Feeding ecology has shaped the evolution of modern sharks

Highlights

- Shark tooth morphologies track changing habitats and resource availability
- Tooth shape correlates with diet in extant shark species
- Declines in lamniform disparity can be linked with dietary “specialization”
- Modern lamniforms are more disparate than coeval carcharhiniforms

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In brief

Bazzi et al. analyze the evolution of lamniform and carcharhiniform shark over the last 83 Ma. These closely related clades are shown to have undergone marked morphological segregation, with a combination of habitat change, prey availability, and feeding strategies influencing their community composition, diversity, and ecology over time.
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SUMMARY

Sharks are iconic predators in today’s oceans, yet their modern diversity has ancient origins. In particular, current hypotheses suggest that a combination of mass extinction, global climate change, and competition has regulated the community structure of dominant mackerel (Lamniformes) and ground (Carcharhiniformes) sharks over the last 66 million years. However, while these scenarios advocate an interplay of major abiotic and biotic events, the precise drivers remain obscure. Here, we focus on the role of feeding ecology using a geometric morphometric analysis of 3,837 fossil and extant shark teeth. Our results reveal that morphological segregation rather than competition has characterized lamniform and carcharhiniform evolution. Moreover, although lamniforms suffered a long-term disparity decline potentially linked to dietary “specialization,” their recent disparity rivals that of “generalist” carcharhiniforms. We further confirm that low eustatic sea levels impacted lamniform disparity across the end-Cretaceous mass extinction. Adaptations to changing prey availability and the proliferation of coral reef habitats during the Paleogene also likely facilitated carcharhiniform dispersals and cladogenesis, underpinning their current taxonomic dominance. Ultimately, we posit that trophic partitioning and resource utilization shaped past shark ecology and represent critical determinants for their future species survivorship.

INTRODUCTION

Macroevolutionary studies aim to explain the drivers of large-scale biological phenomena and often rely on fossils to elucidate the complex processes determining modern biodiversity. 1–3 One frequently documented pattern is for closely related clades to manifest different radiation profiles over their extended geologic history. 4–6 A notable example is the extreme asymmetry in species richness evident between modern lamniform and carcharhiniform sharks. These lineages are globally distributed in today’s oceans but have starkly skewed diversities of 15 versus >290 living species, respectively. 7 By contrast, the global fossil record shows that lamniforms were taxonomically more diverse than carcharhiniforms before the end-Cretaceous mass extinction, 66 Ma. 8,9 The possible causes of this dramatic turnover include widespread oceanic cooling and intensified competition, which potentially promoted carcharhiniform diversifications during the Cenozoic. 10 Conflictingly, though, assessments of dental disparity (morphological variability) suggest that trophic cascades and prey availability moderated the evolution of lamniforms and carcharhiniforms across the Mesozoic–Cenozoic transition. 9

Sharks (Selachimorpha) are an optimal group for reconstructing macroevolutionary patterns because their dentitions are continuously replaced via a sequentially regenerative process termed polyphyodonty. 11–13 The subsequent prolific production of shed teeth, which incorporate a decay-resistant dentine and enameloid composition, has resulted in an abundant and ubiquitous fossil record. 14 To date, research on shark macroevolution has focused mainly on taxon counts and phylogenetic inferences to interpret diversity trends. 8,10,15,16 Corresponding evaluations of dental disparity have also hitherto been limited in their geographic 17 and chronostratigraphic 9 sampling.

Here, we undertake the first global-scale examination of lamniform and carcharhiniform dental disparity and morphology across the last 83 Ma. This time frame encompasses the Campanian–Maastrichtian ages preceding the Cretaceous/Paleogene (K/Pg) mass extinction event and the entire Paleocene–Holocene interval leading up to the present day. We combine fossil and extant shark teeth to reconstruct deep-time patterns of evolution and evaluate the role of biotic and abiotic factors in driving diversity asymmetries between lamniforms and carcharhiniforms. With these data, we test the hypotheses that (1) sea level and temperature 15 have functioned as major abiotic regulators of selachimorph evolution and (2) the covariation between tooth shape and diet, 16,17 as extrapolated
from living sharks, can be used to infer biotic mediators of past ecological change.

RESULTS

We compiled images of 3,837 extinct and extant shark teeth and converted them to a 2D landmark dataset for geometric morphometric analysis (Figures 1A, 1B, S1A, and S1B; Data S1). A Procrustes variance metric and a principal component analysis were used to estimate dental disparity and reconstruct morphological successions, respectively (STAR Methods).20,21 Additionally, we correlated tooth shape with published information on gut contents from living shark species 22 to establish covariation between tooth shape and diet as a basis for inferring feeding ecology across our morphospace time series. Finally, to evaluate environmental correlates, 23,24 a series of first-order autoregressive, AR(1), generalized least-squares (GLS) models compared disparity against sedimentary measures of sea-level and benthic oxygen isotope ($\delta^{18}O$) paleotemperature estimates.

Dental morphospace

Approximately 86% of the total variance in the dataset is explained by principal components (PC)1–PC3 (Figures 1C, S2A, and S2B). PC1 (64.8% of the total variance) describes a range from apicobasally tall and mesiodistally narrow to short and wide tooth morphologies (Figure 1C). PC2 (11.43%) captures distally recurved teeth with reduced cusplets to upright and mesially angled teeth with prominent shoulders that bear denticles or cusplets. Lastly, PC3 (9.54%) represents mesiodistally expanded and apically rounded teeth to mesiodistally tapered teeth incorporating well-developed lateral cusplets (Figure S2C). Lamniforms and carcharhiniforms share similar distributions across this morphospace but are statistically differentiated along PC1 and PC2 (Figure 1C), although not on PC3 (Data S2A).

Disparity dynamics

Lamniform disparity was high throughout the Late Cretaceous and across the K/Pg interval (Figure 2A; Data S2B). However, a significant long-term decline commenced in the late Paleocene (Maastrichtian versus Thanetian: $p_{RRPP} = 0.041$) and persisted until the late Eocene (Priabonian). Lamniform disparity was seemingly high during the Oligocene and early Miocene. However, this peak diminished by the Rupelian once heterodonty was integrated into the disparity estimation model (Figure S3A; Data S2C–S2E). A significant disparity decrease (Figure 2A) from the late Oligocene-early Miocene (Aquitanian/Burdigalian) to the mid- to late Miocene and on into the early Pliocene (Zanclean) was similarly lost when diagnathic, but not monognathic, heterodonty was integrated (Data S2C–S2E). Family-level partial disparity assessments (Figure S3B) specifically correlated low mid-Miocene (Langhian) disparity with lamnids and otodontids (lineages including the white shark, *Carcharodon carcharias*, and “megatooth shark,” *Otodus megalodon*, respectively). Late Pliocene (Piacenzian) lamniform disparity was otherwise comparatively high (Figure 2A; Data S2B), despite the small sample size ($n_{Lamniformes} = 12$), and peaked in the Holocene ($n_{Lamniformes} = 340$). Most conspicuously, Holocene lamniform disparity exceeds that of carcharhiniforms ($PV_{Lamniformes} = 0.070$; $PV_{Carcharhiniformes} = 0.059$; $p_{RRPP} = 0.004$), even after “specialized” filter-feeding lamniforms were excluded ($PV_{Lamniformes} = 0.067$; $PV_{Carcharhiniformes} = 0.059$; $p_{RRPP} = 0.023$).
Carcharhiniform disparity peaked during the Late Cretaceous (Campanian), declined across the Campanian–Maastrichtian interval ($p_{RIPPP} = 0.04$), and then remained largely static throughout most of the Cenozoic (Figure 2 A; Data S2 F). Langhian disparity estimates are notably high, though not significantly different from preceding or succeeding ages. Family-level partial disparities also showed that Paleocene triakids failed to recover their pre-K/Pg disparity levels, with carcharhinids becoming the principal contributors to carcharhiniform disparity by the early Eocene (Ypresian)—a pattern that continued to the Holocene (Figure S2 C).

Generalized least-squares models revealed significant correlations between lamniform disparity and both global sea level and $\delta^{18}$O paleotemperature estimates (Figure 2 B), with greater likelihood support than the intercept-only null models (Table 1). Notably, each 1 m sea level increase (Data S2 G and S2 H) corresponded to a 0.00025 increase in lamniform dental disparity ($p_{GLS} = 0.014$). Carcharhiniform disparity was best explained by a temperature model, where each 1 degree increase in temperature correlates with a 0.0018 increase in dental disparity ($p_{GLS} = 0.002$) but still with some sea-level interaction (Table 2; Data S2I and S2J).

Examination of the Cenozoic-only time bins using alternative environmental datasets ($n_{Lamniformes} = 1,452; n_{Carcharhiniformes} = 1,833; Data S2K and S2L) yielded more ambiguous results, with lamniform disparity best fitting the sea-level and null models, along with some support for the $\delta^{18}$O model (Data S2K). By contrast, Cenozoic-only carcharhiniform disparity was best explained by the sea-level model, with some support for both the $\delta^{18}$O and null models (Data S2L).

**Morphospace dynamics**

PC1 showed a shift toward taller, narrower teeth and the development of lateral cusplets among lamniforms across the K/Pg boundary ($p_{ANOVA} = 0.005$; Figure 3 A). Carcharhiniforms otherwise maintained their characteristically low-crowned, distally curved tooth morphologies, with a positive shift from the late Eocene to late Oligocene (Priabonian versus Chattian:...
pANOVA = 0.032; Figure 3A). Lamniforms and carcharhiniforms then converged on PC1 by the Holocene (Figure 3A; Data S2M–S2V).

PC2 and PC3 illustrated a reduction in negative morphospace representing distally recurved and high-crowned triangular tooth morphologies among lamniforms across the K/Pg interval but with some recovery by the mid-Eocene (Lutetian–Bartonian: Figures 3B and 3C; Data S2M, S2Q, and S2R). Carcharhiniforms otherwise remained largely stable but showed a significant negative shift (pANOVA = 0.008) at the Paleocene/Eocene boundary on PC2 (Figure 3B; Data S2U).

Lamniform morphospace redistributed along PC3 at the Oligocene/Miocene boundary. Initial average positive scores during the Aquitanian–Burdigalian became strongly negative by the Langhian and continued until the Holocene (Figure 3C; Data S2R). Negative scores along PC3 also coincided with a positive shift on PC1 (Figure 3A) and an overall decrease in lamniform disparity during the Miocene (Figure 2A; Data S2B).

### Dietary correlations

A phylogenetic two-block partial least-squares (pPLS) covariance analysis supported a significant covariation between tooth shape and diet in extant lamniform and carcharhiniform species (nSpecies = 55; τPLS = 0.577; p = 0.002; Z = 3.023; Figures 4 and S4). Lamniforms exhibited a high correlation coefficient (nSpecies = 9; τPLS = 0.862; p = 0.073; Z = 1.729) yet did not reach statistical significance. The lamniform morphocline extremes along PLS1 (block 1) reflected a dichotomy of macrophagous vertebrate and cephalopod predators with pointed teeth versus microphagous zooplanktivores with bulbous conical teeth (Figures 4C and 4D). Exclusion of specialized filter feeders produced comparable statistical results (nSpecies = 7; τPLS = 0.873; p = 0.09; Z = 1.382) but with contrasting tooth shapes ranging from mesiodistally narrow, pointed teeth in piscivores to shorter, triangular teeth in cephalopod hunters (Figures S5A and S5B).

The shape-diet pPLS correlation coefficient for carcharhiniforms (Figures 4E and 4F) was lower but significant (nSpecies = 46; τPLS = 0.584; p = 0.002; Z = 3.538). PLS1 captured durophytory on decaped crustaceans versus predation on bony fish and other chondrichthyans (Figures 4E and 4F). Except for piscivorous and, to some extent, chondrichthyan-based diets, dietary loadings for carcharhiniforms on PLS1 contrasted with lamniforms (Figures 4C and 4E), especially when lamniform filter feeders were removed (Figures SS5A and SS5B).

The ordered dietary breadth (ODB) analysis57 of extant lamniforms and carcharhiniforms suggested that these groups utilize broadly similar prey breadths, with positively skewed ODB distributions indicating a general bias toward dietary specialization (Figures 5A and 5B). Phylogenetic ANOVA and Tukey’s HSD tests failed to reject ODB uniformity between lamniforms and carcharhiniforms at order- or family-level groupings (Figure 5C; Data S2W). Nevertheless, ODB distributions indicated a greater concentration of low ODB values in lamniforms (modal value = 0.1), reflecting stronger biases toward dietary “specialization”—e.g., toward cephalopods in thresher sharks (Alopiidae), piscivory in the goblin shark (Mitsukurina owstoni), and zooplanktivory in the basking shark, Cetorhinus maximus. By comparison, carcharhiniforms tended toward higher ODB values (modal value = 0.25; Figures 5A and 5B), with notable examples including the tiger shark (Galeocerdo cuvier) and blue shark (Prionace glauca), which utilized all but one of the dietary categories designated here.

Significant ODB associations were also found on PC2 (Pearson’s product-moment correlation: r = 0.402; p = 0.001; Figures S5C and S5D). This correlated “generalist” diets (high ODB values) with upright, mesiodistally narrow teeth flanked by prominent cusplets (positive PC2 scores) versus specialist diets with distally recurved teeth flanked by reduced cusplets. Finally, we detected no association between ODBs and intraspecific disparity in either lamniforms or carcharhiniforms (Figures S4E–S4G) or with monogastric heterodony in lamniforms as estimated from a subset of complete tooth rows representing 10 species (Data S2X).

### DISCUSSION

Our results demonstrate that (1) lamniforms and carcharhiniforms maintained comparable levels of disparity during the Late Cretaceous despite lamniform taxic dominance; (2) these clades morphologically diverged after the K/Pg boundary; (3) carcharhiniform disparity has been largely stable since the Mesozoic, but with some family-level increases in the early Eocene; (4) lamniform disparity declined during the mid-Miocene–early Pliocene; (5) lamniforms exhibited high disparity...
during the late-Pliocene–Holocene; and (6) lamniform disparity currently exceeds that of carcharhiniforms despite their contrasting taxonomic richness (Figure 2A).

Despite static lamniform disparity across the K/Pg boundary, GLS model-fitting confirms that their decline throughout the Cenozoic can be linked to global sea-level and temperature reductions initiated across the K/Pg boundary (Data S2G and S2H). Specifically, the depletion of lamniform species diversity and a shift toward more “generalized” diets characterized by mesiodistally narrow tooth morphologies coincides with widespread epicontinental regression (e.g., the Western Interior Seaway of North America). Smaller-bodied lamniforms were especially dominant in these epeiric settings and thus may have been disproportionately impacted by latest Cretaceous eustatic change.9,30

Alternatively, the morphological divergence of lamniforms and carcharhiniforms (Figure 3A) does not support ecological competition as a regulator of their Cenozoic evolution.10 Indeed, such community interactions are difficult to discern from the time-averaged fossil record.31–34 Moreover, while some lamniforms and carcharhiniforms of equivalent body size do co-occur in modern habitats (e.g., C. carcharias with Carcharhinus brachyurus, Iaurus oxyrinchus with many carcharhiniform species, and Iurus paucus with Carcharhinus longimanus),7,35 most exhibit ecological differentiation into predominantly pelagic versus neritic environments.14,36 Combined with our discovery of Cenozoic ecomorphological divergence (Figures 3A and 4B), we therefore find no compelling reason to assume that population-level competition was a primary driver of lamniform and carcharhiniform macroevolution in the deep past.37

Implications for ecological evolution

Sharks are usually portrayed as the quintessential generalists, capable of ingesting a wide variety of food items. While this may be true for species like the modern Galeocerdo cuvier (OBD = 0.86), most selachimorphs display varying degrees of dietary specialization (Figure 5).39,42 Approximately 85% (47 out of 55) of extant lamniform and carcharhiniform species that we analyzed have ODBs < 0.5 (Figures 5A–5C), indicating a tendency toward more selective feeding. Unfortunately, without direct evidence from preserved gut contents,41 ODBs cannot be estimated for fossil sharks. Nonetheless, we detect a significant positive correlation between ODBs and morphospace loading along PC2 (Figure S5F). This correlation corresponds with positive loadings of lamniforms on PC2 following the K/Pg boundary and perhaps implies a short-term selection toward more generalized diets across the extinction event. Conversely, negative loadings on PC2 after the Paleocene–Eocene thermal maximum (PETM) implicates lamniform and carcharhiniform specialization. The PETM triggered major extinctions among actinopterygians (e.g., tetradontiforms) and presumably affected niche diversity among selachimorphs (e.g., Synchondrodoniformes).43 However, a shift toward more specialized diets during the PETM also coincided with the proliferation of biotically productive scleractinian coral reefs.44 These habitats are favored by carcharhiniforms today, especially carcharhinids,7,44 which underwent a disparity increase across the PETM (Figure S3C), potentially reflecting an adaptive accommodation of changing prey resource availability.45

While various developmental (e.g., palatoquadrate structure in lamniforms),46 reproductive (e.g., mating), behavioral (e.g., foraging habits), size-related (e.g., prey-size, body-size, and gape),36,43 and environmental (e.g., prey availability) constraints have influenced the evolution of shark dentitions, the primary function of their teeth is to capture and process prey, suggesting a link between tooth morphology and diet.9,48 Concomitantly, we evince a significant association between piscivory and mesiodistally compressed cusipidate anterior teeth (Figures 4C, 4D, and S4A–S4F), such as those of the sand tiger shark (Carcharias taurus), as well as with elongate “needle-like” teeth exemplified by Mitsukurina owstoni. The radiation of odontaspidid and mitsukurinid lamniforms during the early Paleogene could therefore correlate with an extinction-mediated shift toward generalized predation on teleosts.9 Piscivorous carcharhiniforms with low-crowned, distally recurved teeth, including catsharks (scyliorhinids), requiem sharks (carcharhids), hound sharks (triakids), and weasel sharks (hemigales),3,47 might have similarly diversified through trophic cascades instigated by changing prey resources and habitats.9

Certainly, these are the most speciose carcharhiniform clades today (Scyliorhinidae comprising 160 species, Carcharhinidae 56 species, and Triakidae 46 species),7 potentially linking their Cenozoic diversification with shifts toward distally recurved tooth morphologies after the K/Pg boundary (Figure S6).

The early Paleogene trend toward increasingly low-crowned, distally recurved tooth morphologies in carcharhiniforms

| Model                  | df | AIC  | BIC  | logLik | AIC.W | AICc  | AICc.W |
|------------------------|----|------|------|--------|-------|-------|--------|
| Without AR(1)          |    |      |      |        |       |       |        |
| null                   | 2  | −95  | −94  | 50     | 0.0060| −95   | 0.0119 |
| sea level              | 3  | −99  | −97  | 53     | 0.0441| −98   | 0.0558 |
| δ18O                  | 3  | −105 | −102 | 55     | 0.5974| −103  | 0.7567 |
| sea level + δ18O      | 4  | −103 | −99  | 55     | 0.2252| −100  | 0.1443 |
| sea level × δ18O     | 5  | −102 | −97  | 56     | 0.1274| −97   | 0.0312 |

With AR(1)

| Model                  | df | AIC  | BIC  | logLik | AIC.W | AICc  | AICc.W |
|------------------------|----|------|------|--------|-------|-------|--------|
| null                   | 3  | −96  | −94  | 51     | 0.0174| −95   | 0.0432 |
| sea level              | 4  | −99  | −96  | 54     | 0.0716| −96   | 0.0898 |
| δ18O                  | 4  | −103 | −100 | 56     | 0.5970| −100  | 0.7490 |
| sea level + δ18O      | 5  | −101 | −97  | 56     | 0.2211| −96   | 0.1061 |
| sea level × δ18O     | 6  | −100 | −94  | 56     | 0.0929| −92   | 0.0119 |
(Figure 3A) also corresponds with feeding on decapod crustaceans (e.g., the extant bonnethead shark, *Sphyrna tiburo*, and common smooth-hound, *Mustelus mustelus*; Figures 4E and 4F). Such morphologies are particularly abundant after the PETM, which we ascribe to an ecological expansion coincident with geographic dispersals and cladogenesis.48 Lamniforms, on the other hand, segregated into only a few distinctive dental morphologies that suffered synchronized disparity contractions during the mid-Miocene–early Pliocene (Figure 2A)—a time frame commenced by the global mid-Miocene climatic disruption.49 These contractions are most evident among lamnids and otodontids (Figures S3B and S7),8 which are lineages typified by apex predators, such as *Otodus megalodon* with an estimated maximum body length of 20 m.50 *Otodus megalodon* possessed high-crowned triangular teeth that closely resemble those of *Carcharodon carcharias* and was probably likewise specialized for feeding on marine tetrapods (e.g., cetaceans and pinnipeds) as an adult. The decline of lamniform apex predators across the mid-Miocene–early Pliocene has been attributed to climate change and competition51–53 but was possibly also influenced by niche specialization limiting their responsive capacity to environmental alterations.

**Implications for modern shark disparity**

The disparity peak among extant lamniforms and carcharhiniforms (Figure 2A) is robust to rarefaction subsampling and

![Figure 3. Dental morphospace through time](image)

Box and whiskers plots depicting lamniform (red) and carcharhiniform (blue) tooth morphologies on (A) PC1, (B) PC2, and (C) PC3. Visualization indicates the median (center line), 25% and 75% hinges (box), and whiskers and outliers (open circles). Tooth outlines (right) depict hypothetical morphologies at the minimum and maximum axis values. Sample sizes for each clade are listed in Data S1B and S1F. See also Figures S2, S6, and S7 for additional morphospace results.
demonstrates that sample size alone cannot explain our results for these groups (see discussion at https://doi.org/10.5061/dryad.h70rxwdjx). However, elevated extant disparity is undoubtedly influenced by the comprehensive sampling of living versus fossil species. For instance, geographic coverage in the fossil record is not equivalent to that of today (Figure S1). Irrespective, we note that our elevated modern disparity follows an increasing trend that commenced during the late Pliocene (Figure 2A): a signal that is unaffected by heterodonty. Heterodonty can enhance food processing and prey utilization (e.g., as observed in the bull shark, *Carcharhinus leucas*). Yet it can also accompany dietary specialization, as in the sicklefin weasel shark (*Hemigaleus microstoma*), which feeds primarily on cephalopods. As a result, monognathic heterodonty (calculated as disparity within a single jaw) does not strictly correlate with our ODB values (Figures S5C and S5D), implying that niche
breadth (as defined by variation in feeding ecology) does not constrain within-jaw dental variation in living lamniforms and carcharhiniforms.

Comparisons with Late Cretaceous lamniforms (Figure 2A) show that extant species manifest less negatively skewed tooth morphologies ($g_{1_{PC1:Maari}} = -0.592$ versus $g_{1_{PC1:Holocene}} = -0.016$; Figure 3A). Positive PC1 values correspond to apex predators, such as Carcharodon carcharias, and zooplanktivores, including Cetorhinus maximus. Therefore, we posit that the decline of lamniform tooth disparity during the Cenozoic was possibly related to an asymmetrical erosion of morphospace, in which extant lamniforms parallel the disparity, but not the ecology, of their more diverse Mesozoic antecedents. Nevertheless, our ODBs (Figure 5B) highlight the dietary specialization of extant lamniforms (note that C. carcharias exhibits ontogenetic dietary changes toward specialization; see discussion at https://doi.org/10.5061/dryad.h70twxj9), and their corresponding extinction susceptibility, as was observed across the K/Pg boundary. Despite their noted asymmetrical diversity, we discovered that extant lamniforms are more disparate than their taxonomically richer carcharhiniform contemporaries, planting lamniforms as “numerical relicts,” the surviving members of a far more speciose group in the distant geologic past.

Conclusions

The drivers of morphological disparity are complex and, in the case of sharks, have involved an interplay of various abiotic and biotic factors. Our geometric morphometric analysis of the exceptionally rich 83-Ma dental fossil record of lamniforms and carcharhiniforms shows that feeding ecology, coupled with sea-level and temperature change, were primary regulators of their deep-time evolution. Despite the confounding biases of geologic sampling and intraspecific variation through heterodonty, we pinpoint the end-Cretaceous mass extinction, PETM, and mid-Miocene climatic disruption as pivotal events affecting lamniform and carcharhiniform dental morphology, ecology, and diversity. In particular, major turnovers among lamniforms can be correlated with dietary specialization and the loss of apex predator lineages at the K/Pg and mid-Miocene transitions. We therefore recognize food resource and habitat utilization as key predictors of extinction sensitivity and underscore their role in shaping the asymmetrical diversity of lamniform and carcharhiniform sharks today.

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**SUPPLEMENTAL INFORMATION**  
Supplemental information can be found online at https://doi.org/10.1016/j.cub.2021.09.028.

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**AUTHOR CONTRIBUTIONS**  
M.B. designed analytical protocols, collected and analyzed the data, created the figures, and wrote the manuscript. N.E.C. designed analytical protocols, contributed data, and wrote the manuscript. B.P.K. contributed to data interpretation and wrote the manuscript. M.B., N.E.C., B.P.K., and P.E.A. contributed data, and wrote the manuscript. B.P.K. contributed to data interpretation and writing. M.B., D.A., Fowler, and Compagno, L.J. (2013). Sharks of the World: A Fully Illustrated Guide (Wild Nature).

**DECLARATION OF INTERESTS**  
The authors declare no competing interests.

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STAR METHODS

KEY RESOURCES TABLE

| REAGENT or RESOURCE | SOURCE |
|---------------------|--------|
| Deposited data      |        |
| Morphometric data sources | This paper | Data S1; https://doi.org/10.5061/dryad.h70xwdjx |
| Table Results       | This paper | Data S2; https://doi.org/10.5061/dryad.h70xwdjx |
| Supplementary results and methods | This paper | Data Dryad Supplementary File 1; https://doi.org/10.5061/dryad.h70xwdjx |
| Landmark coordinate data | This paper | Data Dryad Supplementary File 2; https://doi.org/10.5061/dryad.h70xwdjx |
| Sliders file        | This paper | Data Dryad Supplementary File 3; https://doi.org/10.5061/dryad.h70xwdjx |
| Maximum clade credibility tree | Stein et al. | Data Dryad Supplementary File 4; https://doi.org/10.5061/dryad.h70xwdjx |
| Environmental dataset 1 | Zachos et al., Prokoph et al., Miller et al. | Data Dryad Supplementary File 5; https://doi.org/10.5061/dryad.h70xwdjx |
| Environmental dataset 2 | Westerhold et al. and Miller et al. | Data Dryad Supplementary File 6; https://doi.org/10.5061/dryad.h70xwdjx |
| Diet compositions and trophic levels of sharks | Cortés, Yano et al., Bizzarro et al., Gonzalez-Pestana et al. | Data Dryad Supplementary File 7; https://doi.org/10.5061/dryad.h70xwdjx |
| Diet of Successive Size Classes of White Shark | Hussey and Dudley | Data Dryad Supplementary File 8; https://doi.org/10.5061/dryad.h70xwdjx |
| R source code       | This paper | R Scripts; https://doi.org/10.5061/dryad.h70xwdjx |

Software and algorithms

| tpsDig2          | Rohlf | http://www.sbmorphometrics.org/ |
|------------------|-------|--------------------------------|
| R                | R Development Core Team | https://www.r-project.org/ |
| geomorph (R package) | Adams et al. | 4.0.0 |
| RRPP (R package)  | Collyer and Adams | 1.0.0 |
| nlme (R package)  | Pinheiro et al. | 3.1-145 |
| ordiBreadth (R package) | Fordyce et al. | 1.0 |
| phytools (R package) | Revell | 0.7-20 |
| geiger (R package) | Pennell et al. | 2.0.7 |

RESOURCE AVAILABILITY

Lead contact
Further information on materials, datasets, and protocols should be directed to and will be fulfilled by the Lead Contact, Mohamad Bazzi (mohamad.bazzi@uzh.ch).

Materials availability
The landmark coordinate data used in all the analyses can be accessed from Data Dryad under DOI: https://doi.org/10.5061/dryad.h70xwdjx.

Data and code availability
All code used in this study and other previously published phylogenetic, ecological, and environmental data are available at the sources referenced in the Key resources table. Landmark data, the sliders file, R scripts, and other metadata are accessioned for open access on the Dryad data repository.
EXPERIMENTAL MODEL AND SUBJECT DETAILS

Conceptual background and limitations
Our taxonomic designations were constrained at order- and family-level. We avoided analytical use of genus and species-level taxa because these classifications can be contentious for fossil sharks. In addition, the biotic drivers of lamniform and carcharhiniform diversity are ambiguous, as opposed to morphological variability, which provides a more direct proxy for ecological macroevolution. While dental morphology represents an unavoidably constrained dataset, it is demonstrably robust to confounding factors, such as sampling and intraspecific variability through heterodonty (see Data Dryad Supplemental File: [https://doi.org/10.5061/dryad.h70rxwdjx](https://doi.org/10.5061/dryad.h70rxwdjx)). Nonetheless, we acknowledge that heterodonty is currently only evidenced in fossil lamniforms. Our disparity estimates for fossil carcharhiniforms may therefore be inflated, yet our lamniform-based results suggest that observed differences between time-bins are likely independent of tooth positional variation in both clades.

Uncertain family-level assignments necessitated some consensus referrals for convenience: (1) *Carcharias taurus* included in Carcharidae rather than Odontaspididae; (2) 73 teeth attributed to the extinct genus *Paranomotodon* were treated as incertae sedis; (3) *Cretomyrhinidae* was limited to the genus *Cretomyrhus*; (4) *Archaeolamnidae* was limited to the genus *Archaeolamna*; (5) *Otodontidae* was limited to the genus *Cretalamna*. While family-level representations are incomplete for some time-bins, we found that these yielded recoverable patterns with which to interpret the coarser order-level analyses.

METHOD DETAILS

Institutional abbreviations
WAM, Western Australian Museum, Perth, Western Australia, Australia. ROM, Royal Ontario Museum, Toronto, Ontario, Canada. FMNH, Florida Museum of Natural History, Gainesville, Florida, U.S.A. NCMNS, North Carolina Museum of Natural Sciences, Raleigh, North Carolina, U.S.A.

Data sources and geographic context
We sourced primary images of fossil shark teeth from the literature (Data S1). Original photographs of teeth from extant species were obtained from specimens held in the collections of the WAM, ROM, FMNH, and NCMNS, with some also sourced with permission from Shark References ([https://shark-references.com/](https://shark-references.com/)) and J-Elasmo ([http://www.elasmo.com/](http://www.elasmo.com/)) (see Data Dryad Supplemental File: [https://doi.org/10.5061/dryad.h70rxwdjx](https://doi.org/10.5061/dryad.h70rxwdjx)). We acknowledge that our geographic and stratigraphic information is subject to historical sampling biases, with assemblages from North America and Europe having disproportionate representation versus those from Asia and South America, which are usually restricted to Cretaceous and Paleogene deposits (Figure S1). Similarly, extant species sampling is largely restricted to catches recorded from the Atlantic, Pacific, and Indian oceans.

Temporal scope
We explored morphological variation over time using an age-based chronostratigraphic scale (Data S2). Initial assessments of our data indicated that sampling varied between time-bins. In particular, the Selandian, Aquitanian, and Serravallian were found to be data deficient. Consequently, to maximize data usage, we combined the Danian/Selandian, Aquitanian/Burdigalian, and Serravallian/Tortonian time-bins, and tested an Epoch-based time-scale for the Eocene, Oligocene, and Miocene (see Data Dryad Supplemental File: [https://doi.org/10.5061/dryad.h70rxwdjx](https://doi.org/10.5061/dryad.h70rxwdjx)).

Environmental data
To evaluate abiotic drivers of lamniform and carcharhiniform dental evolution, we obtained a published dataset of sedimentary sea-level measures combined with δ18O values converted to deep-sea paleotemperatures ([Environmental Dataset 1](https://doi.org/10.5061/dryad.h70rxwdjx)). We also evaluated a second dataset of δ18O estimates compiled for the entire Cenozoic time interval ([Environmental Dataset 2](https://doi.org/10.5061/dryad.h70rxwdjx)). These well-established climatic and eustatic proxies are recognized regulators of marine vertebrate biodiversity change, and have been employed in previous studies on lamniform and carcharhiniform macroevolution.

Geometric morphometrics
Shape variables were acquired using two-dimensional landmark-based geometric morphometrics (GM). Fixed landmark and sliding landmark designations followed published protocols for shark tooth image digitization. Digitization procedures were standardized to trace the outermost perimeter of each crown from its junction with the root toward the apex along the mesial and distal margins (Figures 1A and 1B). This resulted in two open curves comprising 157 sliding semi-landmarks and three fixed landmarks located at the tooth apex and the mesial and distal crown-root junctions (Figure 1B). Landmark digitization was performed in tpsDig2 v.2.32 with point equidistance and standardization routines written in R. Non-shape attributes of scale, orientation, and position were filtered from the landmark dataset using a generalized Procrustes analysis (GPA) implemented in the R package.
A bending energy criterion was computed during the GPA procedure to determine the final position of the sliding semi-landmarks.

**QUANTIFICATION AND STATISTICAL ANALYSIS**

All procedures and analyses were performed in the *R* statistical environment v.4.0.5.

**Multivariate ordination and visualization**

Tooth shape variation was visualized using a PCA of the covariance matrix obtained from the Procrustes shape coordinates. Morphants was assessed across PC1–PC7; however, only PC1–PC3 were retained for further analysis (Figures S2, S6, and S7). Collectively, these axes summarize ~86% of the variation and correspond to recognizable tooth morphotypes. A description of minimum and maximum morphologies associated with these axes is provided in the main text. Differences between time-bins and taxa were evaluated using a non-parametric analysis of variance (ANOVA) incorporating a randomized residual permutation procedure (RRPP) for hypothesis testing. A false discovery rate method was used to limit the rate of false positives in p value statistics.

**Randomization test**

Randomization tests of the extant sample used sub-sampling at predetermined intervals (n = 10, 20, 30, 40) with replacement determined from the frequency distributions for each major clade. We limited testing to PC1 because this component captured the highest percentage of the morphological signal.

**Disparity and rarefaction**

Procrustes variance (*PV*) was used to measure disparity directly from the Procrustes shape coordinates. *PV* is defined as the summed squared distances of all observations to the grand mean divided by the degrees of freedom; this is equivalent to the covariance matrix trace divided by the number of observations, and computed using the `morphol.disparity` function in *geomorph*. Pairwise differences between time-bins were evaluated using the permutation procedure in *geomorph*. We applied rarefaction subsampling to investigate the effects of sample size on clade-specific and temporal disparity. Sub-sampling levels were determined from the lowest sampled time-bin in the data. Bootstrap prediction intervals were computed for raw and rarefied disparity using 999 pseudoreplicates.

**Generalized least-squares models**

Sedimentary sea-level and δ¹⁸O paleotemperature values were correlated with disparity using generalized least-squares (GLS) linear models with a maximum likelihood criterion for parameter estimation. Serial autocorrelation was assessed using complete (ACF) and partial autocorrelation functions (PACF), as well as Ljung-Box tests. ACF calculates the correlation between lagged residuals, whereas PACF determines the number of autoregressive *p* orders needed for suitable model construction. Autocorrelations from both GLS and ordinary least-squares (OLS) model residuals were inspected by plotting a predetermined number of lags (pₖ = 12) against their corresponding autocorrelations. Significant autocorrelations were deemed to be those outside 95% confidence intervals. Although we failed to detect any significant autocorrelations, models that included an autoregressive correlation structure received better support. We provide both GLS and OLS results.

To accommodate for contrasting time-binning schemes, we used arithmetic means for sedimentary sea-level and δ¹⁸O values allocated to each geochronological age. Five separate GLS models were fit to calculated disparities and evaluated using the Akaike Information Criterion corrected for small sample sizes (AICc) and likelihood ratio tests. GLS analyses were carried out over the latest Mesozoic–Cenozoic time bins and again over a Cenozoic-only time-bin subset. All GLS models were run using the `glS` function in *nlme* v 3.1-145.

1. ‘Null model’. Provides a single parameter intercept-only model whereby shark disparity was described as stochastic (= Brownian motion).
2. ‘Sea-level model’. Provides a two-parameter slope + intercept model that described a relationship between changing sea-level and shark disparity.
3. ‘δ¹⁸O model’. Provides a two-parameter slope + intercept model that described a relationship between changing δ¹⁸O paleotemperature values and shark disparity.
4. ‘Sea-level+δ¹⁸O model’. Provides a three-parameter slope + slope + intercept model that described a multivariate relationship between changing sea-level, δ¹⁸O paleotemperature values, and shark disparity.
5. ‘Sea-level x δ¹⁸O model’. Provides a four-parameter slope + slope + interaction + intercept model that described a multivariate relationship between changing sea-level, δ¹⁸O paleotemperature values, and the interaction between changing sea-level and δ¹⁸O paleotemperature values, together with shark disparity.
**Effects of heterodonty on disparity**

We evaluated the potential effects of heterodonty in lamniforms via direct disparity and pairwise comparisons between time-bins (Figure S3). We used three separate morphological models:

1. A monognathic model (n\textsubscript{Lamniformes} = 1345) with standardized tooth positions following the literature (Data S1). Monognathic positional categories include ‘parasymphyseal’, ‘anterior’, ‘lateroposterior’, and ‘posterior’\textsuperscript{30}. These were designated for convenience and do not correspond to strict anatomical configurations.\textsuperscript{46}

2. A dignathic model (n\textsubscript{Lamniformes} = 1196) differentiating the upper and lower dental units.

3. A combined-heterodonty model (n\textsubscript{Lamniformes} = 907) allowing for interaction between tooth positions and dental units.

**Ecomorphological analyses**

Lamniform and carcharhiniform diets were categorized using published gut contents recorded from live-caught specimens of 9 lamniform and 46 carcharhiniform species.\textsuperscript{22,63–65} Food item classifications included: tetrapods (mammals, birds, reptiles); osteichthyans; chondrichthyans; cephalopods; benthic molluscs; decapod crustaceans; other invertebrates; zooplankton; and plants. These data were transformed using arcsine square-roots and then subjected to a covariation analysis with the Procrustes coordinates via a two-block partial least-squares (PLS) analysis.\textsuperscript{80} Correlation coefficients (r\textsubscript{PLS}) were used to evaluate PLS strengths with a permutation procedure (RRPP at 999 iterations) for significance testing.\textsuperscript{69,81} Alternate correlations were also performed for all taxa, and again at order-level (Figures 4 and S4). In addition, we re-ran the lamniform-only analysis excluding specialized filter-feeding lamniforms, *Cetorhinus maximus* and *Megachasma pelagios*, which skewed the initial PLS analysis (Figure S5). Shape changes at the PLS axial extremes were visualized using thin-plate splines (TPSs).\textsuperscript{82,83}

Finally, we computed a phylogenetic two-block PLS using the *phylo.integration* function in *geomorph*. Our preferred tree topology was pruned from the most inclusive molecular phylogeny\textsuperscript{58} and calibrated across 200 random trees with the maximum-clade credibility tree derived using *phytools* v. 0.7-20\textsuperscript{1} and the *geomorph* libraries.\textsuperscript{68}

**Ordinated dietary niche breadth**

Ordinated dietary niche breadths (ODBs) were calculated at species-level (n\textsubscript{species} = 100) using a pairwise Jaccard dissimilarity matrix of ‘presence/absence’ food item category scores,\textsuperscript{22} and analyzed with the *ordiBreadth* v.1.0 package.\textsuperscript{27} The dissimilarity matrix was then ordinated through a principal coordinates analysis (PCoA) from which ODB\textsubscript{rawi} is expressed as:

\[
\text{ODB}_{\text{rawi}} = \frac{1}{v} \sum_{k=1}^{v} \sum_{j=1}^{p} \sqrt{(x_{ijk} - c_{ik})^2}
\]

where \(p\) represents the total number of food items per species \(i\), \(v\) is the number of ordinated axes, \(c_{ik}\) is the centroid for the \(i\)th species on axis \(k\), and \(x_{ijk}\) is the position for the \(j\)th food item used by the \(i\)th species on the \(k\)th axis.\textsuperscript{27}

Significant differences between order- and family-level clades were determined using a phylogenetic ANOVA available through the *aov.phylo* function in *geiger*\textsuperscript{27} v. 2.0.7 and via a posthoc Tukey’s (HSD) test (\(z = 0.05\)). Lastly, we compared ODBs between these clades using a non-parametric Kolmogorov–Smirnov (K–S) test.