comparatively few studies have focused on bone SIA (Vander Zanden et al., 2015), and only a small subset of these studies has examined isotopic variation among different bones from the same organism (Table 1). Bone tissue is slow to grow and regenerate, thus incorporating and reflecting diet isotopic signatures at a slower rate than other tissues (Newsome et al., 2007; Vander Zanden et al., 2015). Controlled feeding studies have been completed that examine bone isotopic values, many with the purpose of coupling isotopic values in bone with soft tissues (Ben-David et al., 1997; Hong et al., 2000; Phillips and Eldridge, 2006). However, these studies only examined matching bones, or the same bones in each study organism. Bone tissue is replaced and repaired at different rates depending on the bone's function, density, and size (Kohn and Cerling, 2002; Lafage-Proust et al., 2015). Due to these differing turnover rates in bone, different bones sampled from the same animal may have different isotopic values. In the case of rare or extinct species, it may be impossible to acquire complete skeletons, compelling researchers to compare isotopic values from different bones among conspecifics. These non-matching bones may suggest different diets, water sources, or other environmental parameters, not because individuals in a population were utilizing different resources, but because the bones reflect dietary or habitat shifts at different rates. Failure to consider isotopic variation among different bones of conspecific individuals may result in erroneous conclusions regarding environmental conditions and dietary habits.

In this study, we investigated intraskelatal stable isotope variation of  $\delta^{13}\text{C}$  and  $\delta^{15}\text{N}$  in 14 cetacean species using skeletons from the National Museums Scotland osteological collection. Cetaceans are a quintessential example of the challenges conservation scientists face: studying cetaceans *in situ* is often invasive, requiring locating the animal, following it, and interacting with it in some way (Dawson et al., 2008; Ballance, 2009). These processes are time consuming and have numerous logistical challenges beyond required permitting. Cetacean skeletons, however, have been collected and housed in museum collections for hundreds of years, providing a large specimen-of-opportunity cache for researchers. By sampling multiple bones from the same skeleton, we can establish an understanding regarding how representative a given bone is of the entire skeleton, thus increasing the power of skeleton-based SIA studies, and providing a valuable contribution to passive habitat use studies.

## MATERIALS AND METHODS

To test for intraskelatal isotopic variation, we sampled the same 8 bone locations from 72 cetacean specimens (14 species) housed in the National Museums Scotland osteological collection (Table 2 and Supplementary Table S1). In order to consistently sample the same location for each bone among individuals and species, we compared bone size and selected the same proportional sampling site. We selected these specimens because they were complete or near-complete skeletons, were well represented in the collection so small-scale destructive sampling

**TABLE 1** | Summary of data from previous studies that investigated intraskkeletal  $\delta^{13}\text{C}$  and  $\delta^{15}\text{N}$  isotopic variation relevant to our research.

References	Species	Sample locations	Number of individuals	Avg. Intraskkeletal Range (‰)	
				$\delta^{13}\text{C}$	$\delta^{15}\text{N}$
Deniro and Schoeniger, 1983	Mink (unknown species)	Femur and humerus	15	0.6	0.7
Deniro and Schoeniger, 1983	Rabbit (unknown species)	Femur, humerus, mandible, radius, scapula, distal tibia, ulna	3	0.3	0.3
Jorkov et al., 2009	<i>Homo sapiens</i>	Rib, femur, temporal	57	0.4	0.8
Jorkov et al., 2009	<i>Homo sapiens</i>	Rib, femur, temporal, molar	16	0.8	1.0
Riofrío-Lazo and Aurióles-Gamboa, 2013	<i>Mirounga angustirostris</i>	Mandible, tooth, maxilla	14	1.8	1.1
Riofrío-Lazo and Aurióles-Gamboa, 2013	<i>Mirounga angustirostris</i>	Mandible and maxilla	17	0.8	0.5
Olsen et al., 2014	<i>Homo sapiens</i>	Rib, fibula, metacarpal	6	0.6	1.6
Webb et al., 2016	<i>Sus domesticus</i>	Rib and femur	48	NA	0.3
Cheung et al., 2017	<i>Homo sapiens</i>	Femur and fibula	11	1.0	0.6
Cheung et al., 2017	<i>Homo sapiens</i>	Femur and radius	6	1.2	0.5
Cheung et al., 2017	<i>Homo sapiens</i>	Femur and tibia	1	1.1	1.2
Cheung et al., 2017	<i>Homo sapiens</i>	Femur and ulna	1	1.8	0.3
Clark et al., 2017	<i>Odobenus rosmarus divergens</i>	Cranium and mandible	11	0.1	0.3
Clark et al., 2017	<i>Pusa hispida</i>	Calcaneus, mandible, femur, humerus, innominate, phalanx, rib, scapula, metatarsal, vertebra	1	0.9	1.2
Clark et al., 2017	<i>Phoca sp.</i>	Cranium, femur, humerus, innominate, phalanx, rib, scapula, metatarsal, vertebra	1	0.5	1.1
Clark et al., 2017	<i>Enhydra lutris</i>	Mandible, femur, humerus, innominate, rib, scapula, metatarsal, vertebra	1	1.2	0.7
Fahy et al., 2017	<i>Homo sapiens</i>	Femur, tibia, rib, radius, occipital, metacarpal, humerus, thoracic vertebrae, pelvis, clavicle	10	0.9	1.6
Bas et al., 2019	<i>Otaria byronia</i>	Atlas, humerus, basioccipital	14	0.8	1.3
Bas et al., 2019	<i>Lagenorhynchus obscurus</i>	Atlas, humerus, basioccipital	15	1.4	0.3

**TABLE 2** | Number of individual animals sampled per species for each skeletal sampling location.

Species	Skeletal sample location							
	Occipital condyle	Mandibular ramus	Thoracic vertebral body	Thoracic vertebral spinous process	Proximal rib	Distal rib	Scapula	Humeral head
<i>Balaenoptera acutorostrata</i>	3	3	3	3	3	3	2*	3
<i>Delphinus delphis</i>	5	5	5	5	5	5	5	5
<i>Globicephala melas</i>	4	4	4	4	4	4	4	4
<i>Grampus griseus</i>	5	5	5	5	5	5	5	5
<i>Hyperoodon ampullatus</i>	5	5	5	5	5	5	5	5
<i>Kogia breviceps</i>	3	3	3	3	3	3	3	3
<i>Lagenorhynchus acutus</i>	5	5	5	5	5	5	5	5
<i>Lagenorhynchus albirostris</i>	5	5	5	5	5	5	5	5
<i>Mesoplodon bidens</i>	10	10	10	10	10	10	10	10
<i>Orcinus orca</i>	4	4	4	4	4	4	4	4
<i>Phocoena phocoena</i>	5	5	5	5	5	5	5	5
<i>Stenella coeruleoalba</i>	5	5	5	5	5	5	5	5
<i>Tursiops truncatus</i>	10	9*	10	10	10	10	10	10
<i>Ziphius cavirostris</i>	3	3	3	3	3	3	3	3

We sampled the same locations by bone among individuals to reduce the effects of natural bone variability, and samples contained a mixture of cortical and trabecular bone. \*One *Balaenoptera acutorostrata* skeleton was without scapulae and one *Tursiops truncatus* skeleton was without mandibles.

would not hinder future studies and encompassed the breadth of physiological and ecological variation in cetaceans. We aimed to limit our sampling to adult ( $n = 49$ ) or subadult

( $n = 11$ ) specimens, but due to the limited number of specimens that contained all sampling locations we included 12 juvenile specimens in order to increase sample size.

We used a battery powered handheld drill to remove 1 g of bone tissue and subsampled 200 mg for collagen extraction. Our extraction protocol was adapted from Ambrose (1990) and Jorkov et al. (2007). We ground subsamples with mortar and pestle and performed lipid extractions in a 2:1 chloroform:methanol solution three times for 30 min each; if the supernatant was not clear after three washes, additional washes were carried out as needed. The mineral component was removed using a 30 min 0.5M HCl bath followed by 3 deionized water rinses, and a 30 min 0.1M NaOH bath followed by 3 deionized water rinses. Previous studies demonstrated that bone tissue lipid extraction and acidification demineralization does not significantly alter  $\delta^{15}\text{N}$  values (Tomaszewicz et al., 2015; Tatsch et al., 2016). We added 7 ml of pH 3 water to each sample and incubated at 80°C for 24 h. The supernatant was collected and freeze-dried, resulting in purified collagen. Between 0.85 and 1.15 mg of collagen was loaded in  $3 \times 5$  mm tin capsules and submitted for C and N stable isotope analysis.

Stable isotope analysis was completed at the Cornell Isotope Laboratory at Cornell University using a Thermo Delta V isotope mass spectrometer interfaced with a NC2500 elemental analyzer (ThermoFisher Scientific, Waltham, MA United States 02451). We calibrated our results using 2 primary reference scales: Vienna Pee Dee Belemnite for  $\delta^{13}\text{C}$ , and Atmospheric Air for  $\delta^{15}\text{N}$ . To ensure accuracy and precision, we analyzed an in-house standard ( $\delta^{13}\text{C}$ :  $-20.16 \pm 0.03\text{‰}$  and  $\delta^{15}\text{N}$ :  $6.35 \pm 0.05\text{‰}$ ) between every 10 samples. As an additional measure of extraction method and analysis accuracy and repeatability, we randomly selected 2 bones, subsampled 4 additional 200 mg samples each, and followed the methods described above to extract collagen and analyze for stable isotope ratios ( $\delta^{13}\text{C}$ :  $-14.93 \pm 0.02\text{‰}$  and  $\delta^{15}\text{N}$ :  $10.98 \pm 0.08\text{‰}$ ;  $\delta^{13}\text{C}$ :  $-13.79 \pm 0.05\text{‰}$  and  $\delta^{15}\text{N}$ :  $11.55 \pm 0.06\text{‰}$ ). We also evaluated collagen sample composition (percent carbon, percent nitrogen, and C/N ratio) and collagen percent yield to monitor sample quality.

We employed descriptive statistics to explore intraskkeletal variation among bone sampling locations for both  $\delta^{13}\text{C}$  and  $\delta^{15}\text{N}$ . Because we are not making comparisons among animals, we did not have to consider the Suess effect, which is long-term incorporation of isotopically light carbon into the marine ecosystem due to fossil fuel use (Keeling, 1979). Analyses were performed using R (R Core Team, 2018) with RStudio (RStudio Team, 2016).

## RESULTS

We found a high degree of variation in the isotopic values among different bones taken from the same animal. For example, internal skeletal ranges for  $\delta^{13}\text{C}$  varied from 0.4 to 7.6‰, with 84.7% ( $n = 61$ ) of skeletons having a range  $>1\text{‰}$ , and 55.5% ( $n = 40$ ) exhibiting a range  $>2\text{‰}$  (Figure 1). Similarly, skeletal ranges for  $\delta^{15}\text{N}$  varied from 0.4 to 5.2‰, with 59.7% ( $n = 43$ ) of skeletons exhibiting a range  $>1\text{‰}$ , and 15.3% ( $n = 11$ ) with a range  $>2\text{‰}$ . For all skeletons, and for both isotopes, at least one bone was  $\geq 1$  SD from the skeletal mean, and in most skeletons multiple bones were  $\geq 1$  SD from the mean. For  $\delta^{13}\text{C}$ , the number

of skeletons with 1, 2, 3, and 4 bones  $\geq 1$  SD from the mean was 23, 31, 14, and 4, respectively. For  $\delta^{15}\text{N}$ , the number of skeletons with 1, 2, 3, and 4 bones  $\geq 1$  SD from the mean was 10, 41, 17, and 4, respectively. In a subset of skeletons ( $n = 31$  for  $\delta^{13}\text{C}$ ;  $n = 17$  for  $\delta^{15}\text{N}$ ), 1 bone was  $\geq 2$  SD from the skeletal mean and one bottlenose dolphin (*Tursiops truncatus*) skeleton had 4 bones  $\geq 2$  SD from the mean for  $\delta^{15}\text{N}$ .

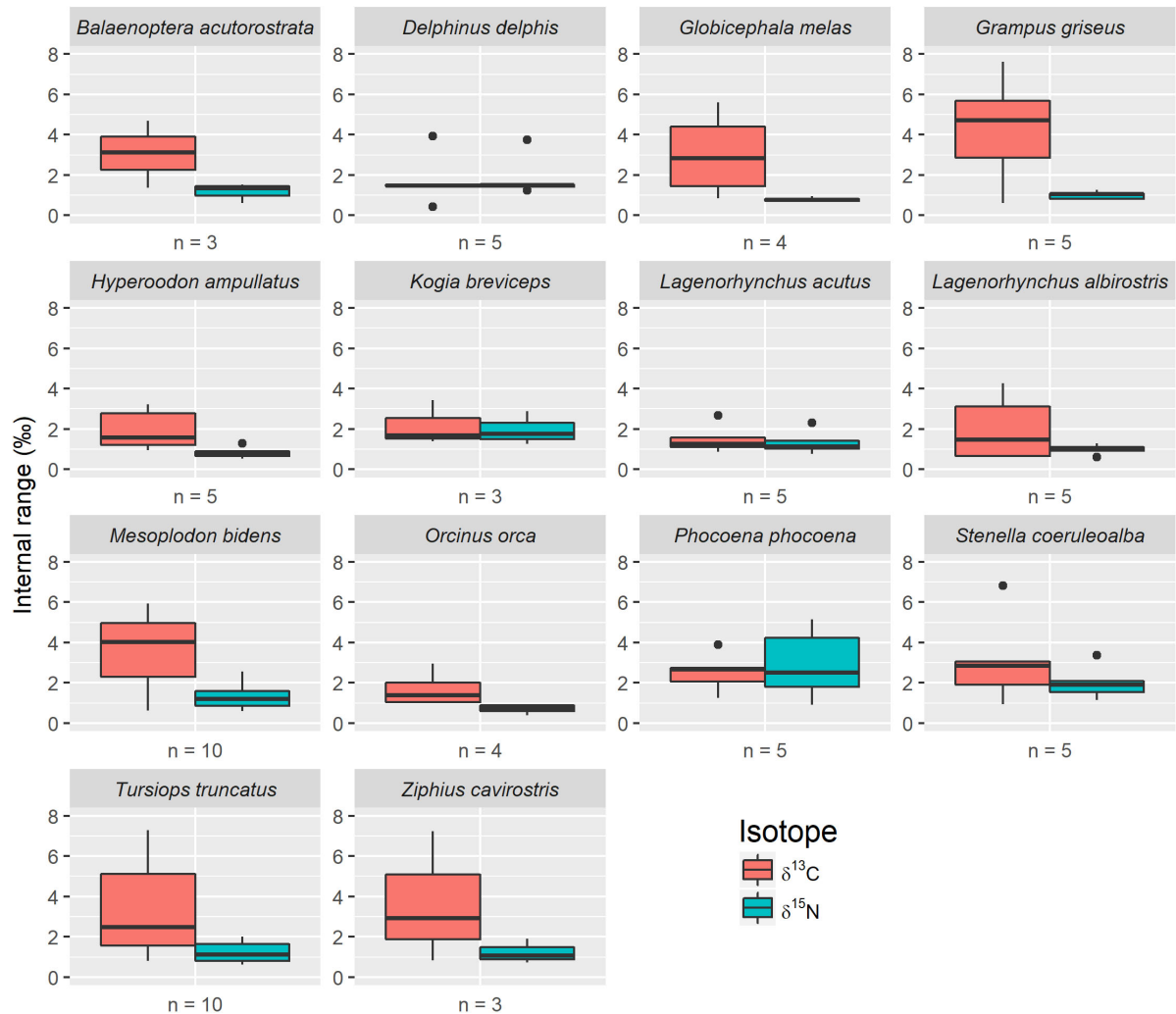
There were no consistent trends regarding which specific bones within an individual animal differed in isotopic values from the skeletal mean across specimens. However, the proximal rib sampling location demonstrated the lowest levels of deviation, with 6 specimens (8.3%)  $\geq 1$  SD from intraskkeletal mean  $\delta^{13}\text{C}$  values, and 7 specimens (9.7%)  $\geq 1$  SD from  $\delta^{15}\text{N}$  intraskkeletal mean values. For  $\delta^{13}\text{C}$ , 50.0% ( $n = 36$ ) of humeral heads were  $\geq 1$  SD lower than the skeletal mean (no humeral heads were  $+1$  SD). Mandibular rami (40.3%;  $n = 29$ ) and scapulae (16.7%;  $n = 12$ ) had the second and third highest rates of deviation from mean skeletal  $\delta^{13}\text{C}$ , and all bone sample locations had at least one representative with  $\geq 1$  SD. We found 56.9% ( $n = 41$ ) of mandibular rami were  $\geq 1$  SD from the skeletal  $\delta^{15}\text{N}$  mean; specifically, of these 22.0% ( $n = 9$ ) were greater than the mean, whereas 78.0% ( $n = 32$ ) were less than the mean. Humeral heads (30.0%;  $n = 28$ ) and occipitals (25.0%;  $n = 18$ ) had the second and third highest rates of deviation from the skeletal  $\delta^{15}\text{N}$  average, and all bones had at least one representative  $\geq 1$  SD from the mean.

Mean collagen yield was 10.6%, with a range of 1.8–37.5%. This includes 37 samples with artificially low percent yield due to a freeze dryer malfunction resulting in partial loss of the sample or producing collagen that was challenging to recover from the vial, which prevented obtaining an accurate weight. Mean %C and %N were 29.56 and 10.19, respectively, with a mean C/N ratio of 3.49.

## DISCUSSION

Intraskkeletal isotopic variation has not been well investigated across a variety of taxa, and only one other study has considered this topic for cetacean skeletons (Vander Zanden et al., 2015; Bas et al., 2019). Studies of this type are challenging due to difficulty in locating large numbers of intact skeletons of the same species. As a result, previous studies typically had small sample sizes, low numbers of sampling locations, or a combination of both (Table 1). We addressed this deficiency by combining a large sample size from five cetacean families with many bone sampling locations per skeleton (Table 2). We documented much greater intraskkeletal isotopic variation than has previously been reported (Figure 1 and Table 1), suggesting that if analyses were to be expanded to other taxa, similar results may be observed.

We identified some noteworthy patterns in stable isotope values that can be used to better inform the design of intraskkeletal isotope studies in cetaceans. The proximal rib demonstrated the lowest rate of deviation from both  $\delta^{13}\text{C}$  and  $\delta^{15}\text{N}$  intraskkeletal means and may be the best bone from our sampling locations to use for comparative studies. The  $\delta^{13}\text{C}$  isotopic value of half of all humeral heads was  $\geq 1$  SD from the skeletal mean, so we



**FIGURE 1 |** Individual intraskkeletal range for  $\delta^{13}\text{C}$  and  $\delta^{15}\text{N}$ , grouped by species ( $n$  = number of individual animals evaluated). Boxes present median and interquartile range and whiskers represent 95% confidence intervals.

advise against using this bone as representative of the skeleton as a whole. The mandibular ramus probably also should be avoided as we documented large, but inconsistent deviation from the skeletal mean for both isotopes. These two sampling locations represent two different bone types with different turnover rates. The humeral head is part of the humeral long bone and forms the shoulder joint with the scapula. However, cetacean skeletal and muscle anatomy studies have found that the humeral head is largely vestigial, and flipper movement is limited and related to maintaining balance and aiding in swimming speed (Cooper et al., 2007; Sanchez and Berta, 2010). As a result, this bone is under less ecophysiological pressure than other more mobile bones and joints. In contrast, the mandibular ramus is part of the dense mandibular irregular bone with a high degree of turnover and remodeling (Matsuura et al., 2014; Shadwick et al., 2017). In cetaceans, the mandible serves as the primary method of interacting with each other and the environment and is more susceptible to damage than other bones. These

two bones represent distinct functions and turnover rates, and this may explain why they exhibit the greatest difference from the skeletal mean.

For 13 of 14 species in our study,  $\delta^{13}\text{C}$  was more variable than  $\delta^{15}\text{N}$  (Figure 1). This trend is similar to cetacean intraskkeletal isotopic variation reported by Bas et al. (2019), who compared  $\delta^{13}\text{C}$  and  $\delta^{15}\text{N}$  isotope values among three sampling locations from 15 specimens (Table 1). They reported  $\delta^{13}\text{C}$  intraskkeletal isotopic variation that fell within the lower range of variation in our study, and we suspect that had additional skeletal elements been compared, then variation found by our two studies may have been similar. Greater  $\delta^{13}\text{C}$  than  $\delta^{15}\text{N}$  intraskkeletal variation is also consistent with Riofrío-Lazo and Auriolles-Gamboa (2013), who found variation in northern elephant seal skeletons (*Mirounga angustirostris*) and for sea otter (*Enhydra lutris*) skeletons in Clark et al. (2017). Many human (*Homo sapiens*) archeological studies also reported this trend (Table 1), but these authors typically compared only two or three sampling locations. The study most



similar in design to ours is Fahy et al. (2017); they compared  $\delta^{13}\text{C}$  and  $\delta^{15}\text{N}$  isotope values between 10 sample locations in 10 humans. They found  $\delta^{15}\text{N}$  intraskkeletal variability was greater than  $\delta^{13}\text{C}$  variation; however,  $\delta^{15}\text{N}$  variation was similar to values in our study. Only a few other studies examined intraskkeletal variation in terrestrial vertebrates (Table 1). The differences observed between terrestrial and marine mammal studies may be due, in part, to different physiological pressures placed on bones in terrestrial versus aquatic and semiaquatic environments.

Newsome et al. (2010) documented that younger marine mammals exhibit higher bone turnover rates of carbon and nitrogen stable isotopes, possibly contributing to intraskkeletal variation. We did not observe this pattern. In fact, adult animals demonstrated some of the highest levels of intraskkeletal variation. For example, harbor porpoises (*Phocoena phocoena*) had relatively low levels of  $\delta^{13}\text{C}$  intraskkeletal variation despite including one subadult individual, and the outlier animal was an adult (Figure 1). Atlantic white sided dolphins (*Lagenorhynchus acutus*) displayed generally lower levels of  $\delta^{13}\text{C}$  intraskkeletal variation compared to white beaked dolphins (*Lagenorhynchus albirostris*), even though we sampled five adult specimens for each species. Amongst beaked whales, Sowerby's beaked whales (*Mesoplodon bidens*) displayed the greatest median  $\delta^{13}\text{C}$  intraskkeletal variation despite including only adult animals, while both northern bottlenose whale (*Hyperoodon ampullatus*) and Cuvier's beaked whale (*Ziphius cavirostris*) samples included juvenile animals. No data on cetacean bone tissue turnover rates is available, but Newsome et al. (2006) estimated complete bone collagen turnover in yearling seals and sea lions at 8–10 months. If young cetaceans exhibit a similar pattern, then the moderate variation we observed in younger animals is logical because their bones are reflecting a shorter time span, and therefore less environmental variability than seen in older age classes. Thus, age class of the specimen does not seem to drive the variation we observed. Likewise, collection date and storage time of the specimens did not contribute to intraskkeletal variation. All our specimens, with the exception of two, were collected since 1989, and the two older specimens demonstrated similar intraskkeletal variation as modern specimens.

We were consistent in our sampling locations in each skeleton to reduce the introduction of additional variation due to natural differences throughout the bone. Each sample contained a mixture of mineralized cortical bone and spongy trabecular bone, and the inherent unequal ratios of these bone types at different sampling sites, and the differences in their turnover rates, may contribute to some of our observed intraskkeletal variation (Manolagas, 2000; Clarke, 2008). However, if this was a major contributing factor, we would expect to see animals of the same species demonstrating similar trends in variation; instead, we saw considerable variation at the individual animal level. This suggests a combination of physiological and ecological factors driving isotopic variation. Carbon and nitrogen isotope ratios in a skeleton reflect habitat and diet, respectively (Ben-David and Flaherty, 2012). Organisms in controlled settings, such as in feeding studies or laboratories, show little isotopic variation when fed a consistent diet, even when considering bone turnover rates (Deniro and Schoeniger, 1983). Therefore,

differences we observed in intraskkeletal isotope ratios suggest differences in foraging behavior and individual-level resource utilization over time.

As animals switch habitats or consume different food sources, the rate isotopes from these sources are incorporated will vary among bones due to bone-specific turnover rates (Newsome et al., 2010). This combination of changing environmental isotope ratios and physiological mechanisms leads to ecologically relevant intraskkeletal isotopic variation – that is, the isotopic values from different bones from the same individual could lead to different conclusions regarding an animal's life history if considered independently. This is especially important for studies forced to use non-matching bones for analyses. The amount of intraskkeletal range that is ecologically relevant depends on the specific questions being asked, but we suggest that  $\delta^{13}\text{C}$  ranges  $>1\text{‰}$  and  $\delta^{15}\text{N}$  ranges  $>2\text{‰}$  are ecologically significant for cetacean studies. Isoscape models built using specific prey resources of the species in this study do not yet exist, but we still can characterize how variation might affect researcher's conclusions by considering isoscapes already available. For example, an isoscape model built from jellyfish collected in waters around the British Isles demonstrates a 1–2‰ difference in  $\delta^{13}\text{C}$  values across the study area (Glew et al., 2019). Based on this, the  $\delta^{13}\text{C}$  variation we observed within the skeletons in our study would indicate different foraging locations along the United Kingdom shelf sea if the bone sample locations were considered independently. If a study is trying to identify important foraging or breeding habitats to make conservation recommendations and must make use of non-matching bones, a 1‰ difference may appear to suggest different regions of importance yet may simply represent differences among bones sampled from the same skeleton. Similarly, nitrogen isotope values in animals are enriched at rate of 3–4‰ for each increase in trophic level (Post, 2002), yet we observed  $\delta^{15}\text{N}$  intraskkeletal range values up to 5.16‰.

Bone turnover rates and changes in habitat use or foraging behavior could explain much of the intraskkeletal variation in carbon and nitrogen isotope ratios we observed, but there is still considerable unexplained variation. This could be due to physiological factors that are beyond the scope of our study, such as metabolic rates or bone disease/injury, both of which can alter bone growth patterns (Manolagas, 2000; Clarke, 2008; Olsen et al., 2014). We did not sample animals that had obvious signs of bone disease or injury remodeling, but there is little information regarding the individual life history of most specimens in our study. Thus, we do not know their movement patterns and habitat use, beyond general species information, or the specifics of their age or health. However, even amongst closely related species, such as Atlantic white sided dolphins and white beaked dolphins, which have overlapping habitats and grow to a similar size, we saw considerable differences in intraskkeletal variation (Weinrich et al., 2001; Galatius and Kinze, 2016). For some species, such as bottlenose dolphins, specimens in our study may have come from different populations, with different foraging behavior and habitat use, further contributing to intraskkeletal isotopic variation. Although we sampled a large breadth of cetacean species, we were limited to relatively small species

sample sizes due to difficulty in acquiring complete skeletons. In two cases, we chose to include animals that were missing one of eight sampling locations to increase sample size for that species; if this study was repeated with much larger species sample sizes, further trends in variation may become apparent. Regardless, to truly understand factors driving individual and species variation, we would need data from hundreds of tagged animals of the same species – where all their life movement data is available – to more systematically evaluate stable isotope variation for the species. Because this is not feasible for most cetacean studies, we instead must acknowledge that considerable variation exists within individual animals.

Specimens of opportunity are a critical resource for ecological studies, but they do present unique challenges that must be considered. Because opportunistically collected skeletons are often incomplete, necessitating comparisons between unmatched bones among animals, there is a need to understand intraskelatal isotopic variation. Our study demonstrates that substantial intraskelatal variation is present for the cetacean species we evaluated. Thence, we recommend that future studies using opportunistic bone tissue for stable isotope analysis conduct species-specific evaluations for intraskelatal variation. Failure to identify or consider this variation could have serious implications for studies that use bone isotope values to explore animal ecology. When the results of such studies are used to inform conservation action, it is imperative to consider that different bones from the same animals may suggest different habitats or resource use when none existed. Accounting for this intraskelatal variation in stable isotopes values produces more robust analyses and thus better-informed conservation management plans.

## DATA AVAILABILITY STATEMENT

The datasets generated for this study are available on request to the corresponding author.

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## AUTHOR CONTRIBUTIONS

KS conceived the idea. KS, JS, and MP designed the methodology. KS and ZT coordinated the data collection. KS and MP analyzed the data. KS led the writing of the manuscript. All authors contributed critically to the drafts and gave final approval for publication.

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## SUPPLEMENTARY MATERIAL

The Supplementary Material for this article can be found online at: <https://www.frontiersin.org/articles/10.3389/fmars.2020.00388/full#supplementary-material>

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