1	Uncovering the mosaic evolution of carnivoran skeletal systems
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10	Abstract
11	The diversity of vertebrate skeletons is often attributed to adaptations to distinct ecological
12	factors such as diet, locomotion, and sensory environment. Although the adaptive evolution of
13	cranial, appendicular, and vertebral skeletal systems is well studied in vertebrates,
14	comprehensive investigations of all skeletal components simultaneously are rarely performed.
15	Consequently, we know little of how modes of evolution differ among skeletal components.
16	Here, we tested if ecological and phylogenetic effects led to distinct modes of evolution among
17	the cranial, appendicular, and vertebral regions in extant carnivoran skeletons. Using multivariate
18	evolutionary models, we found mosaic evolution in which only the mandible, hindlimb, and
19	posterior region of the vertebral column showed evidence of adaptation towards ecological
20	regimes whereas the remaining skeletal components reflect clade-specific evolutionary shifts.
21	We hypothesize that the decoupled evolution of individual skeletal components may have led to
22	the origination of distinct adaptive zones and morphologies among extant carnivoran families
23	that reflect phylogenetic hierarchies. Overall, our work highlights the importance of examining

24	multiple skeletal components simultaneously in ecomorphological analyses. Ongoing work
25	integrating the fossil and paleoenvironmental record will further clarify deep-time drivers that
26	govern carnivoran diversity we see today and reveal the complexity of evolutionary processes in
27	multicomponent systems.
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29	Key words: adaptive landscape; Carnivora; ecomorphology; macroevolution; Ornstein-
30	Uhlenbeck modeling; phylogenetic comparative methods
31	
32	Introduction
33	The diversity of animal forms is one of the most salient patterns across the tree of life. In
34	mammals, morphological innovations in the cranial, appendicular, and axial skeletal systems
35	facilitate the incredible diversity found today, ranging from bats with winged forelimbs to the
36	biggest animals to have ever lived on earth. Many researchers have examined how variation in
37	the skull [1–6], limbs [7–10], or vertebrae [11–15] serve as crucial adaptations to their evolution.
38	These skeletal systems are traditionally examined independently and are rarely investigated
39	simultaneously even though these anatomical regions comprise a single, functionally integrated
40	system to support movement, sensation, and other life functions. When considered wholistically,
41	the observed variation across the different components of organismal anatomy is generally
42	explained by multitudinous factors, some that are potentially incongruous [16–18]. While this
43	evolutionary push-and-pull between anatomical regions may characterize the process of
44	evolution, the hypothesis can only be tested when the different skeletal components are explored
45	simultaneously rather than piecemeal. Simultaneous investigation of integrated components is
46	critical to our understanding of the role of developmental and/or functional integration in

47 canalizing macroevolutionary trajectories [19–21]. Here, we use carnivorans to investigate how 48 ecological and phylogenetic factors correspond to evolutionary changes in the cranial, 49 appendicular, and axial skeletal systems. Carnivorans (bears, cats, dogs, seals, and their relatives) 50 are a productive model system to examine skeletal evolution because of their high species 51 richness and vast distribution across most biomes in all continents and oceans, along with broad 52 ecological diversity in locomotor traits and feeding adaptations. 53 Components of carnivoran skeletal systems are well studied individually. In the skull, 54 craniomandibular diversity is influenced by several ecological factors and phylogeny [22–26]. 55 The skull exhibits decoupled evolutionary modes: cranial shape follows clade-specific

56 evolutionary shifts, whereas mandibular shape evolution is linked to broad dietary regimes

57 [6,27]. In the appendicular skeleton, ecomorphological divergence exists between the hindlimbs,

58 which are adapted primarily for locomotion, and the forelimb, which are adapted for multiple

59 functions ranging from running to grappling prey to manipulating objects [28–31]. Additionally,

60 more recent work using phylogenetic comparative methods found that scaling and phylogeny

61 exhibit stronger effects on limb evolution than do ecological parameters [32–34]. In contrast to

craniomandibular and appendicular ecomorphology, research on the axial skeleton is in nascent

stages. Initial research indicates that distinct regions of the vertebral column are under different

64 evolutionary pressures. The anterior region exhibits low disparity due to phylogenetic constraints

or ecological conservatism, whereas the posterior region exhibits higher disparity that may be

due to adaptations to various locomotor ecologies [12,35]. In contrast to these morphologically-

67 localized studies, analyses of the evolution of whole-body traits like body mass, skeletal size,

and body shape often follow a Brownian motion model or clade-based shift model rather than

69 being associated with ecological regimes [26,36,37].

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70 Compared to skeletal system-specific findings, simultaneous investigation of skulls, 71 limbs, vertebrae, and overall body plan are rarely conducted, likely because of the enormous 72 amount of data that would need to be collected and the complexity of the multivariate analyses 73 required. However, a more comprehensive approach to quantifying skeletal evolution is essential 74 to elucidate its complexity more fully. The search for system-level trends and variations is 75 further obscured by the disparate methods employed to test the effects of ecology and phylogeny 76 on different skeletal systems by different researchers. In this study, we address both issues in our 77 investigation of the mosaic evolution of carnivoran skeletons by creating a new phenomic dataset 78 that encompasses all major components of the skeletal system and using a unified set of 79 multivariate evolutionary models to test the ecological and phylogenetic effects influencing the 80 modes of evolution of these skeletal components.

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82 Methods

83 Skeletal and ecological traits

84 We collected 103 linear measurements to capture the skeletal morphology of 119 carnivoran species (208 osteological specimens; Fig. S1; Table S1). This dataset includes seven 85 86 cranial traits, seven mandibular traits, 13 forelimb traits, 13 hindlimb traits, and seven traits in 87 third cervical, fifth cervical, first thoracic, middle thoracic, diaphragmatic thoracic, last thoracic, 88 first lumbar, middle lumbar, and last lumbar vertebrae. Because carnivorans exhibit differing 89 degrees of sexual dimorphism [38,39], we use only male specimens. To remove size effects, we 90 calculated log shape ratios by dividing each skeletal trait by the geometric mean of all 103 traits 91 [40,41]. We then used principal component analyses (PCAs) to reduce the dimension of each 92 skeletal component (i.e., cranium, mandible, forelimb, hindlimb, and each of the nine vertebrae)

and retained a number of PC axes that corresponded to >90% of the explained variance. We also

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94 conducted a PCA on the entire dataset as our proxy of the whole-skeleton phenome and retained 95 the first six PC axes (~75% of explained variance) for subsequent analyses. We classified the 96 119 carnivoran species into distinct locomotor modes, hunting behaviors, and dietary regimes 97 following [37]. 98 99 *Phylogenetic comparative methods* 100 We tested whether each skeletal component evolved as adaptation to specific ecological 101 regimes or exhibited clade-specific evolutionary shifts by fitting multivariate evolutionary 102 models on the retained PC axes of each skeletal component [42–44]. For the adaptive ecological 103 models, we fit three multivariate multi-optima Ornstein-Uhlenbeck models (i.e., mvOUM_{diet}, 104 mvOUM_{hunting}, and mvOUM_{locomotion}) to test if dietary, hunting behavioral, or locomotor regimes 105 influenced the evolution of each skeletal component using mvMORPH [44]. The models were fit 106 across 500 stochastically mapped trees to account for uncertainty in phylogenetic topology and 107 ancestral character states (see electronic supplementary materials). We also calculated the 108 phylogenetic half-lives of the best supported adaptive ecological model [42]. A short 109 phylogenetic half-life relative to the age of Carnivora (48.2 myr) would suggest that skeletal 110 traits are strongly pulled toward distinct ecological optima across the adaptive landscape. For the 111 clade-based model, we fit a multi-optima OU model (mvOUM_{phyloEM}) without a priori ecological 112 regimes with PhylogeneticEM [45]. We also fit a single-rate multivariate Brownian motion 113 model (mvBM1) and a single-optimum OU model (mvOU1). We assessed the relative support of 114 models using small sample-corrected Akaike weights (AICcW). Lastly, we assessed the 115 covariation among skeletal components using partial least squares with geomorph [46].

Preliminary results revealed that phenotypic differences between pinnipeds (i.e., seals and sea lions) and terrestrial carnivorans are often the greatest source of variation for most skeletal components. These results are unsurprising considering pinnipeds exhibit derived morphologies that enable them to be fully aquatic. Therefore, we repeated our analyses using a reduced dataset with no pinnipeds. Results of the full dataset with pinnipeds are presented in the electronic supplementary material.

122

123 **Results and Discussion**

124 We found mosaic evolution of the carnivoran skeleton in which ecology and phylogeny have differing influences on the evolutionary mode of the various skeletal components. 125 126 Consistent with [6,27], the cranium and mandible exhibited decoupled evolutionary modes. In 127 the cranium, the clade-specific shift model exhibited overwhelmingly greater support 128 (mvOUM_{phvloEM}; AICcW>0.99) compared to adaptive ecological models (Fig. 1; Table S2). We 129 found eight evolutionary shifts in cranial morphology that correspond to carnivoran clades (Fig. 130 2A). In contrast, the adaptive dietary model was the best supported model ($mvOUM_{diet}$; 131 AICcW=0.96) for the mandible with a short phylogenetic half-life of 2.52 myr (Fig. 1; Fig. S2B; 132 Table S2; see Supplementary Results for optima distribution in phylomorphospace). These 133 results are congruent with findings revealing that mandibular shape is evolutionarily labile with 134 respect to dietary evolution whereas cranial shape is partitioned among families rather than 135 among dietary groups [6]. Despite their covariation (r = 0.73; Table S3), decoupled evolutionary 136 modes between the cranium and mandible may be explained by their functions. Diet is often 137 found to have had a strong influence on mandibular evolution because of its direct role in feeding 138 [3,47–51]. In contrast, the cranium has multiple sensory functions in addition to feeding that

influence its evolution [52–54], and therefore, the signal from dietary adaptations in itsmorphology may be obscured.

141 The appendicular system exhibited decoupled evolutionary modes between forelimbs and 142 hindlimbs. The forelimb was best supported by the mvOUM_{phyloEM} model (AICcW>0.99; Fig. 1; 143 Table S2). Seven shifts in forelimb evolution occur primarily along familial branches (Fig. 2B), 144 indicating that the complexity and variation of carnivoran forelimb morphology cannot be 145 captured effectively by dietary, hunting behavioral, or locomotor categories. Instead, these shifts 146 suggest that clade-specific adaptations enabled the diversity of forelimb skeletons for tasks such 147 as grappling or manipulating prey, swimming, or digging [28–31,34,55,56]. For example, most 148 felids use their prehensile forelimbs to ambush and subdue prey, most canids and hyaenids 149 pounce and pursue prey, and some mustelids use their powerful forelimbs to dig out prey while 150 other more derived mustelids (i.e., weasels) pursue prev in tight crevices and burrows [57]. In 151 contrast, the hindlimb was best supported by the mvOUM_{locomotion} model (AICcW=0.83) in the 152 hindlimb with a short phylogenetic half-life of 5.05 myr (Fig. 1; Table S2), supporting 153 hypotheses that the hindlimb is adapted primarily for locomotion as typically found in 154 quadrupedal mammals [58]. Although the forelimb and hindlimb covaries (r = 0.87; Table S3), 155 previous work found that this integration is weaker than expected in carnivorans that do not 156 specialize in cursoriality [34]. This work together supports the hypothesis of functional 157 divergence between the forelimbs and hindlimbs of carnivorans. 158 The axial skeleton exhibits distinct evolutionary modes between the anterior and 159 posterior regions of the vertebral column: cervical and most thoracic vertebrae tended to be best 160 supported by clade-specific shift or single-peak OU models, whereas the last thoracic and all

161 lumbar vertebrae were best supported by $mvOUM_{hunting}$ or $mvOUM_{locomotion}$ models (Fig. 1; Fig.

162 2C-F; Table S2). Our findings strengthen the coalescing hypothesis that anterior vertebrae 163 exhibit lower disparity, higher evolutionary constraints, and more subtle adaptations to 164 locomotion whereas posterior vertebrae exhibit the opposite patterns in carnivorans [35] and 165 broadly across mammals [13]. We posit that high evolutionary constraints of the anterior 166 vertebrae are associated with clade-specific shifts in the cervical and most thoracic vertebrae. 167 Importantly, subtle adaptations in these anterior vertebrae could be masked by many-to-one or 168 one-to-many mappings, making it difficult to uncover the form-function associations with 169 evolutionary models [59]. In contrast, relaxed evolutionary constraints of the posterior vertebrae 170 facilitate the evolution of disparate lumbar vertebrae across the entire carnivoran order. These 171 disparate vertebrae adapt to diverse locomotor modes or hunting behaviors based on the mobility 172 of the posterior backbone and irrespective of clade origins. The short phylogenetic half-lives 173 (1.47–5.12 myr) further suggests strong pulls towards these different adaptive optima. More 174 broadly, this increased mobility of the lumbar region over evolutionary time is hypothesized to 175 be an innovation characterizing crown mammals [13,60,61]. Correspondingly, the posterior 176 vertebrae are tightly integrated (r = 0.84-0.96; Table S3).

Lastly, we found that the clade-specific shift model ($mvOUM_{phyloEM}$; AICcW > 0.99) best 177 178 described the overall skeletal phenome (Table S2), a pattern that is consistent with previous 179 investigations of whole-body proxies such as body size and body shape [26,36,37]. The 180 mammalian body plan is comprised of cranial, axial, and appendicular components; therefore, its 181 multidimensionality transcends one-to-one mapping relationships between morphology and 182 ecological function. Instead, individual skeletal components within distinct body plans can adapt 183 to specific ecological factors independently from each other, enabling species with distinct body 184 plans to exhibit similar ecological or functional regimes and vice versa.

185 Overall, we elucidate the mosaic evolution of the carnivoran skeleton, finding that 186 different skeletal components exhibit distinct modes of evolution. Our results suggest that 187 different methodologies and taxonomic samples do not necessarily explain previously reported 188 region-specific macroevolutionary patterns; rather, complexity in explanatory factors of skeletal 189 diversity is a key feature of Carnivora. The ability of individual skeletal components to adapt to 190 specific ecological factors independently from each other may have contributed to the clade's 191 hierarchical [62,63] evolution. As previously hypothesized [26,36], the restriction of carnassial 192 shear to the P4/m1 pair may have been the key innovation that facilitated the initial carnivoran 193 diversification early in the clade's evolutionary history. Subsequent evolution led to the 194 continual partitioning between clades, resulting in the origination of extant carnivoran families as 195 discrete phylogenetic clusters that occupy different adaptive zones [64] with distinct 196 morphologies including body size and shape [37,65] and various components of the skeleton 197 ([6]; Fig. 2). Within-clade variation then arises to reflect resource partitioning among 198 ecologically similar taxa, leading to adaptations in morphologies such as the mandible, hindlimb, 199 and posterior region of the vertebral column (Fig. 1). These traits were strongly pulled toward 200 distinct ecological peaks across the adaptive landscape as revealed by their short phylogenetic 201 half-lives (1.47–5.12 myr) relative to the clade's age (48.2 myr).

Our research statistically revealed the mosaic evolution of carnivoran skeletons. These distinct evolutionary modes demonstrate the importance of examining multiple skeletal components in ecomorphological analyses. Nevertheless, key questions remain: What spurred the evolutionary transitions towards the evolutionary shifts or adaptations of the various skeletal components? When in the 55 million years of carnivoran evolutionary history did these evolutionary events occur? And what developmental and genetic phenomena underlie the

208	evolutionary	dissociation o	f various	skeletal	elements?	Ongoing	work integr	rating the	e fossil	and

- 209 paleoenvironmental record will further elucidate the carnivoran diversity we see today and reveal
- the complexity of evolutionary processes in multicomponent systems.
- 211

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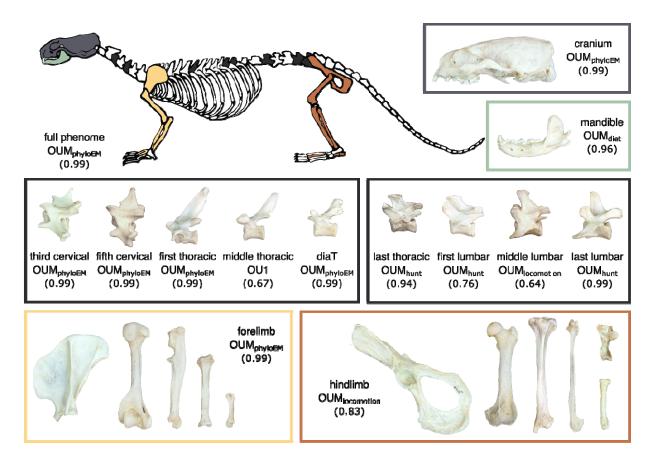
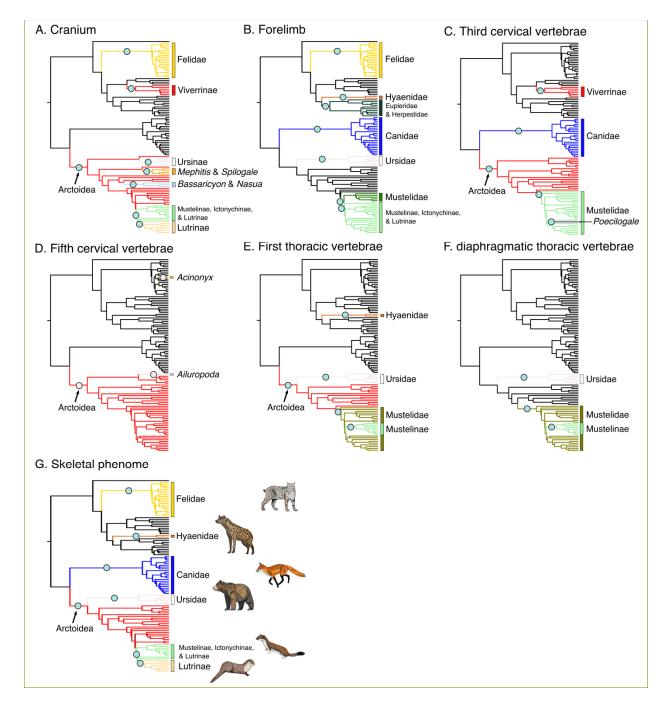


Fig. 1. Diagram of the skeletal components and their best-fitting evolutionary model on *Lontra canadensis*. AICcW are in parentheses. See Table S2 for full AICc table. diaT = diaphragmatic

thoracic vertebrae



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- Fig. 2. Clade-specific evolutionary shifts in skeletal components across terrestrial carnivorans
- 230 identified by PhylogeneticEM. Shifts are represented as pink circles, and branches on the
- 231 phylogenies are colored according to each regime.

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