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Dental Microwear Texture Analysis of Pliocene Bovids from Four Early Hominin Fossil Sites in Eastern Africa: Implications for Paleoenvironmental Dynamics and Human Evolution Dental Microwear Texture Analysis of Pliocene Bovids from Four Early Hominin Fossil Sites in Eastern Africa: Implications for Paleoenvironmental Dynamics and Human Evolution

> A dissertation submitted in partial fulfillment of the requirements for the degree of Doctor of Philosophy in Environmental Dynamics

# By

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> May 2012 University of Arkansas

## ABSTRACT

Many researchers have suggested that Pliocene climate change was a motive force for human evolution. The basic idea is that a shift toward drier, more open settings, led to adaptations for bipedality and the consumption of savanna resources, including large grazing mammals. However, more recent paleoenvironmental reconstructions suggest that Pliocene hominins occupied variable or mosaic habitats, including both open and closed settings. Many techniques have been used to refine our understanding of the paleoenvironments of eastern Africa; however these have not led to consensus reconstructions. At Kanapoi, ecological diversity analysis indicates that at least part of the site was composed of closed woodland forest; however, taxonomic uniformitarianism of bovid taxa suggests a dry, arid habitat. Similarly contradictory reconstructions exist for Allia Bay, with paleosol analysis and palynology suggesting a mosaic habitat dominated by savanna, and taxonomic uniformitarianism of faunal assemblages suggesting an environment composed of gallery forest, open woodland, floodplains and edaphic grasses. The Laetoli faunal assemblages have also led to varying reconstructions, with some suggesting habitats as disparate as open grassland and closed woodland. Hadar has been reconstructed as a shifting mosaic environment, with various proxies supporting different levels of habitat fluctuation. This dissertation aims to test these opposing hypotheses by bringing a new, independent dataset for the inference of diet, and by extension habitats of actual individuals in the days before death. I use dental microwear texture analysis to reconstruct ratios of graze to browse in the diet and therefore ecological contexts of fossil bovids from Kanapoi, Allia Bay, Laetoli and Hadar. This dissertation tests competing hypotheses concerning early hominin habitats at Kanapoi, Allia Bay, Laetoli and Hadar and how ecological settings may have

changed over the temporal span of *Australopithecus anamensis* and *A. afarensis*. It also serves as an important test of the principle of taxonomic uniformitarianism, often applied to fossil fauna.

High resolution casts of 220 fossil bovids from the four sites and 575 extant African bovids were scanned for dental microwear textures using a white-light confocal profiler. Four adjacent scans were collected from each specimen, resulting in a total work envelope of 204 x 276  $\mu$ m. The scans were then analyzed using Toothfrax and Sfrax software packages and compared to a database of extant bovids with known diets. The extant bovids collected for use in this study include 25 extant African taxa, representing the full obligate grazer-browser-frugivore continuum.

The extant bovids showed significant variation that separated the taxa predictably by known dietary category, with the exception of generalists and browser-grazer intermediates. Some variation was also noted within the dietary categories, suggesting seasonal and/or geographic variation. In general, browsing taxa had significantly higher values for complexity, heterogeneity and fill volume than grazing taxa, which evince higher values for anisotropy.

The microwear textures of the fossil taxa were compared to the extant database and classified by diet. The Kanapoi and Allia Bay samples indicated that the bovids were primarily browsers or browser-grazers intermediates, suggesting the presence of more wooded habitats. The Laetoli sample is dominated by various levels of mixed feeding, although the presence of grazing taxa suggests a complex mosaic habitat at the site. Finally, the Hadar sample, divided into three hominin-bearing members, showed an increase in the number of grazing taxa over time. This suggests that there may have been gradual aridification at the site during the occupation of the australopiths.

This dissertation is approved for

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# DEDICATION

This dissertation is dedicated, with love, to my father.

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# **Chapter Outlines**

This dissertation is presented as a series of three prepared articles, the standard format for the Doctoral Program in Environmental Dynamics. The first chapter introduces the project and its primary objectives. It also provides the requisite background material on the fossil sites included in the study and the use of dental microwear for dietary reconstruction. The subsequent chapters contain the three articles that comprise the body of the dissertation. Each article is presented in its entirety, including reference sections.

The first article, presented in Chapter Two, is entitled "*Dental microwear texture analysis* of extant African Bovidae". This article presents the comparative database developed to interpret the microwear signals of the fossil taxa. It addresses the significant differences among modern bovids with known diets and the amount of overlap that exists between dietary categories. This article also discusses the ability of dental microwear to detect seasonality and various levels of mixed feeding. It is the first comprehensive study on dental microwear textures of bovids, or indeed any cetartiodactyls. The paper is currently in press in the De Gruyter journal Mammalia: International Journal of the Systematics, Biology and Ecology of Mammals.

Chapter Three presents the second article, "Dental microwear texture analysis of fossil bovids from Hadar, Ethiopia: implications for the paleoenvironment of Australopithecus afarensis". In this article, the diets of the fossil bovids from the Sidi Hakoma, Denen Dora and Kada Hadar Members of the Hadar Formation are reconstructed. The paper also addresses environmental change over the timespan during which *Australopithecus afarensis* lived in the Afar triangle. This paper has been submitted to The Journal of Vertebrate Paleontology, published by Taylor and Francis. The last of the articles is presented in Chapter Four and titled, "*Paleoenvironmental change in Pliocene eastern Africa as inferred from dental microwear texture analysis of fossil Bovidae*". This article expands on the previous paper by adding dietary reconstructions of the bovids from Allia Bay, Kanapoi and Laetoli to the Hadar data. The paper explores what these reconstructions tell us about local environments at the four sites and the implications of this project for paleoenvironmental change during the middle Pliocene, when the early australopiths roamed eastern Africa. This paper will be submitted to The Journal of Human Evolution, published by Elsevier.

#### **Chapter One: Introduction**

My research to date has focused broadly on late Neogene habitat reconstruction and the ways in which organisms interact with their local environments. While I have worked with a wide variety of taxa, including living and fossil primates, carnivores, modern humans and early hominins, I selected bovids for my dissertation work because of their frequent use as paleoenvironmental proxies. Due to their abundance at hominin fossil sites, these taxa have been used to infer habitat based on presumed ecological similarity to their closest living relatives. The goal of this project was to reconstruct diets of fossil bovids from some of the most important early hominin sites in eastern Africa, using a non-genetic dietary signal. The resulting dietary reconstructions were then used tomake inferences about the types of habitats that would have been available to the local bovid populations.

In order to understand relationships between hominin evolution and environmental change, it is first necessary to have the best possible interpretations of the habitats occupied by our early forbearers. Techniques for reconstructing local paleoenvironments include the study of floral remains and geochemical analyses of paleosols. There have also been various types of studies focused on the faunal assemblages as proxies for paleohabitat, including taxonomic uniformitarianism, stable isotope analyses, ecomorphology and dental mesowear. These have unfortunately not offered consistent reconstructions for paleoenvironments at many important fossil sites, including the four Pliocene localities that are the focus of this dissertation. This dissertation uses dental microwear analysis of local fauna to evaluate the competing hypotheses. Microwear offers direct evidence of diet and can therefore provide important indications of the types of resources available during the occupation of the early hominins.

# **Objectives**

The goal of this dissertation project was to reconstruct the diets of bovids from homininbearing deposits at Kanapoi, Allia Bay, Laetoli and Hadar using dental microwear texture analysis to better understand the environmental contexts of the early hominins *Australopithecus anamensis* and *A. afarensis*, a putative lineage in eastern Africa persisting between about 4.2 and 3.0 million years ago. In order to interpret the microwear textures of these fossil bovids, the results were compared with an extant database that includes a broad range of modern ruminants with known dietary and habitat preferences. This has allowed me to evaluate previous reconstructions of the habitats occupied by the homininsand how their environmental milieu may have changed through the middle Pliocene. The results have been used in conjunction with previously published paleoecological and paleoclimatological data to provide finer resolution habitat reconstructions.

In sum, this project had three primary objectives. First, I sought to develop a large database of extant bovid dental microwear that could be used to interpret the microwear signatures of the fossil samples and be applied to other sites and time periods in Africa during future projects. Second, I set out to use this comparative database to reconstruct the diets of the fossil bovids from Kanapoi, Allia Bay, Laetoli and Hadar. An important component of this objective was to infer local habitat types based on the dietary preferences of the associated fauna and to compare these results with previously published habitat reconstructions. Finally, I aimed to test the principle of taxonomic uniformitarianism and determine whether or not it could be reliably applied to the fossil bovids from eastern Africa.

#### **Objective 1. Dental microwear texture analysis of extant African bovids**

Pilot work on extant bovids suggested that microwear texture analysis could separate not only grazers from browsers, but that it also held the promise of finer dietary distinctions, including obligate versus occasional graze, leaf versus fruit browse, etc. However, microwear texture analyses of living and fossil bovids have been limited to date (Ungar et al, 2007), and a larger baseline series was necessary to improve the resolution enough to reliably place fossil bovids into the number of diet categories typically used to separate and categorize living ruminants.

The resulting database includes 25 extant taxa representing all of the dietary classifications for bovids proposed by Gagnon and Chew (2000). The comparative sample also includes taxa reflecting seasonal and/or geographic variation in diet, in the hope of recognizing these patterns in the fossil species. In addition to extensive dietary variation, the taxa also represent a wide variety of habitat preferences, ranging from closed forest to open savanna grasslands. While it is always possible that environments existed in the past that are no longer present today, the baseline includes specimens from most modern African habitat types and should therefore provide a reliable basis for paleoenvironmental reconstruction.

#### **Objective 2. Diet and habitat reconstruction of fossil bovids**

The four sites selected for this study represent the temporal span of *A. anamensis* and *A. afarensis*. The fossil bovids from the hominin bearing deposits have been previously studied with techniques including taxonomic uniformitarianism and ecomorphology, with the aim of better understanding the adaptations of the taxa (for examples, see Kappelman et al., 1997; Spencer, 1997; Reed, 1997, 2008; Bobe et al., 2002; Kovarovic and Andrews, 2007). The results of these

studies, while valuable, are more reflective of the evolutionary adaptations of the bovid taxa than the actual diet, which can change rapidly with environmental conditions. Nongenetic signals, including dental mesowear and stable isotopes, have also been previously studied, but have generally been restricted to single sites and thus preventing a more regional look at *Australopithecus* paleohabitat (for example, see Schoeninger et al., 2003; Kingston and Harrison, 2007). This study aimed to reconstruct the diets of these taxa using anongenetic signal that reflect the actual diet of the animal during its lifetime.

To interpret the diets of the fossil bovids, the dental microwear signatures were compared to the extant comparative database. All of the fossil taxa analyzed fell within the range of one of Gagnon and Chew's six dietary categories: obligate and variable grazers, browser-grazer intermediate, generalist, browser or frugivore. This suggests that while different resources may have been available in the past, the bovids at these sites were processing foods in similar ways to those consumed by living ruminants.

The dietary reconstructions of the fossil taxa were also used to infer local paleoenvironment. While the relationship between dental microwear and habitat type is not exact, an abundance of browsing or grazing taxa does imply the presence of woodland forest or open grassland, respectively. The taxa from Kanapoi, Allia Bay, Laetoli and Hadar all suggest the presence of mosaic habitats during the occupation of the early australopiths. The earlier sites have higher numbers of bovids with microwear signatures similar to modern browsers, suggesting more heavily wooded habitats. Even at these sites though, at least some grazing taxa are present, indicting the inclusion of, or proximity to, more open grassland. These graze-adapted bovids are more common at the later sites associated with *A. afarensis*, suggesting a gradual aridification trend may have been present. However, the evidence presented here is not

consistent with the dominance of wide open grassland habitats, as previously suggested by 'the savanna hypothesis' (Dart, 1925).

#### **Objective 3. Testing taxonomic uniformitarianism of fossil bovids**

The principle of taxonomic uniformitarianism is often utilized by studies that rely on faunal assemblages for environmental reconstruction. This approach relies on the assumption that fossil species share the ecological preferences of their closest living relatives. The advantage of dental microwear is that it gives insights into what animals actually ate in the past rather than what they were capable of eating, as suggested by the sizes, shapes and structures of their teeth. Microwear mitigates the basic adaptationist assumptions that animals today eat the foods to which their ancestors were evolved, and that morphology reflects dietary preference.

Previous work by Sponheimer et al. (1999) at Makapansgat, South Africa, tested this assumed relationship using stable isotope analysis, also a non-genetic signal for diet. Their results suggested that the assumed diet was inconsistent with the isotopic signature in approximately 30% of the analyzed taxa. The results presented here suggest that this number may be even higher at the sites in eastern Africa. While most inconsistencies were only a matter of degree (i.e., variable grazer instead of obligate grazer), several were much more dramatic. For example, the Alcelaphini sample from Allia Bay is presumed to be most closely related to modern alcelaphines that practice obligate and variable grazing. The dental microwear signature of the fossils, however, is most similar to that of modern browsers.

## **Paleoclimate and Human Evolution**

Researchers have long hypothesized a relationship between Pliocene hominin evolution and global climate change (for examples, see Dart, 1925; Robinson, 1963; Howell, 1978; Laporte and Zihlman, 1983; Vrba, 1985, 1988, 1992, 1995a, 1995b, 2000; Stanley, 1992; deMenocal, 1995; Hill, 1995; Potts, 1996, 1998, 2007; Bobe et al., 2002; Trauth et al., 2005). The "savanna hypothesis" (Dart, 1925, 1953; Bartholomew and Birdsell, 1953; Washburn, 1960; Robinson, 1963; Jolly, 1970; Laporte and Zihlman, 1983; Vrba et al., 1989) suggested that bipedality and other morphological features of early hominins reflected adaptations for open, arid savanna habitats. However, more recent paleoenvironmental reconstructions have revealed a shift towards more variable environments (Kingston et al., 1994; Kappelman et al., 1997; Reed, 1997; Schoeninger et al., 2003).

*Australopithecus anamensis* is the earliest identified member of its genus, first appearing in the Turkana Basin 4.2 mya (Leakey et al., 1995, 1998) and represented by more than 80 fossil specimens from the Kenyan sites of Kanapoi and Allia Bay (Figure 1). *Australopithecus afarensis* first appears in the fossil record 3.6 mya, and possibly as early as 3.85-3.89 mya at Belohdelie, Ethiopia (Asfaw, 1987; Renne et al., 1999). The species is represented by more than 400 fossils, over 95% of which were found at the sites of Laetoli, Tanzania and Hadar, Ethiopia (Leakey and Harris, 1987; Lockwood et al., 2000; White et al., 2000) (Figure 1). *Australopithecus anamensis* is accepted by many paleoanthropologists as being ancestral to *A. afarensis* (Leakey et al., 1995; Wolpoff, 1999; Ward et al., 2001; Kimbel et al., 2006), and some have even argued that the two taxa represent a single chronospecies (Senut, 1996; Wolpoff, 1999). This view is primarily supported by the facts that *A. anamensis* predates *A. afarensis* with no known overlap between species and that the former is the more plesiomorphic taxa (Ward et al., 2001; Kimbel et al., 2006).

Most of the characteristics reflecting evolutionary change in Australopithecus between 4.1 and 3.0 mya are related to dental and locomotor adaptations. The molarization of premolars associated with A. anamensis and A. afarensis has been linked to Pliocene expansion of dry, open savanna; however, modern studies of marine sediment cores suggest that widespread C4 grasslands did not develop in eastern Africa until after 3.0 mya and would not have influenced the habitats of A. anamensis and A. afarensis (deMenocal, 1995; Leakey et al., 1995). Recent paleoenvironmental models instead suggest that a wide range of habitats were available to early hominins, including closed and open woodland, gallery forest, bushland, and wet and dry grassland habitats (Leakey et al., 1995; Reed, 1997; Wynn, 2000; Kingston and Harrison, 2001; Bonnefille et al., 2004; Campisano and Feibel, 2007). Therefore, the morphological adaptations of A. anamensis and A. afarensis could instead reflect increasingly fluctuating seasonal conditions across a range of locally available habitats, where the ability to utilize lower-quality foods as needed during "crunch times" would be advantageous (Teaford and Ungar, 2000; Ungar, 2004; Reed and Fish, 2005; Kimbel et al., 2006). The mosaic nature of reconstructed Pliocene sites also accords well with the inferred locomotor adaptations of the australopiths. Both A. anamensis and A. afarensis are characterized by postcranial adaptations allowing for both terrestrial bipedality and arboreal climbing (e.g., Rodman and McHenry, 1980; Johanson et al. 1987). This unique locomotor adaptation has been hypothesized to be the result of an environmental shift towards mosaic habitats, where the ability to walk bipedally and yet retain climbing proficiency would have been beneficial.

The hominin fossils themselves have been studied in an attempt to reconstruct their trophic adaptations and by extension, their paleoenvironments. Previous examination of the dental microwear of *A. anamensis* and *A. afarensis*, for example, has revealed no consistent changes in diet over time and suggested a narrow range of fracture properties (Grine et al., 2006; Ungar et al., 2010). This is surprising given the paleoclimatic reconstructions of Pliocene eastern Africa as a world of increasing climate variability, which would suggest an accompanying change in resource use. There are several possible explanations for the lack of variation in the dietary signature of these early hominins. It is possible that *A. anamensis* and later *A. afarensis* showed no significant directional change in diet during this period of climatic instability. It is also possible that the environmental change previously reported for middle Pliocene eastern Africa was not entirely regional, and that the basins occupied by early hominins may not have been as strongly affected by the climatic fluctuations. Finally, *A. anamensis* and later *A. afarensis* may have adopted a foraging strategy similar to modern chimpanzees, maintaining large territories and ranging long distances to acquire preferred resources.

## **Previous Paleoenvironmental Reconstructions of Hominin Habitats**

The Kanapoi fossil site is located on the southwest side of Lake Turkana, Kenya and includes fluvial and deltaic sediments accumulated during a major lacustrine phase dated between 4.2-4.17 mya (Harris and Leakey, 2003). As the earliest known site for *A. anamensis*, an understanding of associated paleoenvironments is important, though reconstructions have sometimes been inconsistent (See Table 1 for a summary). A study of paleosol stratigraphy by Brown and Feibel (1991) suggested that the Omo River flowed through the basin and into the Indian Ocean for much of the Pliocene, and that a series of temporary lakes were created when

tectonics disrupted the river outflow. The presence of this river system, combined with the results of stable isotope and palynological studies, has led researchers to interpret Kanapoi as a mosaic habitat (Brown and Feibel, 1991; Cerling, 1992; Bonnefille, 1995; Wynn, 2000). Ecological diversity analysis of the faunal assemblages from the hominin-bearing strata indicate that at least a portion of the site was closed woodland forest during its accumulation, similar to the reconstructed environment for the slightly older hominin *Ardipithecus ramidus* from Ethiopia (Reed, 1997). However, taxonomic uniformitarianism of the fossil bovids recovered from Kanapoi has suggest a drier and more arid habitat, and studies of the associated paleosols liken the paleohabitat to the savanna environment of the modern Omo Delta (Harris and Leakey, 2003).

The Allia Bay hominin site is also located in the Turkana Basin of Kenya, on the eastern shore of the modern lake. The site is dated to approximately 3.9 mya and the fossils were likely accumulated during a relatively short period of time (Leakey et al., 1995). As at Kanapoi, previous attempts to reconstruct the paleoenvironment have suggested a mosaic habitat with the presence of floodplain grassland, woodland forest and dry bushland around the site margins (Cerling et al., 1988; Feibel et al., 1991; Coffing et al., 1994; Leakey et al., 1995) (See Table 1 for a summary). Stable isotope analysis of the associated paleosols suggests a highly seasonal environment (Raymo et al., 1996; Wynn, 2000), with rainfall levels that remained virtually unchanged through the Pliocene and Pleistocene (Feibel, 1999). In fact, a study comparing stress lines in the teeth of fossil bovids to modern bovids from a variety of environments suggests that the paleoenvironment at Allia Bay was likely similar to the modern seasonal mosaic ecosystem of the Masai Mara (Macho et al., 2003).

The Laetoli site is located in northern Tanzania and consists of sediments that span the time range between 4.3 mya and 120 kya. *Australopithecus afarensis* has been found in the Upper Laetolil Beds dated to 3.7-3.5 mya. Several paleoecological reconstructions of Laetoli have been conducted and the results have been very inconsistent (See Table 1 for a summary). Reconstructions based on taxonomic uniformitarianism of mammal assemblages found in association with *A. afarensis* indicate fairly closed woodland (Andrews, 1989; Reed, 1997) or woodland with patches of forest (Kingston and Harrison, 2007), thoughinterpretations of the bovid assemblages recovered from the site have been confusing. Kaiser and Solounias (2003) used dental mesowear of fossil bovids to suggest multiple habitat types but few arid-adapted grazing taxa. However, other researchers have argued that the presence of grazing bovids including dik-diks and wildebeest, indicates a more open, arid habitat, as does the complete absence of aquatic species at the site (Harris, 1987).

The Hadar hominin site is located in the Afarregion of Ethiopia and has yielded more than 370 numbered specimens of *A. afarensis* to date, the most found at any site in eastern Africa. The site has four identified geologic members associated with early hominins: Basal (~3.8-3.42 mya), Sidi Hakoma (~3.42-3.26 mya), Denen Dora (~3.26-3.2 mya), and Kada Hadar (~3.2 mya) (Taieb et al., 1976; Campisano, 2007), each bounded by radioisotopically dated tephras. Horizons associated with *A. afarensis* have been dated to 3.4-3.0 mya and contain the last known individuals of the species. Previous paleoenvironmental reconstructions of Hadar have suggested a mosaic of microhabitats with slight fluctuations throughout the temporal range of *A. afarensis* that indicate a general trend towards drier and more open settings (See Table 1 for a summary). This reconstruction is supported by a wide variety of studies, including those of palynological remains (Radosevich et al., 1992; Bonnefille et al., 2004), ecomorphological and

diversity analyses of faunal assemblages (Bobe and Eck, 2001; Reed, 2008) and stable isotope data (Hailemichael, 2000). Isotopic analysis of paleosols at Hadar also suggest periodic shifts from wetter to drier environments suggestive of, according to Hailemichael (2000), a general trend from more closed to open habitats.

In sum, the reconstructions for these four important Pliocene hominin sites have been inconsistent and have, in some cases, yielded inconsistent results. These local habitat types have been used to explain the adaptations that distinguish *A. anamensis* and *A. afarensis* from one another, as well as from other hominins, and it is therefore critical that the reconstructions are as accurate as possible. Additionally, given the morphological and apparent dietary stasis of *A. anamensis* and *A. afarensis*, evaluating variation in food availability may improve our understanding of the adaptations of these anagenetic taxa. Therefore, if we hope to develop a clearer understanding of the adaptations of the early australopiths, one of the first steps must be clarifying our interpretations of their local environments, and one way of doing this is to look to the local fauna.

# **Bovids as Paleoenvironmental Indicators**

Bovids are generally considered to be reliable indicator species for environmental reconstruction both because they are ubiquitous at fossil hominin sites, and because modern ruminants fall into discrete dietary categories that reflect habitat preferences. These dietary categories refer to the percentages of monocotyledons (grasses, sedges and roots) and dicotyledons (fruits, seeds, flowers, buds, leaves, tubers and shoots). The commonly used dietary classifications are: obligate grazers, which inhabit open environments and eat >90% dicotyledons; browsers, which inhabit more closed environments and eat >90% dicotyledons;

and mixed feeders, which include species with highly localized and/or seasonal diets that alternate between grazing and browsing (Fortelius and Solounias, 2000). Actual bovid diets are often more complex than these categories suggest though, and many intermediate classifications have been proposed, including the subdivision of mixed-feeding taxa and the recognition of a separate frugivore category (Gagnon and Chew, 2000). Even given the oversimplification of a three category system (grazer versus browser versus mixed feeder), such a classification has proven valuable for interpreting the diets and habitat preferences of fossil bovids (e.g., Fortelius and Solounias, 2000; Merceron and Ungar, 2005; Schubert et al., 2006). Finer dietary distinctions for fossil taxa may be possible through dental microwear texture analysis, as suggested by a recent study on the fossil bovids from Langebaanweg by Ungar et al. (2007).

Habitats occupied by extant bovids range from deserts and rainforests. These allow us to determine microwear-environment associations for a broad variety of habitats. The use of bovid assemblages to make inferences about Pliocene hominin habitats is widespread in the paleoanthropological literature, and has been applied to sites in both eastern and South Africa (Kappleman, 1984; Vrba, 1980, 1985; Shipman and Harris, 1988; Harris, 1991; Plummer and Bishop, 1994; Spencer, 1997; Sponheimer et al., 1999; Merceron and Ungar, 2005; Reed, 2008).

Fossil bovids recovered from eastern African hominin sites include members of 12 biological tribes, often the most specific identification available for bovid specimens not accompanied by the distinctive horn cores. Aepycerotini (impala), Tragelaphini (kudu and browsing antelope), Reduncini (reedbucks and lechwe) and Bovini (cattle and spiral horned antelope) are typically browsers and associated with closed, moist habitats dominated by woodlands and fresh grasslands (Kingdon, 1982). Alcelaphini (wildebeest), Antilopini (gazelles), and Hippotragini (oryx antelopes and their kin) are predominantly grazers considered to be

indicators of open and seasonally arid environments dominated by grasslands (Kingdon, 1982; Gagnon and Chew, 2000). The other tribes, Cephalophini (duikers), Neotragini (dwarf antelope), Caprini (sheep and goats), Boselaphini (four-horned antelope), and Ovibovini (takin) are mixed feeders associated with various types of mixed habitats (Kingdon, 1982).

# Taxonomic uniformitarianism

Studies of faunal assemblages as proxies for the paleoenvironment of a fossil site typically apply the principle of taxonomic uniformitarianism, which is based on the assumption that fossil species share the same ecological preferences as their closest living relatives. Although this assumption is rarely questioned for recent faunas, its utility for reconstructing the paleoecology of long extinct species is uncertain. The primary problems with taxonomic uniformitarianism are that it is not applicable to fossil taxa with few or no living relatives, and that it assumes that groups remain constant in their ecological preferences over long periods of time. While taxonomic uniformitarianism is relatively straightforward and requires minimal effort beyond a quick glance at a faunal list, critics have questioned many of its inherent assumptions (Solounias et al., 1988; Reed, 1996; Sponheimer et al., 1999; Schubert et al., 2006). For example, Sponheimer et al. (1999) demonstrated that more than one-third of the Makapansgat fossil bovids had diets that differed from the assumed diets based on taxonomic uniformitarianism. While Aepyceros sp. and Gazella vanhoepeni were both assumed to be mixed/seasonal feeders like their closest living relatives, for example, isotopic signatures and ecomorphological data suggested that both were obligate browsers, and showed no evidence of C4 grass consumption. Due to these discrepancies between inferred and actual diet, it is critical

that ecological similarities of extant and fossil taxa be tested in some reliable way in order to have confidence in these assumed relationships.

#### **Dietary Reconstruction**

Diet is the most direct way that an organism interacts with its environment. Attempts to reconstruct diet in fossil forms have included the use of tooth size, shape and structure. However these are all genetic signals that reflect only what a species capable of eating, not what individuals actually ate. The diet of an individual may vary over its lifetime, but it can take many generations for the morphology of species to change in response to selective pressures. In order to reconstruct what an animal actually ate at a moment in time in the past, a non-genetic signal is needed. Dental microwear, or the microscopic pits and scratches created on the enamel surface during mastication, is one such line of evidence.

Differences in the dental microwear between grazing and browsing ungulates have been noted in previous studies of extant and fossil species (for examples, see Solounias et al., 1988; Solounias and Moelleken, 1993; Merceron and Ungar, 2005; Merceron et al., 2005; Schubert et al., 2006; Ungar et al., 2007), with browsers having more pits and grazers having more scratches. Mixed-feeding taxa tend to have intermediate microwear signatures, or patterns overlapping with browsers and grazers. This has important implications for reconstructing paleoenvironments, as while the connection between diet and habitat are not exact, grazing and browsing microwear signatures do imply the availability of grasses and woody plants respectively.

#### **Dental Microwear Analysis**

Conventional methods of studying dental microwear, including scanning electron microscopy and light microscopy, are subject to high rates of observer measurement error and a lack of depth data in two dimensional surface characterizations. Dental microwear texture analysis was developed to be a solution to these problems. It combines white-light confocal profilometry with scale-sensitive fractal analysis, resulting in the quantitative description of surfaces at a range of scales. This offers a suite of objective and repeatable measures of microwear surface textures in three dimensions that can be compared between groups with differing diets. Further, white-light confocal profilometry is quicker, easier to use, and less costly than scanning electron microscopy, and automated analysis requires less time and effort from researchers than does identification and measurement of individual features. The most important advance of texture analysis is the reduction of observer error in measurement, which allows direct comparisons between studies, as well as the establishment of a large database that researchers can access for interpretation of their results. Dental microwear texture analysis has been successfully used to distinguish dietary differences between species of extant and fossil taxa including humans and non-human primates, carnivores, bovids, kangaroos and wallabies (Scott et al., 2005, 2006; Ungar et al., 2008; Prideaux et al., 2009; Scott et al., 2009; Schubert et al., 2010).

Teeth are relatively common in the fossil record, as they are almost entirely mineralized and more durable than any other part of the skeleton. For this reason, diet reconstructions of extinct species, including early hominins, center on what data can be gathered and inferred from teeth. Teeth provide the most direct evidence of diet since they are the only part of the body to

come into direct contact with food. Additionally, much information can be gained from teeth in relation to jaw movement and diet. For example, form-function relationships have been described between tooth shape and diet in bovids. Grazing species tend to be more hypsodont, or high crowned, and this has been linked to offsetting the added wear caused by abrasives in grass forage (Fortelius, 1985; Janis, 1988, 1995; Mendoza and Palmqvist, 2008). Studies of these form-function relationships have also been conducted on primates. Species with sharp molar crests are associated with folivory and the need for shearing of tough foods, whereas those eating hard or brittle foods typically have flat, thick enameled surfaces for crushing and grinding (Kay, 1975; Kay and Hylander, 1978; Lucas, 1979, 2004). These studies have been complicated by the difficulty in studying teeth with worn cusps, yet wear often begins very early in life- sometimes even in utero (Teaford and Walker, 1983). Ungar and M'Kirera (2003) proposed a solution to this problem using GIS software, but even with this technology to reconstruct the occlusal surface, simply knowing what a species was capable of eating does not definitely indicate that this was their primary source of food. The diet of a species may change relatively quickly, but it can take many generations for the morphology of the tooth to follow.

One of the most reliable methods for reconstructing what an individual actually ate in the past is dental microwear analysis. This consists of examining the occlusal or buccal surface of a tooth with a microscope to document the scratches, pits, and gouges caused by foods of different properties during the mastication process. Microwear is created when the teeth come into occlusion during chewing. Scratches are formed on the enamel during the mastication of tough foods like fibrous plant material, much in the same way a pair of scissors would be scratched by abrasives on paper dragged between its blades during slicing. In this case, teeth come into contact at a steeper angle relative to the wear facet for shearing. Both maxillary and mandibular

molars can be examined for microwear. Previous studies have demonstrated no significant difference between them as an indicator of diet related wear (Teaford and Walker, 1984). It is standard to look for bovid microwear on the disto-buccal enamel band of the mesial cuspid of the  $M_1$  and  $M_2$  or the mesio-buccal enamel band of the mesial cusp of the  $M^1$  or  $M^2$ . (Janis, 1990; Merceron et al., 2005b).

# **Microwear Signatures on Living Species**

Dental microwear studies have been conducted on many living species, including bovids, and a direct connection has been made between the type of wear found on the teeth and the known and observed diets of the species. Moderate pit counts on the surface of a tooth indicate that the animal was likely a frugivore or specialized in soft foods. Scratches can also be found on the teeth of these animals if the soft food they eat is coated in dirt or other gritty material (Silcox and Teaford, 2002). If the pit counts are very high, the animal was probably a hard-object feeder, specializing in items like seeds or nuts (Strait, 1993; Silcox and Teaford, 2002). Folivores are usually distinguished by high counts of scratches on the molar surface from plant material being scraped across the tooth by the opposing molar (Teaford and Walker, 1984; Teaford, 1988a).

Because of the connection between food type and microwear pattern, the study of the dental microwear of fossils can reveal important clues about the diet and ecology of extinct species. Studies have been conducted on many types of fossil species, including rodents (Rensberger, 1978), ungulates (Solounias and Hayek, 1993; Solounias and Moelleken, 1992a, b, 1994; Solounias and Semprebon, 2002), carnivores (Van Valkenburgh et al., 1990), horses (MacFadden et al., 1999), giraffids (Solounias et al., 1988), tyrannosaurids (Schubert and Ungar, 2005), non-human primates (Teaford and Walker, 1984; Lucas and Teaford, 1994; Ungar, 1996;

Ungar and Teaford, 1996; King et al., 1999; Leakey et al., 2003; El-Zaatari et al., 2006), and human bioarchaeological populations (Rose and Marks, 1985; Harmon and Rose, 1988; Krueger and Ungar, 2010). There have also been studies of microwear in early hominin species in an attempt to answer questions about our own dietary heritage (Walker, 1981; Grine, 1981, 1986; Puech et al., 1983, 1986; Puech, 1986; Grine and Kay, 1988; Kay and Grine, 1989; Grine et al., 2006a, 2006b; Ryan and Johanson, 1989; Ungar and Grine, 1991; Lalueza Fox and Pérez-Pérez, 1993; Lalueza et al., 1996; Pérez-Pérez et al., 1999, 2003; Ungar et al., 2006; Teaford et al., 2007).

# The Development of Dental Microwear Studies

An interest in the markings left on teeth during mastication developed in 1933, when Simpson noted a correlation between the movement of the jaw and the principles of molar occlusion in mammals (Simpson, 1933). He suggested that the wear evinced on the teeth should correspond to specific movements of the jaw during mastication and should be indicative of diet. Later work demonstrated that molar facets, created by the occlusion of the upper and lower teeth, can be used to reconstruct the motions of the jaw. The preferred orientations of striations (if any) created on these surfaces reflect the direction of jaw movement (Butler, 1952; Mills, 1955, 1963, 1967; Butler and Mills, 1959).

While the first studies of microwear on the occlusal surfaces of teeth were focused on understanding jaw movements during mastication, studies soon began to emphasize the association between food types eaten by a species and microscopic wear patterns. These early microwear studies were conducted using binocular light microscopy and primarily qualitative analysis, and emphasized the link between diet and wear on the enamel (Baker et al., 1959;
Dahlberg and Kinzey, 1962; Walker, 1976). Baker et al. (1959) studied the microwear of caprine teeth to determine the impact of silica in grass and found that these abrasives contributed significantly to wear. Young (1986) later confirmed the contribution of abrasives by conducting SEM analysis on the incisors of Canadian moose. Despite the lack of quantitative data, these studies showed the potential of microwear for the reconstruction of diet in fossil species (Ungar et al., 2007).

Dahlberg and Kinzey (1962) suggested that studies of dental microwear might be used to study interspecific and intraspecific variation among populations. It was not until the late 1970s, however, that interest in microwear and diet really began to take off. Walker (1976) hypothesized that arboreal and terrestrial monkeys exhibited different microwear patterns because or foods available in different substrates. Combined with studies by Rensberger on rodents (1978) and Walker (1978) on hyraxes, a strong connection was inferred between microwear on the occlusal surfaces of teeth and the mechanical properties of foods.

With all of this potential, it quickly became apparent that a better approach than light microscopy, with its limited depth of focus, was needed. Rensberger (1978) and Walker et al. (1978) began using scanning electron microscopy (SEM) to look at microwear. Early SEM images taken of microwear surfaces on the teeth of small mammals were much clearer and allowed forobservation of more detail on the enamel surfaces. Researchers soon began to measure wear features on photomicrograph images, and standardized procedures were developed in an attempt to make these studies comparable (Peters, 1982; Gordon and Walker, 1983; Kay and Covert, 1983). These methods included using calipers or a digitizing table and later PC based measuring software (Ungar, 1995) to count and measure individual microwear features.

Small mammals were not the only subjects of early SEM studies. The hypothesized connections between dental microwear and food properties were being investigated using the teeth of fossil hominins (Grine, 1977, 1981; Puech, 1979; Ryan, 1980a, 1980b; Walker, 1981). While these studies were still qualitative, comparisons were made between fossil species and often included modern primate taxa. Walker (1981) speculated that similarities between early hominin microwear and that found on modern chimpanzees and orangutans suggested a frugivorous diet for*Paranthropus boisei*. Similar studies conducted on anterior tooth wear (Ryan, 1980a, 1980b; Ryan and Johanson, 1989) also included comparative samples, both nonhuman primates and recent human populations (Inuit and Micronesians), and suggested that early hominins might have used their incisors to strip away gritty plant parts like roots and seeds.

While much about the diets of extinct species has been learned from SEM-based microwear studies, these studies suffer from two primary criticisms. First, identification and measurement of wear features are highly subject to observer error. Intraobserver measurement error rates have been estimated at 7% and interobserver error rates at 9% (Grine, 2002). Secondly, SEM studies produce two dimensional images based on three dimensional surfaces, which causes the image to be affected the angle of the specimen on the microscope stage relative to the light or electron source (i.e., depth is sensed by shadows, so the dip and strike of the surface will affect the pattern of shadows) and the instrument settings (Gordon, 1988). Finally, these SEM studies are time-consuming due to the necessity of identifying and measuring up to hundreds of individual features on a single surface.

In the 1980s, several studies critiqued microwear techniques and their ability to differentiate between dietary patterns. Covert and Kay (1981) conducted a controlled experiment and attempted to control for diet. They used test groups of opossums, feeding them cat food

supplemented with insect chitin, plant fibers, or no additives. After the study, Covert and Kay were unable to distinguish between the groups eating different foods. They used this study to cite the inability of microwear techniques to distinguish between herbivory and insectivory. These conclusions were later questioned by Gordon and Walker (1983), who critiqued the Covert and Kay study due to inappropriate dietary mediums, the use of additives and insufficient quantification of the results. Gordon (1982, 1984) predicted that future refinements of microwear techniques would reveal differences in the opossums that were not seen by Covert and Kay and noted that her own studies of chimpanzee teeth had revealed that variation in microwear features could be partially accounted for by facet type, tooth position and individual information like age (Gordon, 1982). The Covert and Kay experiments called for a reevaluation of the limitations of dental microwear studies and the importance of a dietary control when categorizing specimens. Later studies focused on the necessity of standardizing the portion to the tooth scanned (Gordon, 1982, 1984), reducing interobserver and intraobserver error through automated practices (Kay, 1987; Grine and Kay, 1987; Walker et al., 1987), and the effect of enamel structure on tooth wear (Boyde 1984; Boyde and Fortelius, 1986; Maas, 1991).

The work of Covert and Kay also challenged researchers to demonstrate the ability of microwear studies to distinguish animals based on their diets. The work of Teaford and Walker (1984) utilized anthropoid primates to demonstrate that frugivorous species had more pits on their "Phase II" wear facets, while folivorous species had more scratches. Among frugivores, these authors recorded even higher percentages of pits for species that consumed hard objects, making three distinct dietary categories distinguishable through microwear studies of primates. Teaford and Oyen (1989) also conducted long-term studies on vervet monkeys, altering their diet and examining the microwear on the teeth. They have recorded differences in microwear signals

between animals eating hard monkey chow and soft wet chow. Animals eating the hard, dry chow showed significantly more microwear features than those eating soft foods. Their research also confirmed that pit frequency is related to the hardness of the food source.

New and more automated methods of image processing were introduced in an attempt to minimize interobserver error (Kay, 1978), but these still relied on SEM photomicrographs and the two dimensional images. A semi-automated analysis technique using these images was developed by Ungar et al. (1991) and Ungar (1995) and has been used consistently over the years, but is not immune to the problems associated with SEM, because it still requires time-consuming observer identification of microwear features (Teaford, 2006). Therefore, studies performed using these methods are still prone to subjectivity and high rates of observer error (Grine et al., 2002).

SEM-based analysis of microwear has been the most commonly used approach to dental microwear study. Recent investigations have used this technique with emphasis on reconstructing the diets of fossil species and other primates by using large comparative databases of modern species with known diets. SEM studies of Eocene omomyids (Strait, 1991), Egyptian Oligocene primates (Teaford et al., 1996), Miocene apes (Walker et al., 1994; Ungar and Teaford, 1996; Ungar et al., 1996), subfossil Malagasy lemurs (Rafferty et al., 2002), and early hominins (Grine et al., 2006a, 2006b; Ungar et al., 2006) have been conducted and yielded new information on the diets of fossil species. However, even with its continued use and the efforts to automate and standardize the technique, it has been suggested that SEM-based analysis has reached its limitations and that new methods must be sought (Teaford, 2006).

As noted by Rose and Ungar (1998), taking a single specimen through the process from preparation to data collection from SEM photomicrographs can take hours. This, predictably,

limits the number of specimens that can be included in a single study. In an effort to improve sample sizes, Solounias and Semprebon (2002) returned to low-magnification light microscopy, which involves the tallying of microwear features while viewing specimens though a binocular light microscope. This approach has been applied to a variety of taxa in recent years, including primates, bovids, equids, mammoths and even squirrels (Solounias and Semprebon, 2002; Kaiser, 2003; Godfrey et al., 2004; Semprebon et al., 2004; Nelson et al., 2005; Green et al., 2005; Godfrey et al., 2005; Rivals and Semprebon, 2006). Merceron and his colleagues have used a variant of this approach, combining light microscopy with the quantification of features on a computer screen (Merceron, 2003a, 2003b, 2004, 2005a, 2005b). The advantage to low magnification light microscopy is that itallows for the processing of large samples comparatively quickly and studies have reported low rates of interobserver error among experiences observers (Solounias and Semprebon, 2002). However, as Ungar et al. (2008) observed, this revived technique has not solved the problems with depth of field and magnification that led researchers away from light microscopy in the 1970s.

Dental microwear texture analysis combines white-light confocal profilometry with scale-sensitive fractal analysis, resulting in the quantitative description of surfaces at a range of scales. This offers a suite of objective and repeatable measures of microwear surface textures in three dimensions that can be compared between groups with differing diets. The technique has been successfully used to distinguish dietary differences between species of extant and fossil taxa including humans and non-human primates, carnivores, and bovids (for examples, see Scott et al., 2005, 2006; Krueger et al., 2008; Ungar et al., 2007, 2008, 2010; Scott et al., 2009; Krueger and Ungar, 2010; Schubert et al., 2010).

## **Limitations of Dental Microwear Research**

A factor considered early on to be a problem for microwear studies of fossils was that surface patterns might have been affected by postmortem or taphonomic damage. Taphonomic processes are those that affect the remains after death. Fossil teeth can sustain several different types of postmortem damage. Acid etching, damage from excavation tools or preservatives, water transport, and weathering are well established taphonomic agents that can easily obliterate microwear features. As the microwear scratches can be less than a single micron deep, they can be easily destroyed by taphonomic processes and even obscured by fingerprints. However this is not the complicating factor it was once thought to be. It is relatively easy for researchers to tell actual microwear from postmortem scratching of the tooth surface by rocks and dirt. Microwear is regular and found predictably on specific areas of the tooth, whereas damage to the enamel caused by being in the ground is randomly placed and not at all uniform (Teaford, 1988; King, 1999). In fact, the problem with postmortem wear is not that it confuses microwear researchers, but that it limits the number of useable samples for study. For example, according to Teaford (2006), only about 20% of the monkey specimens found at Koobi Fora, and only 60% of Olduvai material have been determined to be useable for microwear. A recent study on early hominin microwear supports this conclusion, suggesting that only approximately 23% of the hominin molars found throughout Ethiopia, Tanzania, Kenya, Malawi and South Africa display useable antemortem microwear (Ungar et al., 2006). This number also corresponds to similar estimates of Plio-Pleistocene monkey fossils from South Africa (El-Zaatari et al., 2005). The percentages could be even lower with bovid taxa, as the narrow bands of enamel that preserve microwear are easily damaged. Thus, the most serious complication of dental microwear studies on fossils is the small sample sizes for many species, and the limited number within these collections that are suitable for microwear analysis.

Dental microwear research has other limitations. While microwear studies have been successfully used to differentiate between short and long term diets and therefore the presence of seasonal diets, the microwear on a single tooth is not necessarily indicative of the overall dietary pattern of the animal's lifetime. It has been demonstrated that microwear only reflects a few days diet because recently created microwear overlays earlier wear (Walker et al, 1978). This is commonly known as the "Last Supper" phenomenon and is a complicating factor which must be considered in all microwear studies (Grine, 1986). Knowing this, seasonal variation in diet and the limited temporal range of what microwear can reveal can be taken into consideration. The potential to reveal dietary breadth makes the collection of large samples especially important in microwear studies, particularly when we can assume limited bias in the sample.

#### Summary

Environmental change is widely accepted to have been a motive force in human evolution and because of this, it is critical that our paleoenvironmental reconstructions are as detailed and robust as possible. Dental microwear, as direct evidence of actual diets of individuals, is an established method used in paleoenvironmental reconstruction free from assumptions inherent with taxonomic uniformitarianism. Further, because bovids are the most common taxa found at fossil sites and are known to have diets that vary with habitat, they are wellsuited to serve as environmental proxies. The results of this study not only test previous assessments of diets of fossil bovids from Pliocene hominin sites, but they provide more robust interpretations of early hominin habitats in eastern Africa based on faunal assemblages.

This project has generated the largest database of microwear textures for both extant and fossil bovids yet assembled, and will provide a substantial comparative collection for future research. It also serves as an important test of the principle of taxonomic uniformitarianism, which is often assumed in the use of faunal assemblages as paleoenvironmental proxies. The results of this study confirm previous work by Sponheimer et al. (1999) which suggests that taxonomic uniformitarianism may not be a consistently reliable predicator of bovid dietary preferences.

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Method of Reconstruction	Hypothesized Paleoenvironment	Source
Kanapoi, Kenya		
Paleosols- carbon isotopes 2000	Mosaic with semi-arid, seasonal cli	mate Wynn,
Paleosols- stratigraphy 1992;	Mosaic with gallery forest and edap	hic Cerling,
	grasslands E 1991	Brown & Feibel, 1986,
Faunal assemblages 1995:	Mosaic with gallery forest and dry, open	Leakey et al.,
	bushlands	Reed, 1997
Bovid Taxonomy Leakey, 2003	Arid savanna grassland	Harris &
Pollen assemblages	Included forest or woodland habitats E	Sonnefille, 1995
Allia Bay, Kenya		
Paleosols- carbon isotopes 2000;	Mosaic, but with dominant open say	vanna Wynn,
Paleosols- stratigraphy Floral assemblages Faunal assemblages 1994;	Raym Mosaic and highly seasonal Open woodland with expanses of grassland Mosaic with gallery forest, bushland, open	o et al., 1996 Brown & Feibel, 1991 Cerling et al., 1988 Coffing et al.,
Organic stable isotopes Reeser,	floodplain and riparian woodland Feibel et al., 1991 Mosaic with woodland and grassland Schoeninger &	
	1991	
Laetoli, Tanzania		
Paleosols- carbon isotopes Paleosols- stratigraphy Paleosols- stratigraphy Faunal assemblages	Closed, deciduous woodland forest At least a seasonal dry, arid climate Closed woodland to bushland Mosaic with heavy woodland A Reed	Cerling, 1992 Hay, 1987 Verdcourt, 1987 andrews, 1989; , 1997
Faunal assemblages Andrews,	Patch woodland and grassland	Harrison, 2005;
		2006, Kingston & Harrison, 2007
Faunal taxonomy 1980; Harris, 1985	Savanna grassland and open woodla	and Hay,
Bovid Taxonomy Pollen assemblages	Open arid savanna Mosaic with grassland and open woodland	Harris, 1987 Bonnefille &
Riollet, 1987 Hadar, Ethiopia		

**TABLE 1.**Summary table of previous paleoenvironmental reconstructions at fossil sites

Paleosols- carbon isotopes 2000	Seasonal habitats and shifts from open and Hailemichael,	
	closed woodlands	
Paleosols- stratigraphy	Grassy woodland and semi-arid climate	
Radosevich et al., 1992	2	
Faunal assemblages	Physiognomic habitat with habitats ranging from scrub to edaphic grassland	Reed, 2008 Bobe & Eck,
2001	1 0	,
Pollen assemblages 1987	Mosaic habitat that trended towards dry	Bonnefille et al.,
	open environments B	onnefille et al., 2004

FIGURE 1.Map of East African fossil sites



# Chapter Two: Dental Microwear Texture Analysis of Extant African Bovidae

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# Abstract

Bovids are often used as paleoenvironmental proxies because they are among the most commonly recovered large mammals at many fossil hominin sites and because modern African bovids occupy a wide range of dietary and environmental niches. This study uses dental microwear texture analysis to examine 25 species of extant African bovids, representing 6 dietary categories and with an emphasis on various levels of mixed feeding. The results show significant differences among the dietary classifications and confirm previous work suggesting that grazing taxa have less complex, more anisotropic surfaces with smaller features than browsing taxa. The results also indicate that dental microwear texture analysis can distinguish beyond the classic grazer-browser-mixed feeder trichotomy and accurately separate variable grazers, generalists, browser-grazer intermediates and frugivores from obligate grazers and browsers, as well as from one another. Some differences among taxa within dietary categories were also found, likely reflecting seasonal and/or geographic differences in diet. In addition to demonstrating the effectiveness of the technique at differentiating between different levels of mixed feeding in bovids, this study also provides a comprehensive comparative dataset of extant bovid microwear textures that can be applied to fossil taxa from sites and time periods across Africa

Key Words: microwear, bovid, grazer, browser, diet

## Introduction

The Family Bovidae is an ecologically diverse group of ungulates represented by more than 45 genera spread across Africa, Asia, Europe and North America (Wilson and Reeder, 2005). African bovids are particularly varied and include 27 extant genera and 79 species, making them one of the dominant herbivorous mammals on the continent (Kingdon, 1997). They inhabit a broad variety of ecosystems- from desert to tropical rainforest (Grzimek, 1990). The diets of African bovids are equally varied and range from habitual grass grazers such as the common tsessebe (*Damaliscus lunatus*), to browse specialists like the gerenuk (*Litocranius walleri*), to highly frugivorous taxa such as the blue duiker (*Philantomba monticola*). Even as ubiquitous as these bovids are today, the fossil record shows that they were even more abundant in the past, with over 100 fossil taxa from the Pliocene and Pleistocene described to date (Gentry, 1990; Vrba, 1995). An understanding of both modern and fossil bovid ecology is therefore crucial if we hope to better understand the paleoenvironments associated with human evolution.

Reconstructions of paleoenvironment that utilize faunal remains are only as strong as the extant baseline used to interpret them. There is a wealth of literature available on modern bovid diets and considerable work has been done to compile this information into a comprehensive system of classification (Gagnon and Chew, 2000). This system distinguishes not only grazing and browsing taxa, but also recognizes several levels of mixed feeding, an important advance that can be useful in informing habitat preference. However, even with clearer dietary categories for extant species, distinguishing between various levels of mixed feeding in the fossil record is difficult using traditional methods of dietary reconstruction. Dental microwear texture analysis, which combines white-light confocal microscopy with scale-sensitive fractal analysis, is an objective and repeatable method for characterizing microwear surfaces. Most importantly, this

technique has the potential to get beyond the classic grazer-browser-mixed feeder trichotomy and distinguish taxa that engage in various levels of mixed feeding, as suggested by a previous study on the fossil bovids at Langebaanweg, South Africa (Ungar et al., 2007). In this paper, I test this hypothesis using a new comparative baseline series of extant bovid dental microwear textures. This dataset includes 25 species of modern African bovids, representing a wide variety of dietary preferences. These data provide a baseline series that can be used to interpret the microwear signatures of fossil bovids from sites and time periods across Africa.

The dietary categories typically used to classify bovids refer to the percentages of monocotyledons (grasses and sedges) and dicotyledons (fruits, seeds, flowers, buds, leaves, twigs, tubers and shoots) that make up their diet. Traditionally, these percentages have been used to divide bovids into three distinct dietary classifications: grazers, browsers and mixed feeders (Vesey-Fitzgerald, 1960; MacNaughton and Georgiadis, 1986; Hoffman, 1989; Estes, 1991; Fortelius and Solounias, 2000). The grazer and browser categories reflect specific dietary and habitat preferences, with the former including taxa that inhabit open environments and eat mostly monocotyledons, and the latter representing taxa that inhabit more closed environments and primarily eat dicotyledons. The generalized mixed feeder category, however, refers to a highly diverse group of bovids that display a myriad of environmental and dietary preferences. These taxa are not static in the foods that they consume and many have highly localized and/or seasonal diets that alternate between grazing and browsing.

Actual bovid diets are more complex than this grazer-browser-mixed feeder category scheme suggests, and several more detailed classifications have been proposed. Hofmann and Stewart (1972), for example, proposed a three category system based on differences in extant bovid stomach structure. Taxa were classified as bulk-roughage eaters (grazers), concentrated herbage selectors (browsers) or intermediate feeders, and were further subdivided based on consumed percentages of each food type. Langer (1988) built on these categories by adding three more levels of intermediate feeding. A later re-classification by Bodmer (1990) recognized the importance of fruit in the diet of many African bovids and proposed a dietary continuum ranging from grazing to dedicated frugivory, with browsing being the intermediate form. More recently, Gagnon and Chew (2000) described a comprehensive system of dietary classification for African bovids based on food type, seasonality, geographic variation and body mass. Their system proposed six primary dietary categories: obligate grazers (> 90% monocots), variable grazers (60-90% monocots), browser-grazer intermediates (30-70% monocots and dicots, < 20% fruit), browsers (> 70% dicots), generalists (> 20% all food types) and frugivores (> 70% fruits). Sponheimer et al. (2003) successfully used stable carbon isotope analysis to test the dietary information for 24 of the taxa classified by Gagnon and Chew and found that, with a few exceptions that could be explained based on sample population differences, the results of the two studies yielded similar dietary classifications.

Diet is the most direct way that an organism interacts with its environment and has had significant influence on the evolutionary adaptations of grazing and browsing ungulates. Previous attempts to identify morphological indicators of diet have included taxonomic uniformitarianism and ecomorphological proxies like the configuration of the masticatory structure, as well as the use of tooth size, shape and structure (Solounias et al, 1995; Pèrez-Barberia and Gordon, 2001; Archer and Sanson, 2002). However, these are all genetic signals that reflect only what a species is capable of eating, not what individuals actually ate. The diet of an individual may vary over its lifetime, but it can take many generations for the morphology of a species to change in response to selective pressures (Solounias and Moelleken, 1999). In order

to reconstruct what an animal actually ate at a moment in time, epigenetic signals must be examined. Dental microwear, or the microscopic pits and scratches created on the enamel surface during mastication, is one such line of evidence.

Differences in the dental microwear of grazing and browsing ungulates have been noted in previous studies of extant and fossil species (e.g. Solounias et al., 1988; Solounias and Moelleken, 1993; Merceron and Ungar, 2005; Merceron et al., 2005; Schubert et al., 2006; Ungar et al., 2007), with browsers having more pits and grazers having more scratches. Mixedfeeding taxa tend to have intermediate microwear signatures, or patterns overlapping with both browsers and grazers. This has important implications for understanding the relationship between bovid diet and habitat preference, as while the connection between diet and habitat are not exact, grazing and browsing microwear signatures do imply the availability of grasses and woody plants respectively.

Bovids are generally considered to be reliable indicator species for environmental reconstruction because modern ruminants fall into discrete dietary categories that reflect habitat preferences. The relationship between bovid diet and habitat has been long established in the literature (e.g., Vrba, 1980, 1985; Wing et al., 1992, Bobe and Eck, 2001) and their fossils are frequently used to infer the ecological context of a site (e.g. Gentry, 1970; Solounias and Dawson-Saunders, 1988; Merceron et al., 2004a, 2005; Merceron and Ungar, 2005; Ungar et al., 2007). This is especially true for early hominin localities where fossil bovids tend to be among the most commonly recovered large mammals (e.g. Vrba, 1980, 1995; Kappelman, 1984; Shipman and Harris, 1988; Harris, 1991; Plummer and Bishop, 1994; Kappelman et al., 1997; Reed, 1997, 2008; Spencer, 1997; Sponheimer et al., 1999, 2003; Schoeninger et al., 2003; Schubert et al., 2006; Kingston and Harrison, 2007).

Most models for paleoenvironmental reconstruction using bovids apply the principle of taxonomic uniformitarianism, the assumption that fossil species share the same ecological preferences as their closest living relatives. Although this assumption is rarely questioned for recent faunas, its utility in reconstructing the paleoecology of long extinct species is uncertain. One of the principal issues with taxonomic uniformitarianism is that it assumes groups remain constant in their ecological preferences over long periods of time. While taxonomic uniformitarianism is relatively straightforward and requires minimal effort beyond a glance at a faunal list, critics have questioned many of its inherent assumptions (Solounias et al., 1988; Reed, 1996; Sponheimer et al., 1999; Schubert et al., 2006). In a study of fossil bovid diets utilizing stable isotopes and ecomorphology, Sponheimer et al. (1999) demonstrated that half of the included taxa had diets that differed from the assumed diets based on taxonomic uniformitarianism. For example, although Aepyceros sp. and Gazella vanhoepeni had been previously assumed to be mixed/seasonal feeders like their closest living relatives, the isotopic signatures and ecomorphological data suggested that both were obligate browsers, and revealed no evidence of C4 grass consumption. Due to these discrepancies between dietary signals, it is critical that ecological similarities of extant and fossil taxa be tested in some reliable way in order to have confidence in these assumed relationships.

### **Materials and Methods**

#### Specimens

575 specimens were included in this study, representing 25 extant bovid taxa. The taxa used in this study were selected because they represent a wide variety of ecological adaptations, with diets ranging from open-country-adapted obligate grazers to closed-habitat browsers to

dedicated frugivores. Special emphasis was placed on including taxa that engage in various levels of mixed feeding, including variable grazers, browser-grazer intermediates and generalists. Finally, most of the included taxa have well-understood diet and habitat preferences that have been documented during field studies. Table 1 lists the included extant taxa and summarizes their dietary preferences.

The specimens examined for this study are housed at the American Museum of Natural History (AMNH), New York; the Field Museum (FMNH), Chicago; the Smithsonian National Museum of Natural History (NMNH), Washington D.C.; and the Royal Museum of Central Africa (RMCA), Tervuren, Belgium. All are wild-shot specimens with known provenience data. Only specimens that preserved unobscured antemortem microwear were included in the dataset. The criteria used to determine suitability for microwear analysis were those of Teaford (1988) and King et al. (1999).

Both upper and lower second molars were included in this study and have been shown in previous projects to yield similar results (Merceron et al., 2004a, 2004b). High-resolution casts were prepared following conventional procedures for microwear analysis (Grine, 1986, Ungar, 1996). The original specimens were cleaned with cotton swabs soaked in acetone and then molded using a polyvinylsiloxane dental impression material (Presidents Jet Regular Body, Coltene-Whaledent Corp.). Casts were then made using high-resolution epoxy (Epotek), which has been demonstrated to produce replicas that are precise to a fraction of a micron (Beynon, 1987; Teaford and Oyen, 1989).

## **Dietary Classifications**

The dietary categories used in this study are those of Gagnon and Chew (2000) and expand on the traditional grazer-browser-mixed feeder continuum. This classification system was selected because it recognizes various levels of mixed feeding and allows for categorization of all African bovid taxa. The categories are determined by the percentages of fruits, dicotyledons and monocotyledons included in the diet, as recorded during observations in the wild. Inarguably, consideration of these various levels of mixed feeding improves the resolution of dietary reconstructions and can aid in our understanding of behavior and habitat preference in fossil taxa. However, it is also important to note that extensive ecological observation and fecal analyses have not been conducted for many taxa classified as mixed feeders (Gagnon and Chew, 2000, Fortelius and Solounias, 2002). Further field work is needed to better determine the influence of seasonal and geographic variation on the diets of these mixed feeding bovids.

The species included in the study were classified using the Gagnon and Chew model (2000) and are listed in Table 1. Taxa classified as obligate grazers have a diet consisting of more than 90% monocotyledons, with no seasonal or geographic variation. Variable grazers consume 60-90% monocotyledons, but vary in diet seasonally and geographically. The diet of bovids classified as browser-grazer intermediates consists of 30-70% monocotyledons and dicotyledons and always includes some fruit, although never more than 20%. Browsers have a diet that includes more than 70% dicotyledons. Fruit accounts for more than 70% of the diet of bovids classified as frugivorous and these taxa rarely, if ever, consume monocotyledons. Taxa classified as dietary generalists are widely variable by season and locality, but always consume more than 20% of each of the three food types.

## **Data Collection**

Conventional methods of studying dental microwear, including scanning electron microscopy and light microscopy are subject to observer measurement error and a lack of depth data due to two dimensional surface characterizations (Grine et al., 2002). Dental microwear texture analysis was developed to be a solution to these problems (Ungar et al., 2003). It combines white-light confocal profilometry with scale-sensitive fractal analysis, resulting in the quantitative description of surfaces at a range of scales. This offers a suite of objective and repeatable measures of microwear surface textures in three dimensions that can be compared between groups with differing diets. Further, white-light confocal profilometry is quicker, easier to use, and can be less costly than scanning electron microscopy; and automated analysis requires less time and effort from researchers than does identification and measurement of individual features. The most important advance of texture analysis is the reduction of observer error in measurement, which allows direct comparisons between studies, as well as the establishment of a large database that researchers can access for interpretation of their results. Dental microwear texture analysis has been successfully used to identify dietary differences between species of extant and fossil taxa including humans and non-human primates, marsupials, carnivores, and bovids (R. Scott et al., 2005, 2006; Ungar et al., 2008a, 2008b, 2010, in press; Krueger and Ungar, 2009; Merceron et al., 2009; Prideaux et al., 2009; J. Scott et al., 2009; Ungar and Scott, 2009; Schubert et al., 2010; Schultz et al., 2010) and to detect differences in diet by season and even gender (Merceron et al., 2010).

As is conventional for studies of bovid dental microwear, texture data were collected on the disto-buccal enamel band of the mesial cuspid of  $M_2$  and the mesio-buccal enamel band of the mesial cusp of  $M^2$  (Janis, 1990; Merceron, 2005; Schubert, 2006; Ungar et al., 2007). These

areas are illustrated in Figure 1. The casts were scanned using a Sensofar PLµ white-light confocal profiler (Solarius Development Inc., Sunnyvale, CA). This instrument collects 3-D point clouds from the wear surface of the tooth to create photo simulations and digital elevation models.

Each specimen was scanned using a 100x objective, resulting in the generation of a point cloud with a lateral sampling interval of 0.18  $\mu$ m, a vertical resolution of 0.005  $\mu$ m, and a field of view of 102 x 138  $\mu$ m. Data were collected for four adjoining fields, resulting in a total work envelope of 204 x 276  $\mu$ m for each specimen. The scans were then leveled using Solarmap Universal software (Solarius Development Inc., Sunnyvale, CA). Prior to analysis, defects such as dust or other adherents were removed from the scan using the erase or thresholding function in Solarmap and/or the slope-filtering function in Sfrax (Surfract Corporation). The resulting data files were saved in ".sur" format in preparation for analysis.

#### **Scale-Sensitive Fractal Analysis**

Point clouds were analyzed using Toothfrax and Sfrax scale-sensitive fractal analysis software packages (Surfract Corporation). Scale-sensitive fractal analysis originates from studies of fractal geometry, and is based on the principle that the texture of a surface changes with the scale of observation. Thus, a surface that appears smooth at a coarse scale may seem rough when viewed at a fine scale. These apparent changes in surface texture across scales can be examined for entire surfaces or for profiles across a surface. Several texture variables generated during these analyses have been identified as particularly informative for dental microwear studies (Ungar et al., 2003; Scott et al., 2005, 2006). Data for five of these variables are presented here. They are area-scale fractal complexity, length-scale anisotropy of relief, scale of maximum
complexity, textural fill volume and heterogeneity of area-scale fractal complexity. These variables have been described in detail elsewhere (Ungar et al., 2003, 2007, 2008; Scott et al., 2005, 2006) and will be summarized briefly here.

<u>Complexity (Asfc)</u> - Area-scale fractal complexity is measured as the change in surface roughness across different scales of observation. *Asfc* is calculated by taking the slope of the steepest part of a curve fit to a plot of relative area over the range of scales at which the measurements are taken. The value for *Asfc* will be higher for steeper slopes, indicating more complex surfaces. For example, a surface dominated by pits of various sizes or pits and scratches overlaying one another will result in high values for *Asfc*.

<u>Scale of Maximum Complexity (Smc)</u>- Earlier studies using dental microwear texture analysis have suggested that range of scales at which *Asfc* is calculated may be informative (Scott et al., 2005, 2006). This is calculated by taking the range of scales where the relative area versus scale curve is the steepest. *Smc* likely corresponds in some measure with the scale of wear-causing particles. Values for this variable will be highest for surfaces that display less wear at finer scales and will be lowest for those with more wear at fine scales.

<u>Anisotropy (epLsar)</u>- Length-scale anisotropy of relief is a measure of the orientation concentration of surface roughness. This variable is calculated by taking profiles of the wear surface at different orientations at a given scale. This study employed a sampling interval of 5° and a scale of 1.8  $\mu$ m. Highly anisotropic surfaces have relative profile lengths that differ with orientation. The relative lengths at given orientations can be defined as vectors and normalized to create a rosette diagram. Values for *epLsar* are calculated by taking the length of the mean vector. Surfaces that feature parallel scratches will have a high value for *epLsar*.

<u>Textural Fill Volume (Tfv)</u>- The textural fill volume algorithm calculates the volume of square cuboids at a given scale that can fill a surface. *Tfv* is defined as the difference in summed volume for fine cuboids (2  $\mu$ m per side) and large cuboids (10  $\mu$ m per side). This removes the impact of overall surface structure, such as facet curvature, and allows for the characterization of the microwear features themselves. Surfaces dominated by large pits or deep scratches will have high values for *Tfv*.

<u>Heterogeneity (HAsfc)</u> - The variables mentioned above all help to provide accurate descriptions of microwear textures; however, the values for adjoining areas of the facet may vary. This variation for a given variable across a surface may be important in characterizing the surface. In order to better study this variation, the heterogeneity of area-scale fractal complexity variable measures the difference in *Asfc* across a surface. *HAsfc* is calculated through the use of the Auto-Split function in Toothfrax. This computation divides the scanned area of the facet into successively smaller sub-regions with equal numbers of rows and columns, beginning with 2 x 2 and ending with 11 x 11. The value for *HAsfc* is defined as the median absolute deviation of *Asfc* divided by the median of *Asfc*.

#### **Statistical Analysis**

Following analysis, median values were calculated for the four scans representing each specimen (Scott et al., 2006). The resulting data were then rank transformed prior to analysis because unranked microwear texture data typically violate assumptions inherent in parametric statistical tests (Conover and Iman, 1981). A nested analysis of variance model was used in this study, with taxon as the subordinate level of classification, nested within assigned dietary category, specifically obligate grazer, variable grazer, browser-grazer intermediate, generalist,

browser, or frugivore (Gagnon and Chew, 2000). The data were then analyzed using a general linear model to determine whether there were overall differences in microwear among the six dietary categories. Analyses of variance for individual texture attributes and pairwise comparisons tests were then used to determine the sources of significant variation both between dietary categories and among taxa within the dietary categories. Both Tukey's Honestly Significant Difference and Fisher's Least Significant Difference tests were used to balance the risks of Type I and Type II errors (Cook and Farewell, 1996).

# Results

Examples of the microwear surfaces of extant bovid taxa are illustrated in Figures 2 and 3. Descriptive and analytical statistics are presented in Tables 2, 3 and 4 and Figures 4 and 5. There is significant variation in the overall general linear estimate model, indicating that the microwear textures varied among the dietary classifications in all six variables. Additionally, some significant variation was also identified among taxa within the dietary categories.

#### **Results by variable**

<u>Asfc</u> – Tukey's HSD pairwise comparisons indicate significant differences in surface complexity among all pairs of dietary categories except browser-grazer intermediates versus generalists. These tests indicate that frugivores had significantly higher microwear complexity values than all other dietary categories. Predictably, browsers had higher complexityvalues than the grazing and mixed feeding taxa. Browser-grazer intermediates and generalists had similar values for complexity and were not significantly different using Tukey's HSD or Fisher's LSD

tests. Of the grazing taxa, variable grazers had significantly higher complexity values than obligate grazers.

<u>epLsar</u> – Significant differences were found among all taxa except browser-grazer intermediates versus generalists, and browsers versus frugivores. Fisher's LSD test also indicated significant differences between browser-grazer intermediates versus generalists. Obligate grazers had the highest values for anisotropy, while frugivores and browsers had the lowest. Among the mixed feeding taxa, browser-grazer intermediates had higher values than generalists and variable grazers.

<u>Smc</u> - Tukey's HSD pairwise comparisons indicate significant differences among all dietary categories, with the exception of browsers versus variable grazers, and frugivores versus generalists, which were distinguished by Fisher's LSD test. Obligate grazers had the highest values for *Smc*, while frugivores had the lowest. Variable grazers had the highest values among mixed feeding taxa, with browser-grazer intermediates and generalists having comparatively low values.

<u>*Tfv*</u>- Significant differences in textural fill volume were found between all pairs of dietary categories when analyzed with Tukey's HSD pairwise comparisons test. Frugivores had the highest values for textural fill volume, followed by browsers. Among the mixed feeding taxa, generalists had higher values for fill volume than browser-grazer intermediates. Obligate grazers had the lowest recorded values.

<u>*HAsfc9*</u>- Tukey's HSD pairwise comparisons indicate significant differences between all pairs of dietary categories except frugivores versus browser-grazer intermediates, and generalists versus variable grazers. Fisher's LSD test also reported significant differences for browser-grazer intermediates versus frugivores. Browsers had the highest values for heterogeneity, while

obligate grazers have the lowest values. Among mixed feeders, browser-grazer intermediates and frugivores have higher values than variable grazers and generalists.

<u>HAsfc<sub>81</sub></u>– Tukey's HSD pairwise comparisons indicate significant variation in heterogeneity between all pairs of dietary categories except browser-grazer intermediates versus frugivores, browser-grazer intermediates versus generalists, frugivores versus generalists, and obligate grazers versus variable grazers. A Fisher's LSD test also indicated significant differences for browser-grazer intermediates versus generalists and obligate grazers versus variable grazers. Browsers have the highest values forheterogeneityand obligate and variable grazers have the lowest values.

#### **Results by diet category**

The trends by dietary category are summarized in Table 5. In general, frugivores have high values for complexity and fill volume, low values for anisotropy and mid-range values for heterogeneity and the scale of maximum complexity. Browsers have high values for complexity, fill volume and heterogeneity, low values for anisotropy and mid-range values for scale of maximum complexity. Obligate grazers have high values for anisotropy and the scale of maximum complexity, and low values for complexity, fill volume and heterogeneity. Variable grazers have low complexity and heterogeneity, and mid-range values for anisotropy, fill volume and the scale of maximum complexity. Browser-grazer intermediates have relatively low values for anisotropy and mid-range values for complexity, the scale of maximum complexity, fill volume and heterogeneity. Generalists have high values for fill volume, low values for anisotropy and the scale of maximum complexity, and mid-range values for complexity and heterogeneity.

#### Variation within dietary category

In addition to clear differences among the dietary categories, some significant variation among taxa within dietary category was also indicated. Significant differences among obligate grazers, variable grazers, browser-grazer intermediates and browsers were present for all texture attributes. There was significant variation among generalists in *epLsar* and among frugivores in *epLsar*, *Smc*, *Tfv*, *HAsfc*<sub>9</sub>.

<u>Obligate grazers</u>- Among obligate grazing taxa, *Redunca fulvorufula*has significantly higher values for complexityand significantly lower values for the scale of maximum complexitythan any other species. *Damaliscus lunatus* and *Redunca fulvorufula*both have significantly higher values for anisotropy and heterogeneity 3 x 3, as well assignificantly lower values for fill volumeand heterogeneity 9 x 9than *Kobus leche*, and *Redunca fulvorufula*also has higher values for this variable than *Redunca arundinum*.

Variable grazers- Significant differences were also present among variable grazers in all texture attributes. Tragelaphus spekii and Gazella granti have significantly higher values for complexity than the other four taxa. Gazella granti, Damaliscus pygargus and Tragelaphus spekii have significantly lower values for anisotropythan the other variable grazing taxa. Gazella granti and Hippotragus nigerboth have significantly lower values for the scale of maximum complexitythan the other four taxa, and Damaliscus pygargus has significantly higher values for this variable than Gazella granti, Hippotragus niger, Kobus ellipsiprymnus and Syncerus caffer. Damaliscus pygargus and Hippotragus niger have significantly higher values for fill volumethan the other four taxa, none of which differ significantly from one another. Hippotragus niger and Kobus ellipsiprymnus group together in heterogeneity 3 x 3and have significantly higher values

than any of the other four taxa. *Damaliscus pygargus* and *Gazella granti* have significantly lower values than *Hippotragus niger*, *Kobus ellipsiprymnus*, and *Syncerus caffer*. *Damaliscus pygargus* and *Gazella granti* have significantly lower values for heterogeneity 9 x 9 than *Kobus ellipsiprymnus* and *Tragelaphus spekii*.

<u>Browsers</u>- Browsing taxa also showed some significant differences among taxa in all variables. *Tragelaphus euryceros* and *Neotragus batesi* have significantly higher complexityvalues than *Sylvicapra grimmia*, *Litocranius walleri*. *Sylvicapra grimmia* has significantly lower values for complexity than any of the other browsers. *Tragelaphus euryceros* has significantly lower anisotropy values than any of the other browsing taxa, while *Sylvicapra grimmia* has significantly higher values for this variable than *Neotragus batesi* or *Tragelaphus euryceros*. *Tragelaphus euryceros* has significantly higher fill volumethan *Neotragus batesi*. *Litocranius walleri* has significantly higher scale of maximum complexity values than *Sylvicapra grimmia*, as well as higher values for both heterogeneity variables than any of the other browsers.

Browser-grazer intermediates- Among browser-grazer intermediates, Aepyceros melampus and Tragelaphus imberbis have significantly lower complexityvalues than the other four taxa, while Raphicerus campestris has significantly lower anisotropyvalues. Tragelaphus imberbis and Antidorcas marsupialis have significantly lower values for the scale of maximum complexity. Raphicerus sharpei has significantly lower fill volumevalues than all five other taxa in this category, while Raphicerus campestris has significantly higher values for this variable than Aepyceros melampus, Antidorcas marsupialis, and Raphicerus sharpei. Finally, Raphicerus sharpei has significantly higher values for both heterogeneity variables than all of the other browser-grazer intermediates.

<u>Generalists and frugivores</u>- There was generally less significant variation among frugivores and generalists than the more dedicated grazing and browsing taxa. Of the generalist taxa, the only significant difference was in anisotropy and all three taxa were significantly different from one another, with *Tragelaphus strepsiceros* having the highest values and *Oreotragus oreotragus* having the lowest. Frugivores differed in anisotropy, the scale of maximum complexity, fill volume and heterogeneity 3 x 3, with *Cephalophus sylvicultor* having significantly higher values for anisotropy and the scale of maximum complexity and significantly lower values for fill volumeand heterogeneity 3 x 3.

#### **Discussion and Conclusion**

The dataset presented here allowed me to test the hypothesis that dental microwear texture analysis can distinguish between taxa that fall into different dietary categories, particularly those that engage in various levels of mixed feeding. By including large sample sizes, by microwear standards, and multiple taxa representing each dietary category, testing the ability of dental microwear texture analysis to differentiate taxa could be done with much more confidence. The results of the analysis confirm that the technique is capable of finer dietary distinctions than previously demonstrated and suggest that, with additional testing, more specific information still may be gleaned. These results have important implications for the use of dental microwear textures to distinguish bovid taxa with different dietary categories and to reconstruct the diets, and by extension habitats, of fossil bovids.

**Bovid diets and dental microwear analysis** – As previous studies have suggested, the classic dietary categories used to describe bovids (grazers, browsers, mixed feeders) are obvious

oversimplifications (Bodmer, 1990; Gordon and Illius, 1994; Gagnon and Chew, 2000). Even so, these classifications do have substantive heuristic value. In addition to differing in their nutritional content, grasses and browse also differ in their physical properties (Shipley, 1999), which are essentially what is being measured by dental microwear analysis. Monocotyledonous grasses are distinguished by dense cell walls composed primarily of cellulose (Demment and Vansoest, 1985) and veins that typically run parallel to the long axes of their blades. This structure causes the grass blades to be highly resistant to stress and difficult to fracture (Vincent, 1990). It has also been long hypothesized that the tissues found in grasses are denser and more homogenous than those found in browse, thus resulting in a difference in the overall fracture pattern between the two food types (Spalinger et al., 1986; Kay, 1993; Van Wieren, 1996). Additionally, grasses tend to have high concentrations of abrasives, such as endogenous phytoliths, and/or exogenous grit adhering to the surfaces of individual blades (Baker et al., 1959; Dougall et al., 1964; NcNaughton et al., 1985; Sanson et al., 2007). While they cannot fuel the body, these abrasives are harder than dental enamel and contribute to the microwear signature of grazing taxa by scratching the surface of the teeth during mastication (Baker, 1959).

Dicotyledonous browse can include a wide variety of foods, making it difficult to characterize. In general, however, items classified as browse have thinner cell walls resulting in greater ease of access to nutrient content than grasses (Spalinger et al., 1986). Browse components are also more variable than grasses in their physical properties (Caswell et al., 1973). Some browse resources tend to be more brittle and therefore less resistant to fracture than grass blades. Others, however, are much harder and require greater stress to fracture, possibly due to the higher percentages of bundle sheaths characteristic of C3 browse (Caswel et al., 1973; Heckathorn et al., 1999).

These contrasts between hard and tough foods have important implications for interpreting patterns of dental microwear because they reflect the biomechanical adaptations needed to process foods with different fracture properties (Lucas, 2004). Tough foods, such as grass blades, tend to require shearing or grinding, with the lateral movement of opposing occlusal surfaces relative to one another. This process results in abrasives being dragged across the surface of the teeth, causing microscopic striations on the enamel (Rensberger, 1973; Walker et al., 1978). Parallel tooth-food-tooth processing can also cause 'prism plucking', or small prism-sized pits on the enamel surface (Walker, 1984; Teaford and Runestad, 1992).

Hard and/or brittle foods, on the other hand, necessitate crushing, or vertical contact of opposing surfaces with food items fractured between them, a process that results in the production of microscopic pits. Because of the range of food types and fracture properties in items categorized as browse, the resulting pits can vary widely in size and shape. Larger pits, for example, are thought to result from concentrated pressure on hard foods between the enamel surfaces, whereas smaller pits may be caused by strict tooth-tooth wear (Walker, 1984; Teaford and Oyen, 1989a, b; Teaford and Runestad, 1992)

The ability of dental microwear analysis to distinguish grazing and browsing bovids is well documented in the literature. The earliest studies of bovid dental microwear were conducted using a scanning electron microscope (e.g. Solounias et al., 1988; Solounias and Moelleken, 1992a, 1992b, 1993). Microwear surfaces were imaged at 500x and individual features were digitized and measured. These studies demonstrated that grazing taxa tend to have occlusal surfaces dominated by scratches, while browsers had more pits. Mixed feeding bovids usually show intermediate percentages of pits and scratches.

More recent studies have approached the characterization of bovid dental microwear using feature-based or texture-based analysis techniques to identify finer subdivisions within the basic grazer-browser-mixed feeder categories. Low-magnification light microscopy, a featurebased microwear technique, requires the examination of wear facets using a standard stereomicroscope at 35x. Microwear features are then quantified and divided into six categories: scratches, pits, scratch textures, cross scratches, large pits and gouges (e.g. Solounias and Semprebon, 2002; Rivals and Deniaux, 2003; Semprebon et al., 2004). Studies employing this method have reported clear differences between dedicated grazing and browsing taxa, and have described two distinct variations in the microwear signatures of mixed feeders- seasonal/regional mixed feeders and meal-by-meal mixed feeders. Other studies have applies combinations of various techniques to successfully distinguish between grazing and browsing bovids, including low magnification light microscopy with semi-automated image analysis (Merceron et al., 2004a, b, 2005, 2007) and white-light confocal microscopy with feature-based measurement techniques has also been applied to the study of bovid dental microwear (Merceron and Ungar, 2005; Schubert et al., 2006).

Most recently, texture-based analysis, which utilizes three-dimensional characterization of the microwear surface and semi-automated image analysis, has been employed in the study of dental microwear. While to date there have been only limited applications of this technique to bovids, the results have clearly demonstrated the effectiveness of texture analysis in distinguishing grazing, browsing and mixed feeding taxa and the technique shows promise for more specific dietary distinction. Ungar and colleagues (2007) have suggested that dental microwear texture analysis has the potential to differentiate among finer dietary categories and called for further study of extant bovids with known subtle differences in diet.

A new database of bovid dental microwear textures- The extant African bovids included in this study show a wide variety of microwear texture patterns that reflect a correspondingly broad range of known diets. The results presented here confirm the previous observation of Ungar and colleagues (2007) that bovids classified as browsers have complex microwear surfaces with low anisotropy, whereas those classified as obligate grazers have highly anisotropic surfaces, with low complexity. In fact, there is no overlap between obligate grazers and browsers in an anisotropy-complexity bivariate space, as illustrated in Figure 4. In addition to supporting the previous work, this study adds data for four additional variables: *Smc*, *Tfv*, *HAsfc*<sub>9</sub>, and *HAsfc*<sub>81</sub>, to that previously described by Ungar et al. (2007). The present work also greatly expands sample sizes, and the number and dietary range of extant bovid taxa available for the baseline to interpret wear signatures of fossil taxa.

Variation among dietary categories- Diet clearly explains a significant fraction of the overall variation in the microwear patterns, with browsers having higher values for complexity, fill volume, and both heterogeneity variables and grazers having higher values for anisotropy and scale of maximum complexity. In fact, when taxa were grouped by dietary category and compared, significant differences were present for all six variables. The only consistent overlap in microwear signature was between taxa classified as browser-grazer intermediates and those classified as generalists.

Beginning with the extremes of the classic browser-grazer continuum, significant differences were present between browsers and obligate grazers for all six variables and in fact, there was no overlap recorded between the ranges of these taxa for any of the texture attributes

(Figure 4). The difference between these two dietary categories is clearly visible when viewed in a bivariate space contrasting complexityand anisotropy (Figure 5), with browsers having higher values for *Asfc* and lower values for *epLsar* and grazers falling at the opposite end of both distributions. Taxa that have been classically categorized as mixed feeders, including browser-grazer intermediates and generalists, fall in between these two extremes.

One of the primary goals of this study was to determine whether or not dental microwear texture analysis could differentiate between various levels of grazing, browsing and mixed feeding, as previously suggested by Ungar et al. (2007). In addressing this question, comparisons between obligate grazers and variable grazers; browsers and frugivores; and browser-grazer intermediates and generalists were of particular import. Beginning with the grazing taxa, some overlap was documented in the ranges of obligate and variable grazers; however, there were significant differences between the two groups for all texture attributes with the exception of  $HASFC_{81}$ . Bovids classified as obligate grazers have more limited ranges than variable grazers, particularly for complexity and anisotropy. This lack of variation in the microwear signature of obligate grazers is unsurprising. These bovids have diets that consist of >90% monocots and are not variable by season or region. Although monocots also make up the largest component of the diets of variable grazers, these taxa consume more dicots and/or fruit than obligate grazers. The diets of variable grazers also change as herds move from one region to another and can also be altered seasonally. The wider value ranges for texture attributes like complexity, anisotropy and fill volume is likely reflective of this seasonal and/or regional shift in food types evinced by variable grazers.

Browser-grazer intermediates and generalists had more overlap in microwear signature than any of the other groups, with no significant differences in complexity or anisotropyand only suggestive differences in scale of maximum complexity or heterogeneity. This lack of variation is not surprising given that both groups regularly consume a combination of hard and tough foods in somewhat equal parts. Browser-grazer intermediates typically consume 30-70% monocots and dicots and slightly less than 20% fruit, while generalists regularly consume >20% of all food types. Even given these dietary similarities, some differences were present. Significant differences in fill volume were found between the two groups, although some overlap was present in the ranges. This may be reflective of differences in hard fruit consumption by the taxa included in this study. Fruit makes up >20% of the diets of all three species classified as generalists, whereas it only makes up 10% or less of the diets of all six browser-grazer intermediates. Since frugivores have the highest values for fill volume recorded in this study, it is possible that higher inclusions of this food type may explain why this is the only significant difference between browser-grazer intermediate and generalist taxa.

It is interesting to note that non-fruit browsers and frugivores, often lumped together into a single dietary classification, show significant differences in microwear signature for all variables, with the exception of anisotropy. In general, the values for the frugivorous bovids were higher than those of the browsing taxa, suggesting that regular fruit consumption by bovids results in even more complex microwear surfaces with larger and deeper features than those resulting from the consumption of other dicot plant parts. Fruit, as a dietary category, includes fruits, bulbs, tubers, succulents, pods, and seeds. Many of these food types are protected by tough or hard shells that require considerable processing to open and would result in increased pitting on the enamel surface. For example, fruit is frequently protected by a tough exocarp and seeds are often surrounded by a thick, hard endocarp. Frugivorous bovids also display lower degrees of seasonal and/or geographic variation than browsing taxa. For example, both frugivores included in this study have documented diets consisting of >75% fruit with no recorded seasonal variation (Kingdon, 1982, Perrin et al., 1992), while browsers like *Sylvicapra grimmia* and *Tragelaphus euryceros* include >75% other dicots and vary by season. The distinct differences in microwear textures recorded here between frugivorous and browsing bovids supports previous arguments by Bodmer (1990) and Gagnon and Chew (2000) that frugivory should be recognized as a distinct dietary category for bovids, as it is for primates, and that bovid diets should be considered as a grazer-frugivore continuum rather than a grazer-browser continuum.

Variation within dietary categories- In addition to considering the higher-order factor "dietary classification", variation among taxa nested within dietary categories was also explored. Some significant differences were found among species classified as obligate grazers, variables grazers, browsers and browser-grazer intermediates.

Among obligate grazing taxa, *Redunca fulvorufula*differed significantly from the other three taxa in with higher values for complexity and scale of maximum complexity, as well as lower values for anisotropy. These differences can be best explained by differences in fruit consumption. This species has been reported to include fruit as nearly 5% of its regular diet, while *Damaliscus lunatus*, *Kobus leche*, and *Redunca arundinum* consume little to no fruit (Hofmann and Stewart, 1972). This addition of hard fruit to a diet of primary grazing may be responsible for the significant increases in microwear surface complexity and feature size, as well as the decrease in feature anisotropy, when compared to the other taxa.

Significant differences were also found among variable grazers in all texture attributes. *Tragelaphus spekii*, *Gazella granti*, and *Syncerus caffer* generally group together away from the other taxa, with higher values for complexity and lower values for anisotropy. This distinction may be related to increased dicot consumption by these species. Dicots make up 20-30% of the diet of these three taxa, whereas they only account for <10% of the diet in *Damaliscus pygargus*, *Kobus ellipsiprymnus* or *Hippotragus niger* (Skinner and Smithers, 1990; Gagnon and Chew, 2000).

Browsing taxa also showed some significant differences among taxa in anisotropy, scale of maximum complexity and heterogeneity 9 x 9. *Tragelaphus euryceros* has significantly lower anisotropy values than *Litocranius walleri*, *Neotragus batesi* or *Sylvicapra grimmia*. *Tragelaphus euryceros* consumes slightly more monocots than the other three taxa and this could explain the lower values for anisotropy. Additionally, *Sylvicapra grimmia* has significantly lower values for scale of maximum complexity and heterogeneity 9 x 9 than the other three taxa. This could be reflective of the somewhat more catholic diet reported for the species. Dicots make up >83% of the diet of *Sylvicapra grimmia*, with a combination of fruits and monocots comprising the rest. This is in contrast to *Neotragus batesi* and *Tragelaphus euryceros*, the diets of which consist of only 75% monocots.

Significant differences among taxa were also recorded for browser-grazer intermediates in anisotropy and fill volume. *Raphicerus campestris* has significantly lower values for anisotropy and higher values for fill volume than the other browser-grazer intermediates. This may reflect the fact that *Raphicerus campestris* has been reported to consume higher percentages of fruits and dicots that most of the other browser-grazer intermediates. Combined, these two food types comprise >65% of the diet for *Raphicerus campestris*, while accounting for

approximately 55% in *Antidorcas marsupialis*, *Raphicerus sharpei* and *Tragelaphus imberbis* and only 50% in *Taurotragus oryx* and *Aepyceros melampus* (Gagnon and Chew, 2000). Interestingly, this supports previous work by Sponheimer et al. (2003), which suggested that *Raphicerus campestris* included considerably less grass in its diets than previously thought.

The differences between taxa within dietary categories were not always as predicted but seem to have some biological relevance when considered alongside known differences in diet. All of this suggests that dental microwear texture analysis has the potential to distinguish between taxa with more subtle differences in diet and the within category differences reported here will certainly be a source of hypotheses for further testing. While this obviously requires further investigation, the possibility of being able to identify these subtle dietary distinctions in the fossil record is a tantalizing one. If further study confirms that the technique can, in fact, identify dietary differences to this level, the application of dental microwear texture analysis to the study extinct bovids could strongly enhance what we know about the niche differentiation and habitat preference of fossil taxa.

**Implications for habitat reconstruction**- Because many of the extant bovids included in this study utilize multiple habitat types seasonally and/or depending on geographic range, comparisons of taxa by habitat type were not practical. However, the results do have implications for the study of bovid habitat preference that can be cautiously applied to the fossil record. Obviously, the connection between diet and habitat are not exact, but grazing and browsing microwear signatures do imply the availability of grasses and woody plants respectively (Schubert et al., 2006). The obligate grazing taxa, for example, all occupy some type of grassland. *Redunca arundinum* lives in both moist lowland savanna grassland and short

shrubland (Kingdon, 1982; Skinner and Smithers, 1997), *Redunca fulvorufula*occupies grassy mountain ridges (Hofmann and Stewart, 1972; Kingdon, 1997; Skinner and Smithers, 1990), *Damaliscus lunatus* is found in both open grassland and floodplain (Kingdon, 1982; Skinner and Smithers, 1997), and *Kobus leche* (Kingdon, 1982; Sayer and Van Lavieren, 1975) typically inhabits wet grassy marshland. Dedicated browsing taxa, on the other hand, occupy forest or shrubland. *Litocranius walleri* inhabits both bushland and woodland (Kingdon, 1982; Leuthold, 1978), *Sylvicapra grimmia* lives in savanna and bushland (Kingdon, 1982; Skinner and Smithers, 1990), while *Neotragus batesi* and *Tragelaphus euryceros* are both typically found in closed forest habitats. Both frugivorous species included here inhabit closed forests (Hofmann and Stewart, 1972; Ralls, 1978; Kingdon, 1982; Emmons et al., 1983).

The relationship between diet and habitat is even less exact for mixed feeding taxa that consume a variety of food types. Even so, dietary reconstruction using microwear analysis can still inform habitat preference. All of the variable grazers included in this study occupy some type of primary or peripheral grassland, with the exception of *Hippotragus niger*, which is usually associated with open woodland habitats (Kingdon, 1992). The taxa classified as browser-grazer intermediates are all found in forest or bushland habitats, except *Aepyceros melampus*, sometimes found in savannas (Skinner and Smithers, 1990; Kingdon, 1992), and *Antidorcas marsupialis*, which occupy grasslands (Hofmann and Stewart, 1972; Kingdon, 1982). The three generalist taxa occupy a combination of bushland, open woodland and closed forest habitats (Dunbar and Dunbar, 1974; Kingdon, 1982, 1992). In sum, while there is no one-to-one relationship between dental microwear and habitat, the ability to accurately classify diet, and especially to classify beyond the traditional grazer-browser-mixed feeder continuum, can help narrow down habitat preferences to a likely range of local environments.

**Summary and concluding remarks**- This study builds on previous work by others, and features 25 extant African bovid taxa from six known dietary categories. Analysis nesting species within dietary classifications proposed by Gagnon and Chew (2000) was employed in order to better understand differences in microwear signatures resulting from the consumption of different combinations of monocots, dicot plant parts and fruits. The results indicate that dental microwear texture analysis can successfully distinguish between bovid taxa with differing diets, including those that reflect different levels of mixed feeding. Additionally, the fact that nearly all dietary categories were significantly different from one another supports the accuracy of the dietary classification system proposed by Gagnon and Chew (2000).

The dataset described here can be used as a baseline for interpreting microwear in fossil bovids from a variety of sites and time periods across Africa. The clear separation between dietary categories indicates that this comparative database will be useful in reconstructing diet and possibly even habitat preference. Significant differences among species within dietary categories confirm the previous assertion by Ungar et al. (2007) that finer dietary distinctions between taxa may be possible with texture analysis. Continued work on extant ruminants and the inclusion of taxa with even finer distinctions in diet and habitat preference will be necessary to assess the ultimate potential resolution of dental microwear texture analysis.

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#### **Figure Legends**

- Figure 1. Occlusal view of bovid M2, illustrating the shearing facets where microwear was examined (Illustration courtesy of Gildas Merceron).  $A = left m^2$ ,  $B = right m_2$
- Figure 2.Examples of dental microwear surfaces for extant bovids from different dietary categories. These images are photosimulations based on data collected using the confocal imaging profiler, and each represents an area of 276 x 204 μm. A) obligate grazer; B) variable grazer; C) browser-grazer intermediate; D) generalist; E) browser; and F) frugivore.
- Figure 3.Examples of dental microwear surfaces for extant bovids from different dietary categories. These images are 3-D meshed axonometrics based on data collected using the confocal imaging profiler, and each represents an area of 138 x 102 μm. A) obligate grazer; B) variable grazer; C) browser-grazer intermediate; D) generalist; E) browser; and F) frugivore.
- Figure 4.Box and whisker plots comparing extant taxa. A) mean complexity (*Asfc*); mean anisotropy (*epLsar*); C) mean scale of maximum complexity (*Smc*); mean textural fill volume (*Tfv*); mean heterogeneity 3x3 (*HAsfc*<sub>9</sub>); mean heterogeneity 9x9 (*HAsfc*<sub>81</sub>). Lines associated with each bar indicate standard error above and below the mean.
- **Figure 5.** Bivariate plot of M<sub>1</sub> shear facet microwear texture anistropy and complexity. Data are plotted for individuals with species as indicated by the markers.

# Tables

# TABLE 1.Extant bovid taxa included in this study, along with associated diets

# TaxonCommon NameDietObligate Grazers (>90% monocots, little to no seasonal or geographic variation)

Damaliscus lunatus	Tsessebe	Savanna and floodplain grasses
Kobus leche	Lechwe	Aquatic marsh grasses
Redunca arundinum	SouthernReedbuck	Tall grass specialist
Redunca fulvorufula	Mountain Reedbuck	Seasonal grasses

# Variable Grazers (60-90% monocots, varies seasonally)

<i>Damaliscus pygargus</i> consumption	Bontebok	Short Highveld grasses, some fynbos
<i>Gazella granti</i> foliage	Grant's Gazelle	Short grasses, seasonal herbs and shrub
Hippotragus niger	Sable Antelope	Grasses in wet season and forbs and foliage
Kobus ellipsiprymnus rushes	Waterbuck	Medium to short grasses, reeds and some
<i>Synceos caffer</i> vegetation during dry	African Buffalo season	Pasture grasses year-round and swamp
<i>Tragelaphus spekii</i> level vegetation	Sitatunga	Swamp grasses, sedges, leaves and low-

# Browser-Grazer Intermediates (30-70% monocots, <20% fruit)

Aepyceros melampus In	mpala	Grazes on grasses when green, also browses on
S	eedpods, foliage, fort	os and shoots
Antidorcas marsupialis	Springbok	Shrubs and succulents in dry season, young
grasses during wet sease	on	
Raphicerus campestris	Steenbok	Low herbaceous plants, seeds and pods
during dry season, youn	g grass in wet season	l
Raphicerus sharpei	Sharpe's Grysbok	Mature plants, fruit pods and roots
Taurotragus oryx E	land	Seasonal young grasses, small herbaceous plants,
roots, and tubers, highly	v seasonal	
Tragelaphus imberbis	Lesser Kudu	Succulents, buds, leaves, pods and grasses,
pronounced seasonal va	riation	

### **Browsers (>70% dicots)**

Litocranius walleriGerenukPure browsers, focuses on flowers andleaves of shrubs, not known to eat grass or herbsNeotragus batesiBate's Pygmy AntelopeGround-level herbaceous vegetation

Sylvicapra grimmiaCommon Duikerinsects, rarely known to consume grassesTragelaphus eurycerosBongoshrubs and creeping plants

Leaves, shoots, fruit pods and seeds, tubers,

High-concentrate green herbage, including

# Generalists (>20% of all three food types, highly variable by season and region)

Oreotragus oreotragus K	lipspringer	Leaves, buds, pods, bark, short grasses and
some fruit		
Tragelaphus angasi Nyala	L	Mixed grasses and leaves
Tragelaphus strepsiceros G	reater Kudu	Herbs, leaves, vines, fruit, tubers and some
grass		

# Frugivores (>70% fruit, rarely consumes monocots)

Cephalophus sylvicultor	Yellowback Duiker	Forbs and shrubs
Philantomba monticola	Blue Duiker	Fallen fruit, leaves, buds, seeds and
mushrooms		

Diet information based on Walker, 1975; Kingdon, 1982, 1997; Grzimek, 1990; Nowak, 1991. Gagnon and Chew, 2000

Taxon	Statistic n	Asfc	epLsar	Smc	Tfv HAsfc9	HASF	C81
A. melampus	Mean 20	1.868	0.004	2.842	5878.293	0.498	0.831
-	Median	1.809	0.004	0.512	5719.810	0.455	0.749
	Standard Deviation	0.482	0.002	10.071	1062.347	0.145	0.237
	Skewness	0.815	0.480	4.467	0.927 1.475	0.377	
A. marsupialis	Mean 21	2.382	0.004	4.082	6038.774	0.510	0.786
	Median	2.064	0.004	0.267	5733.700	0.480	0.799
	Standard Deviation	1.138	0.001	17.085	1269.053	0.136	0.138
	Skewness	1.896	-0.271	4.581	0.424	0.725	-0.115
C. sylvicultor	Mean 25	4.617	0.003	0.253	12405.420	0.494	0.801
	Median	4.628	0.003	0.267	13081.060	0.498	0.811
	Standard Deviation	1.263	0.001	0.048	1790.455	0.052	0.052
	Skewness	0.396	-0.140	-0.788	-0.375	-0.204	-0.481
D. lunatus	Mean 22	0.952	0.007	1.474	2174.730	0.435	0.657
	Median	0.909	0.007	1.417	1912.030	0.440	0.672
	Standard Deviation	0.247	0.001	0.437	944.932	0.065	0.074
	Skewness	0.164	0.505	0.580	0.698	0.036	-0.171
D. pygargus	Mean 22	1.281	0.005	1.316	5013.254	0.381	0.608
	Median	1.265	0.005	1.330	4917.000	.0389	0.609
	Standard Deviation	0.390	0.001	0.207	1663.663	0.091	0.076
	Skewness	0.443	-0.070	-0.170	0.625	-0.469	0.334
G. granti	Mean 25	1.680	0.004	0.525	3193.148	0.358	0.676
	Median	1.816	0.004	0.532	3070.000	0.364	0.692
	Standard Deviation	0.448	0.001	0.309	1243.082	0.073	0.118
<u>.</u>	Skewness -0.27	6 0.623	0.623		0.747 0.068	0.102	
H. niger	Mean 22	1.464	0.006	0.271	5001.706	0.525	0.735
	Median	1.530	0.006	0.267	4929.280	0.536	0.740
	Standard Deviation	0.478	0.001	0.102	1047.802	0.087	0.097
	Skewness075	1 -0.243	0.206		0.127 -0.123	0.045	
K. ellipsypry.	Mean 23	1.299	0.006	0.803	3608.550	0.622	1.110
	Median	1.361	0.006	0.816	3473.530	0.535	0.803
	Standard Deviation	0.452	0.001	0.125	1077.223	0.267	0.768
	Skewness	0.352	-1.264	-0.182	0.007	1.365	2.709
K. leche	Mean 24	0.986	0.006	1.598	2913.420	0.379	0.790
	Median	0.954	0.006	1.558	2975.175	0.378	0.751
	Standard Deviation	0.267	0.001	0.544	980.589	0.056	0.119
	Skewness	0.823	0.353	0.452	-0.007	-0.072	0.793
L. walleri	Mean 25	3.410	0.002	0.990	11136.860	0.754	1.076
	Median	3.265	0.002	0.965	10685.150	0.699	0.994
	Standard Deviation	1.099	0.001	0.350	1874.815	0.181	0.196
	Skewness	0.473	-0.001	1.036	0.476	0.830	0.841
N. batesi	Mean 22	3.833	0.002	0.833	10254.340	0.601	0.964
	Median	4.121	0.002	0.764	9875.320	0.599	0.932
	Standard Deviation	1.099	0.001	0.283	1603.675	0.112	0.127

TABLE 2.Descriptive microwear texture statistics for extant African Bovidae

	Skewness -0.1	255 -0.1	04 1.259		0.389 0.	338 1.067	
O. oreotragus	Mean 24	2.239	0.004	0.344	8975.312	0.435	0.791
0	Median	2.223	0.003	0.313	8587.210	0.446	0.742
	Standard Deviation	n 0.573	0.001	0.129	1851.004	0.084	0.183
	Skewness	0.851	0.339	0.470	0.678	-0.703	1.088
P. monticola	Mean 21	4.526	0.002	0.201	14629.710	.0551	0.813
	Median	4.215	0.002	0.203	14597.540	0.570	0.836
	Standard Deviation	n 1.036	0.001	0.046	1881.079	0.089	0.119
	Skewness	0.400	0.198	0.177	-0.270	-0.483	0.027
R. campestris	Mean 21	2.296	0.003	0.690	8376.412	0.456	0.863
1	Median	2.355	0.003	0.668	7420.810	0.488	0.874
	Standard Deviation	n 0.543	0.001	0.181	1617.150	0.088	0.155
	Skewness	0.138	0.023	1.056	-0.188	-0.649	-0.296
R. sharpei	Mean 25	2.350	0.004	0.527	4816.287	0.577	0.937
1	Median	2.523	0.004	0.433	4862.960	0.583	0.932
	Standard Deviation	n 0.429	0.001	0.218	1354.920	0.100	0.150
	Skewness	-0.417	2.586	1.213	-0.186	-0.235	1.221
R. arundinum	Mean 25	0.922	0.006	1.585	3349.817	0.371	0.784
	Median	0.865	0.006	1.698	3497.230	0.367	0.794
	Standard Deviation	n 0.216	0.001	0.530	1001.484	0.041	0.088
	Skewness	0.666	0.137	0.092	-0.307	0.336	-0.087
R. fulvorufula	Mean 24	1.199	0.008	1.084	1717.802	.0413	0.845
	Median	1.184	0.007	1.192	1657.615	0.414	0.620
	Standard Deviation	n 0.241	0.001	0.236	515.621	0.094	1.155
	Skewness	-0.097	0.729	-0.164	0.670	0.487	4.851
S. grimmia	Mean 25	3.076	0.003	0.629	11690.790	0.725	0.884
	Median	3.201	0.003	0.567	11599.350	0.608	0.884
	Standard Deviation	n 0.774	0.001	0.242	3233.051	0.289	0.157
	Skewness -	-0.128 -0	0.086	1.136	0.039	0.813	1.026
S. caffer	Mean 22	1.490	0.005	0.775	2847.686	0.434	0.719
	Median	1.483	0.005	0.743	2872.690	0.440	0.731
	Standard Deviation	n 0.454	0.001	0.259	1184.979	0.085	0.139
	Skewness	0.397	0.557	1.771	0.146	-0.106	-0.035
T. oryx	Mean 21	2.123	0.004	0.455	7101.973	0.539	0.950
	Median	2.133	0.004	0.416	6829.630	0.504	0.842
	Standard Deviation	n 0.453	0.001	0.289	1289.774	0.164	0.218
	Skewness	0.540	-0.048	0.845	0.260	0.841	0.501
T. angasi	Mean 22	2.091	0.004	0.296	9741.029	0.446	0.810
	Median	1.876	0.004	0.267	9882.410	0.441	0.794
	Standard Deviation	n 0.730	0.001	0.117	1506.152	0.089	0.153
	Skewness	0.574	0.635	0.253	-0.200	0.180	0.801
T. euryceros	Mean 22	3.779	0.002	.0816	12568.670	0.611	0.986
	Median	3.799	0.002	0.854	12740.510	0.624	0.969
	Standard Deviation	n 0.817	0.000	0.265	2306.309	0.110	0.183
	Skewness -0.0	92 -0.93	57	-0.144	-0.037	0.126	0.301
T. imberbis	Mean 23	2.038	0.004	0.234	7206.095	0.481	0.884

	Median	1.960	0.004	0.267	7127.730	0.461 0.873
	Standard Deviati	on 0.606	0.001	0.057	2062.956	0.088 0.227
	Skewness	1.316	-0.386	-0.009	-0.151	0.195 0.232
T. spekei	Mean 22	2 1.897	0.005	1.046	2773.355	0.406 0.917
	Median	1.885	0.004	1.009	2545.770	0.400 0.879
	Standard Deviati	on 0.450	0.001	0.206	1020.237	0.061 0.131
	Skewness	0.344	-0.331	0.405	0.314	0.501 .0514
T. strepsicero.	os Mean 25	5 2.328	0.004	0.263	8836.616	0.450 0.882
	Median	2.388	0.004	0.267	9027.370	0.465 0.866
	Standard Deviati	on 0.654	0.001	0.087	1914.357	0.070 0.176
	Skewness	-0.263	0.331	0.432	-0.194	0.269 0.943

Source	SS	df	MS	F	,	р	
Complexity (Asfc r	anked data)						
Diet	11114250	5	22228	350	296.4	188	0.00
Species (diet)	5405	529.7	19	28448.9	30 3.7	795	0.00
Error	4115	5996		549 7	,497.261		
Anisotropy (epLsar	r ranked data)						
Diet	9967934	5	19935	587	228.1	61	0.00
Species (diet)	6042	23.02	19	60423.02	6.9	915	0.00
Error	4796	5959		549 8'	737.631		
Maximum Complex	xity ( <i>Smc</i> ranke	d data)					
Diet	8697416	5	17394	483	249.8	368	0.00
Species (diet)	3253	3916	19	171258.8	3 24.6	500	0.00
Error	3921917			549 6961.597			
Fill Volume ( <i>Tfv</i> ra	inked data)						
Diet	12784997	5	25569	999	662.0	002	0.00
Species (diet)	9088	375.9	19	47835.57	12.3	385	0.00
Error	2120	)527		549 3	862.526		
Heterogeneity 3x3	(HAsfc9 ranked	data)					
Diet	5026810	,	5	1005362		64.357	0.00
Species (diet)	2037	7744		19 10	07249.7	6.865	0.00
Error	8573	3277		549 1:	5621.63		
Heterogeneity 9x9	(HAsfc <sub>81</sub> ranked	d data)					
Diet	3473674	,	5	694734.8	39.81	0.0	0
Species (diet)	2685	5031		19 14	41317.4	8.098	0.00
Error	9580	)677		549 1'	7451.14		

# TABLE 3.Nested analyses of variance

Categories		Asfc	epLsar	Smc	Tfv	HAsfc9	Hasfc81
B-G Int.	Browser	-147.829 **	142.73 **	-136.633 **	-171.974 **	-137.736 **	-88.847 **
B-G Int.	Frugivore	- 208.329* *	114.949 **	129.234 **	-225.374 **	-40.521	43.049
B-G Int.	Generalis t	-7.037	-32.485	84.865 **	-105.897 **	77.087 **	42.203
B-G Int.	Obl. Grazer	236.792 **	-245.575 **	-272.592 **	209.901 **	153.011 **	146.922 **
B-G Int.	Var. Grazer	124.517 **	-145.656 **	-112.54 **	142.105 **	86.074 **	107.406 **
Browser	Frugivore	-60.500 **	-27.781	265.867 **	-53.400 **	97.215 **	131.896 **
Browser	Generalis t	140.792 **	-175.216 **	221.498 **	66.076 **	214.824 **	131.05 **
Browser	Obl. Grazer	384.621 **	-388.305 **	-135.959 **	381.874 **	290.748 **	235.769 **
Browser	Var. Grazer	272.346 **	-288.386 **	24.093	314.079 **	223.811 **	196.254 **
Frugivore	Generalis t	201.292 **	-147.435 **	-44.369	119.476 **	117.608 **	-0.846
Frugivore	Obl. Grazer	445.121 **	-360.524 **	-401.826 **	435.274 **	193.532 **	103.873 **
Frugivore	Var. Grazer	332.846 **	-260.605 **	-241.774 **	367.479 **	126.595 **	64.358 *
Generalis t	Obl. Grazer	243.829 **	-213.09 **	-357.457 **	315.798 **	75.924 **	104.719 **
Generalis t	Var. Grazer	131.554 **	-113.171 **	-197.405 **	248.003 **	8.987	65.203 *
Obl. Grazer	Var. Grazer	-112.275 **	99.919 **	160.052 **	-67.795 **	-66.937 **	-39.516

TABLE 4.Pairwise comparisons test of dietary categories

\* indicates significant differences using Fisher's Least Significant Difference test, \*\* indicates significant differences using both Tukey's Honest Significant Difference Test and Fisher's Least Significant Difference test.
	Complex ity ( <i>Asfc</i> )	Anisotro py ( <i>epLsar</i> )	Scale of Maximu m Complex ity ( <i>Smc</i> )	Fill Volume ( <i>Tfv</i> )	Heterogen eity 3 x 3 ( <i>HAsfc9</i> )	Heterogen eity 9 x 9 ( <i>HAsfc81</i> )
				2306.851		
Obligate Grazer	0.9845	0.0065	1.3425	6	0.3871	0.698
				3691.862		
Variable Grazer	1.5015	0.0053	0.7741	9	0.4291	0.7368
Browser-Grazer				6248.325		
Intermediate	2.0632	0.0037	0.4173	3	0.4968	0.8655
				2736.054		
Generalist	2.2553	0.0038	0.2671	9	0.4482	0.8174
				10975.14		
Browser	3.6115	0.0022	0.7665	46	0.6215	0.9513
				13569.55		
Frugivore	4.3993	0.0023	0.2346	72	0.5162	0.8117

# TABLE 5. Median values for all microwear variables by dietary category.

Chapter Three: Dental microwear texture analysis of fossil bovids from Hadar, Ethiopia: implications for the paleoenvironment of *Australopithecus afarensis* 

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## Abstract

Considerable paleoecological research has focused on the Pliocene locality of Hadar in Ethiopia, due to the long occupation by Australopithecus afarensis. In order to better understand the adaptations of these early hominins, it is necessary that reconstructions of the paleoenvironment be as accurate as possible. This study employs dental microwear texture analysis to reconstruct the diets of the local bovid fauna, frequently used as indicator taxa for habitat. To interpret the microwear signatures, the Hadar taxa are compared to a database including 25 extant bovid species with well-documented diet and habitat preferences. This paper also addresses the paleoenvironments of the Sidi Hakoma, Denen Dora and Kada Hadar Members, as inferred from the dietary data. Clear differences in bovid microwear textures were found between members. The Sidi Hakoma bovids are reconstructed as browsers or browsergrazer intermediates, suggesting the presence of woodland resources, and closed habitats. The Denen Dora bovids evince microwear textures ranging from obligate grazers to browser-grazer intermediates, indicating access to both browse and graze and a mosaic habitat. The Kada Hadar bovids primarily have variable or obligate grazing signatures, although a few browsing taxa are present. This suggests open habitats for grazing, although the presence of browsing taxa supports a mosaic reconstruction for this member. The results presented here are consistent with a trend toward increasing aridification over time at Hadar. The microwear signatures of the fossil bovids suggest a transition from closed habitats during accumulation of the Sidi Hakoma Member to a more open environment by the time of Kada Hadar deposition.

Keywords: Australopithecus afarensis, microwear, paleoenvironment, diet, Hadar, bovid

# Introduction

An important goal of paleoanthropology is to understand the ecological backgrounds in which early hominins lived, and the climatic and environmental changes that likely influenced their evolution. In order to understand the relationships between hominin evolution and environmental change, it is first necessary to have the most robust possible interpretations of the habitats occupied by our early forbearers. Techniques for reconstructing these paleoenvironments include the use of floral and faunal assemblages as proxies and geochemical analysis of paleosols. However, these techniques address paleoecological questions of different temporal and geographic scales and frequently do not offer consistent results.

Considerable recent research has focused on reconstructing local habitats at the Hadar in Ethiopia, during the long occupation period of *Australopithecus afarensis*. Although a wide variety of studies, including those utilizing palynological remains (Radosevich et al., 1992; Bonnefille et al., 2004), ecomorphological and diversity analyses of faunal assemblages (Bobe and Eck, 2001; Reed, 2008) and stable isotope data (Hailemichael, 2000), have suggested a mosaic of microhabitats at Hadar throughout the temporal range of *A. afarensis*, some disagreement exists about the amount of fluctuation in habitat type, rainfall and humidity.

In this paper, I address paleoecological change at Hadar by reconstructing the diets of fossil bovids from the Sidi Hakoma, Denen Dora and Kada Hadar Members using dental microwear texture analysis. The microwear data for the fossil taxa are compared with a large database of texture signatures for extant African bovids, with results discussed in relation to other paleoecological reconstructions of the site. I also address variation in the diets of Hadar bovid taxa found in multiple members and its implications for habitat change over time. Finally,

I discuss the reconstructed habitats for Hadar as compared to the *A. afarensis* localities at Laetoli and Koobi Fora.

## Hadar

The Hadar hominin site in the Afar Region of Ethiopia dates to ~3.42-2.9 mya and includes the last known locality associated with *A. afarensis* (Walter et al., 1984; Walter and Aronson, 1993; Walter, 1994; Campisano and Feibel, 2007). Approximately 370 specimens of *A. afarensis* have been recovered from the site, along with more than 7500 other vertebrate specimens (Reed, 2008). The Hadar faunal assemblage is both diverse and extensive, including camels, hippos, antelopes, pigs, giraffes, rhinos, true and false saber-toothed cats, jackals, lions, hyenas, mongooses, weasels, rabbits, horses, civets, monkeys, elephants, deinotheres, and rodents.

The Afar triangle was occupied by *A. afarensis* for more than 500 ky and the remains of fossil hominins have been recovered from three of the four recognized members: Sidi Hakoma, Denen Dora and Kada Hadar. Each of the Hadar members is composed of geologic layers bound by a radioisotopically dated tephra (Taieb et al., 1976; Campisano, 2007). These members are then subdivided into smaller stratigraphic units based on the presence of marker beds. There are ten submembers that make up the Hadar Formation: Basal, Sidi Hakoma 1-4, Denen Dora 1-3, and Kada Hadar 1-2. Studies of local geology have revealed that these strata were deposited during periods of high fluvial activity of a large river system associated with the site, as well as incursions of a reoccurring paleolake (Aronson and Taieb, 1981).

The Basal Member has been reliably dated to 3.8-3.42 mya and has not produced hominin remains, although they have been recovered from the formation and the same time

period at the nearby site of Dikika, Ethiopia (Alemseged et al., 2005). The composition of faunal remains from this member has suggested a mosaic habitat composed of woodland and shrubland. The bovid assemblages contain congeners of modern taxa adapted to both open and closed environments. This includes abundances of alcelaphins, typically adapted to open grassland today; tragelaphins, which inhabit a variety of partially or totally closed environments; and aepycerotins, which occupy woodlands (Skinner and Smithers, 1990). Palynological studies have inferred a closed forest habitat, although it is possible that floral sampling was limited to a forested area of a more complex environment. Regardless, most reconstructions agree that the local habitat was more closed during the formation of the Basal Member than is evidenced later at the site (Bonnefille et al., 2004).

The Sidi Hakoma Member was deposited between ~3.42-3.26 mya (Campisano, 2007). Previous studies suggest some subtle differences in the paleoenvironments between the four submembers. The rarity of grazing bovid taxa suggests a more closed habitat and the reconstructed locomotor and dietary adaptations of the local mammalian fauna suggest that there was higher annual rainfall and less seasonality during the deposition of the lower Sidi Hakoma submembers than any other time at the site (Reed, 2008). A study of the local mollusks by Hailemichael (2000) supports this reconstruction, with carbon and oxygen isotope analyses consistent with a three month annual dry season during this time. The upper Sidi Hakoma Member has been reconstructed as more mosaic, including scrub woodlands and floodplain grasslands. The local faunal assemblages include both grazing and browsing bovids, as well as wetland taxa like reduncins and hippotamids (Reed, 2008). Additionally, the pollen record from these upper members suggests slightly drier conditions than those recorded for SH-1 and SH-2 and is consistent with closed to open woodland and wet grassland (Bonnefille et al., 2004).

Overall, both the palynological data and the faunal record from throughout the member have suggested a shift from wetter, more closed conditions to slightly drier, more open ones (Hailemichael, 2000; Bonnefille et al., 2004; Reed, 2008).

The Denen Dora Member of the Hadar Formation dates between ~3.26-3.2 mya and is composed of three submembers. In general, composition of the faunal assemblages within Denen Dora suggests that Hadar became drier and more seasonal over this comparatively brief interval (Reed, 2008). The oldest fauna in the Denen Dora Member, including reduncin, aepycerotin and tragelaphin bovids, are associated with highly mosaic habitats, including gallery forest, bushland and woodland. Evidence suggests that there was a lake incursion during the DD-2 phase and the faunal assemblages reflect an increase in the abundance of wetland adapted taxa. Finally, a slight faunal shift is evident in the more recent layers of the Denen Dora Member, with open woodland and wooded grassland adapted taxa like Alcelaphini becoming more common (Reed, 2008). Analysis of fossil pollen has revealed a corresponding increase in C3 grasses and a decrease in annual precipitation (Bonnefille et al., 2004). All of this suggests that Hadar was mosaic and fluctuating during the formation of the Denen Dora Member.

The Kada Hadar Member dates from ~3.2 mya-2.94 mya and contains the last known occurrences of *A. afarensis*. The member is divided into two submembers, both of which have been generally reconstructed as drier and more open than the earlier deposits at Hadar. Fauna from the hominin-bearing strata share adaptations for habitats ranging from open woodlands to savanna grasslands (Reed, 2008). For example, the predominant bovid genera found in the deposit are Antilopini and Alcelaphini, both associated with dry, open environments (Vrba, 1980; Kingdon, 1982a,b). In contrast, however, studies of fossil pollen from the Kada Hadar Formation have suggested a shift back to a wetter woodland habitat (Bonnefille et al., 2004).

Reed (2008) suggested that this discrepancy might be indicative of multiple closed physiognomies within the geologic member.

## Previous paleoecological reconstructions at Hadar

Paleoenvironmental reconstructions based on associated fauna, sedimentology and paleosols at Hadar have generally indicated a temporary mosaic of habitats that fluctuated between open and closed and woodland and grassland (Johanson et al., 1982; Radosevich et al., 1992). The presence of these different habitat types can indicate the presence of subterranean water, changes in soil type, or even incursions from river systems or lakes (Puhakka et al., 1992; Dodds, 1997). All of these factors, related to the source of water and the amount available, can alter vegetation patterns at a site dramatically and account for modern sites where disparate habitat types are found in the immediate vicinity of one another (Archibold, 1995; O'Brien, 1998, 2006; Andrews, 2006; Reed, 2008).

The examination of the faunal assemblages from each individual member provides a unique opportunity to assess paleoecological change during the occupation of *A. afarensis*. However, reconstructions of mammalian paleoecology have traditionally relied upon taxonomic uniformitarianism, a principle based on the assumption that fossil taxa shared the same ecological preferences as their closest living relatives. In most cases, this is probably accurate; however, Sponheimer et al. (1999) have suggested that the principle deserves further scrutiny. In their study on dietary reconstruction of the Makapansgat bovids using stable isotopes and ecomorphology, these authors concluded that two of the seven fossil bovid taxa had isotopic signatures consistent with different diets than those suggested using only phylogenetic similarities. Thus, while taxonomic uniformitarianism may be generally reliable, it must be

independently verified in some reliable way in order to have confidence in the assumed relationships.

In addition to comparisons within the site, studies of the Hadar fauna in relation to other Pliocene localities have also been useful for understanding the local paleoenvironment. Reed (2008) used the absence and presence of specific mammalian taxa to characterize the local community ecology at Hadar and compared the results to reconstructions of other known A. afarensis localities, including Maka and Laetoli. Adaptations of fossil taxa were assumed based on taxonomic uniformitarianism and/or previous dietary reconstructions using stable isotope analysis. In general, these data suggest that the environment at Hadar was highly mosaic and included intermediate cover habitats like scrubland, bushland, and open woodland, with occasional incursions of wetlands or edaphic grasslands over time (See Table 1 for a description of the habitat categories referenced in this study). Bobe and Eck (2001) also suggested a mosaic habitat at the site. They compared the abundance of bovid remains from Hadar and compared them to those recovered from the Shungura Formation in the lower Omo Valley, Ethiopia, where a distinct shift toward more open, arid habitats has been recorded. The study concluded that there is no evidence for a faunal shift towards open grassland habitats at Hadar. Instead, the assemblage of bovid taxa suggests that the overall habitat at the site was composed primarily of open grassland and woodland.

Hadar, like the other early hominin habitats in eastern Africa, has a paleoenvironmental signature that is not completely understood. Different analyses have returned results ranging from primary wetlands to a mosaic habitat of open savanna and closed woodland forest. Improving the paleoecological assumptions made about the fauna recovered from the site is an

important step in achieving an accurate understanding of the Pliocene environment at Hadar and determining how it might have shaped the evolutionary adaptations of *A. afarensis*.

## **Bovid diets and paleodiets**

African bovids have long been used as paleoenvironmental indicator species because modern taxa can be grouped into categories based on specific diet and habitat preferences. Even though the literature on bovid ecology is extensive, it is also often contradictory, resulting in considerable debate about how to best classify bovids by diet. One early attempt to improve classification of bovids was based on consumption of roughage versus herbage. This system, described by Hofmann and Stewart (1972), divided taxa into three dietary categories: grazers, browsers and intermediate feeders. Taxa in the latter category were then subdivided based on the relative percentages of food types included in the diet. Bodmer (1990) argued that fruit is a staple dietary component for many bovid species and proposed a new continuum ranging from grazer to frugivore, with browsing as an intermediate category. In an effort to address the lack of consensus on the dietary classifications of bovids, Gagnon and Chew (2000) offered a system based on a synthesis of field research on 78 African bovid diets. Their six dietary classifications were based on the observed percentages of monocotyledons and dicotyledons included in the diet, with fruit considered separately from other dicots. The Gagnon and Chew system also takes into account the effect of seasonality and geographic changes on diet.

In this study, I employ the system of Gagnon and Chew (2000) because it not only distinguishes grazing and browsing, but also fruit from leaf browse, and recognizes various levels of mixed feeding. These categories have been determined by the percentages of fruits versus non-fruit vegetation, and dicotyledons versus monocotyledons included in the diet, as

recorded during observations in the wild. While the inclusions of these various levels of mixed feeding cannot help but improve the resolution of dietary and habitat reconstructions of fossil taxa, extensive ecological observations and fecal analyses have not been conducted for many mixed-feeding taxa (Gagnon and Chew, 2000, Fortelius and Solounias, 2002). Therefore, further field work will eventually be required if we are to fully understand the influence of seasonal and geographic variation on the diets of these bovids.

Gagnon and Chew (2000) distinguish obligate grazers, variable grazers, browser-grazer intermediates, browsers, generalists and frugivores. Obligate grazing taxa have a diet made up of more than 90% monocotyledons, with no seasonal or geographic variation. Taxa classified as variable grazers consume 60-90% monocotyledons, but may vary in diet seasonally and geographically. Browser-grazer intermediates have diets consisting of 30-70% monocotyledons and dicotyledons, always including some fruit, although never more than 20%. Browsing bovids consume more than 70% dicotyledons. Fruit accounts for at least 70% of the diet of frugivorous bovids and these taxa rarely, if ever, consume monocotyledons. Finally, generalist taxa classified are highly variable by both locality and season, but they consistently consume more than 20% of each of the three food types.

## Implications of diet for habitat reconstruction

The selection and consumption of food is the most direct way that an organism interacts with its environment; therefore the reconstruction of diet holds the potential to tell us much about that animals preferred habitat. However, many extant bovids utilize multiple habitat types seasonally and/or depending on geographic range, so it is not reasonable to expect a one-to-one correspondence between diet and habitat. However, dietary reconstruction does have

implications for bovid habitat preference that can be cautiously applied to the fossil record. Even though the correlation between diet and habitat is not exact, grazing and browsing microwear signatures do have implications for availability of grasses and woody plants respectively (Schubert et al., 2006). For example, all obligate grazing taxa occupy some type of grassland, although this category includes dry and moist savanna, floodplain grassland and marshland. Dedicated browsing taxa, on the other hand, always live in or near forest or shrubland, which can also include open and closed woodland, bushland and closed forest. Dedicated frugivorous species either inhabit or have ready access to closed forests or open woodlands (Hofmann and Stewart, 1972; Kingdon, 1982a,b; Skinner and Smithers, 1990).

The relationship between diet and habitat is less exact for taxa that consume a variety of food types. Even so, dietary reconstruction of these "mixed feeders" can still inform habitat preference. Variable grazers are typically found in some type of grassland, although a few inhabit open woodland (Kingdon, 1997). Browser-grazer intermediate taxa all occupy forest or bushland habitats for at least part of the year, although some range into savannas and grasslands during dry seasons. Generalist taxa are unsurprisingly the most difficult to classify, occupying a wide range of environments, including grassland, bushland, open woodland and closed forest habitats (Hofmann and Stewart, 1992; Dunbar and Dunbar, 1974; Kingdon, 1982a,b, 1997; Skinner and Smithers, 1990). In sum, while there is no exact correlation between diet and habitat, the ability to accurately classify diet beyond the traditional grazer-browser-mixed feeder continuum can help us narrow down habitat preferences to a likely range of local environments.

## **Dental microwear analysis**

Conventional methods of studying dental microwear include scanning electron microscopy and light microscopy, both subject to observer measurement error and a lack of three-dimensional surface characterizations. Dental microwear texture analysis was developed to address these issues and provide an objective and replicable method for characterizing surfaces. The technique combines white-light confocal profilometry and scale-sensitive fractal analysis, allowing for quantitative descriptions of microwear surfaces over a range of scales. White-light confocal profilometry is also faster, less expensive and easier to use than scanning electron microscopy; and the automated analysis is less time consuming and requires less effort for quantification than identification and measurement of individual features. Most importantly, the automated nature of texture analysis reduces the amount of observer error in measurement, which allows direct comparisons between studies, as well as the establishment of a large database that researchers can access for interpretation of their results. Dental microwear texture analysis has previously been used to successfully identify dietary differences between species of extant and fossil taxa including humans and non-human primates, marsupials, carnivores, and bovids (R. Scott et al., 2005, 2006; Ungar et al., 2008a, 2008b, 2010, in press; Krueger and Ungar, 2009; Merceron et al., 2009; Prideaux et al., 2009; J. Scott et al., 2009; Ungar and Scott, 2009; Schubert et al., 2010).

## **Materials and Methods**

# Specimens

This study includes 121 fossil bovid specimens from the Sidi Hakoma, Denen Dora and Kada Hadar geologic members of the Hadar Formation (summarized in Table 2). The fossil bovids represent 12 taxonomic tribes, often the most specific identification available for bovid specimens not accompanied by distinctive horn cores. Taxa identifiable to the level of genus were considered both inclusive and exclusive of other members of the tribe. Molds of upper and lower first and second molars were collected at the Ethiopian National Museum (NME) in June 2010. Microwear analysis of these different tooth types has been shown to yield similar results (Merceron et al., 2004a, b). All available bovid specimens from the Hadar Formation were examined and the vast majority had to be excluded due to postmortem damage, following the criteria of Teaford (1988) and King et al. (1999). Unworn teeth and those with extreme wear were also excluded from the study. Ideally, comparisons would have been made among the individual Hadar submembers rather than the members themselves. Unfortunately, the sample sizes were not consistently large enough when divided out by submember, so the analysis presented here contrasts the Sidi Hakoma, Denen Dora and Kada Hadar bovids exclusive of submember.

In order to interpret the microwear signatures of the fossil taxa, the study also includes a comparative sample of 575 specimens, representing 25 extant African bovid species (Scott, in press). These extant specimens are housed at the American Museum of Natural History (AMNH), New York; the Field Museum (FMNH), Chicago; the Smithsonian National Museum of Natural History (NMNH), Washington D.C.; and the Royal Museum of Central Africa (RMCA), Tervuren, Belgium. All extant specimens were wild-shot and have known provenience data. The extant taxa were selected because they vary widely in ecological adaptations, ranging

from open-country-adapted obligate grazers to closed-habitat browsers to dedicated frugivores. The included species have well-documented diet and habitat preferences, and special emphasis was placed on including taxa that engage in various levels of mixed feeding, including variable grazers, browser-grazer intermediates and generalists. Table 3 lists the included extant taxa and summarizes their dietary preferences.

High-resolution casts were prepared following conventional procedures for microwear analysis, a process described in detail elsewhere (Rose, 1983; Grine, 1986, Ungar, 1996). The original specimens were cleaned with acetone or alcohol soaked cotton swabs and then molded using a polyvinylsiloxane dental impression material (Presidents Jet Regular Body, Coltene-Whaledent Corp.). The casts were then made using a high-resolution epoxy polymer (Epotek, Epoxy technologies Inc.) This process has been previously demonstrated to produce replicas that precisely reproduce microwear features with a resolution to a fraction of a micron (Beynon, 1987; Teaford and Oyen, 1989).

## **Data Collection**

A Sensofar PL $\mu$  white-light confocal profiler (Solarius Development Inc., Sunnyvale, CA) was used to collect the microwear texture data. This instrument allows for the collection of data points which form a 3-dimensional point cloud reproducing the surface of the tooth. These point clouds can be used to create photo simulations and digital elevation models of the wear surface. As in previous studies of bovid dental microwear, texture data were collected on the disto-buccal enamel band of the mesial cuspid of M<sub>2</sub> and the mesio-buccal enamel band of the mesial cusp of M<sup>2</sup> (Janis, 1990; Merceron, 2005; Schubert, 2006; Ungar et al., 2007), as illustrated in Figure 1.

The occlusal surface of each specimen was scanned with 100x objective, resulting in a point cloud with a lateral sampling interval of 0.18  $\mu$ m, a vertical resolution of 0.005  $\mu$ m, and a field of view of 102 x 138  $\mu$ m. Texture data were collected for four adjoining fields of view for a total work envelope of 204 x 276. Solarmap Universal software (Solarius Development Inc., Sunnyvale, CA) was then used to level each scan. Surface defects, including dust and other adherents, were removed from the scan using the erase or thresholding function in Solarmap.

The resulting point clouds were analyzed using Toothfrax and Sfrax scale-sensitive fractal analysis software (Surfract Corporation). Scale-sensitive fractal analysis is based on the principle that surfaces may appear different when viewed from different scales. Thus, a surface that appears smooth at a coarse scale may seem rough when viewed at a finer scale. Previous studies employing texture analysis have identified several variables as particularly informative for dietary reconstruction (Ungar et al., 2003; Scott et al., 2005, 2006). I present data for five of these variables here. They are area-scale fractal complexity, length-scale anisotropy of relief, scale of maximum complexity, textural fill volume and heterogeneity of area-scale fractal complexity. These variables will be summarized briefly here, as they have been described extensively elsewhere (Ungar et al., 2003, 2007, 2008; Scott et al., 2005, 2006).

Area-scale fractal complexity (*Asfc*) is a measure of change in surface roughness across a range of scales. This variable calculated by taking the slope of the steepest part of a curve fit to a plot of relative area over the range of scales at which the measurements are taken. Specimens with steeper slopes will have correspondingly higher values for *Asfc*, indicating a more complex surface. For example, surfaces dominated by overlying features of various sizes will result in high values for *Asfc*.

The scale of maximum complexity (*Smc*) refers to the fine scale limit of the *Asfc* line (Scott et al., 2005, 2006). This variable is calculated as the fine-scale limit of the steepest part of the curve described for the *Asfc* measure. It has been suggested that *Smc* corresponds with the size of wear-causing particles. Surfaces with less wear at finer scales will have the highest values for *Smc* and those with more wear at fine scales will have the lowest values.

The length-scale anisotropy of relief variable (*epLsar*)measures the orientation concentration of surface roughness. *EpLsar* is calculated by taking profiles of the wear surface at different orientations at a given scale. The sampling interval employed for this study was 5° and a scale of 1.8  $\mu$ m. Anisotropic surfaces have relative profile lengths that differ with orientation and these relative lengths at given orientations are defined as vectors and normalized, resulting in a rosette diagram. Mean values for the mean vector lengths are then calculated for *epLsar*. Microwear surfaces characterized by parallel scratches will have higher values for *epLsar* than those dominated by pits or scratches with different orientations.

Textural fill volume (*Tfv*)describes the volume of cuboids of various sizes that can digitally fill the microwear surface. The *Tfv* variable is the difference in summed volume for 2  $\mu$ m cuboids and 10  $\mu$ m cuboids, which removes the impact of facet curvature and allows for the characterization of the microwear features themselves. Microwear surfaces that have high values for this variable are typically those characterized by moderate size features.

While all the variables help to describe the overall texture of the microwear surface, surfaces themselves vary in their textures. The heterogeneity (*HAsfc*)variables can help quantify the extent of this variation and may be important in accurately characterizing the surface. This variable is calculated through the use of the Auto-Split function in Toothfrax and measures the difference in *Asfc* across a surface. The scanned area of the facet is divided into successively

smaller sub-regions with equal numbers of rows and columns, beginning with 2 x 2 and ending with 11 x 11. The value for *HAsfc* is defined as the median absolute deviation of *Asfc* divided by the median of *Asfc*.

# **Statistical Analysis**

Following scale-sensitive fractal analysis, median values were calculated for each of the four scans collected for each specimen (Scott et al., 2006). Resulting data were rank transformed prior to statistical analysis because unranked microwear texture data typically violate assumptions inherent in parametric statistical tests (Conover and Iman, 1981). I employed a nested analysis of variance model in this study, with taxon as the subordinate level of classification, nested within Hadar Member- Denen Dora, Sidi Hakoma and Kada Hadar. The data were then analyzed using a general linear model to determine whether there were overall differences in microwear among the taxa within the members, as well as among the members themselves. Analyses of variance for individual texture attributes and pairwise comparisons tests were then used to determine the sources of significant variation both among taxa within members and among the individual members. Both Tukey's Honestly Significant Difference and Fisher's Least Significant Difference tests were used to balance the risks of Type I and Type II errors (Cook and Farewell, 1996).

# Results

Examples of the microwear textures of the extant and fossil taxa are pictured in Figures 2 and 3. Median values for each texture variable by taxon are presented in Table 3. Descriptive and analytical statistics are presented in Tables 4, 5 and 6, as well as Figures 4, 5 and 6. To summarize, there are significant differences among the taxa, among the members, and among some of the taxa within each of the members. There is also significant variation between members for some individual taxa found in more than one member.

The general linear estimate model revealed significant differences among the three Hadar members in all texture variables except *Smc*. When considered as a group, the bovid taxa within the Sidi Hakoma Member have significantly higher values for *Asfc*, *Tfv* and *HAsfc*<sub>9</sub>, and the lowest recorded values for *epLsar*. The Denen Dora bovids have the lowest values for *Tfv*, and intermediate values for *Asfc*, *epLsar and HAsfc*<sub>9</sub>. The bovids associated with the Kada Hadar Member have significantly lower values for *Asfc* and *HAsfc*<sub>9</sub>, intermediate values for *Tfv*, and the highest recorded values for *epLsar*.

# **Results by member**

Sidi Hakoma Member- Significant variation was found among the eight bovid taxa from the Sidi Hakoma Member for all texture variables, with the exception of  $HAsfc_{81}$ .

*Asfc*- Alcelaphini, *Gazella* sp. and *Kobus* sp.cluster together and have significantly lower values for this variable than the other Sidi Hakoma taxa. *Tragelaphus kyaloae* and *Ugandax coryndonae* have similar *Asfc* values and are only significantly different from *Aepyceros* sp., Antilopini and Neotragini using a Fisher's LSD test.

*epLsar-Gazella* sp. has significantly higher values for this variable than any other taxon, while *Aepyceros* sp., *Kobus* sp. and Neotragini have the lowest values. *Tragelaphus kyaloae*, Alcelaphini and Antilopini all group together with intermediate values and are significantly different from the other taxa.

*Smc- Aepyceros* sp., Alcelaphini and *Ugandax coryndonae* have significantly lower values for this variable than the other taxa, while *Kobus* has the highest recorded value. Antilopini, *Gazella* sp., Neotragini and *Tragelaphus kyaloae* all cluster together with intermediate values.

*Tfv- Kobus* sp. has significantly lower *Tfv* values than any other taxon, while *Aepyceros* sp.,Antilopini, Neotragini and *Tragelphus* sp.have the highest values. Alcelaphini, *Gazella* sp. and *Ugandax coryndonae* all have intermediate values for this variable.

*HAsfc*<sub>9</sub>- The Sidi Hakoma bovids vary significantly in the *HAsfc*<sub>9</sub> variable, though this can primarily be explained by a few taxa with extreme values. There are not significant differences between most of the individual taxa, with the exception of *Kobus* sp., which has the lowest recorded values for this variable and differs significantly from all other taxa. The other notable taxa are Antilopini and Neotragini, both of which differ significantly from *Gazella* sp.

 $HAsfc_{81}$ - Although there is no significant variation in the overall model for this variable, significant differences in  $HAsfc_{81}$  are present between some of the individual taxa. *Gazella* sp. has the lowest recorded values for this variable and is significantly different than those of Neotragini and Antilopini, the taxa with the highest values for  $HAsfc_{81}$ .

<u>Denen Dora Member</u>- Significant variation was found among the ten bovid taxa from Denen Dora for all variables.

Asfc- Aepyceros sp., Alcelaphini, Hippotragini, Tragelaphus kyaloae and Ugandax coryndonae group together with intermediate values for this variable, with Antilopini and Gazella sp. having significantly higher complexity values and Reduncini, Pelorovis sp. and Kobus sp. having significantly lower values than the other taxa.

*epLsar-Aepyceros* sp., Antilopini, Alcelaphini and *Kobus* sp. have significantly lower values for this variable than the other taxa, with Antilopini having the lowest recorded values. *Gazella* sp., *Tragelaphus kyaloae* and *Ugandax coryndonae* all cluster together with intermediate values and Hippotragini, *Pelorovis* sp. and Reduncini have the highest values.

*Smc- Aepyceros* sp., Alcelaphini, Antilopini, *Gazella* sp., Hippotragini and *Ugandax coryndonae* cluster together with significantly lower values than Kobus sp., Pelorovis sp., Reduncini and *Tragelaphus kyaloae* 

*Tfv*- Antilopini has the highest values for this variable, while *Pelorovis* sp. and Reduncini have significantly lower values for this variable than any of the other taxa. *Aepyceros* sp., Alcelaphini, *Gazella* sp., *Tragelaphus kyaloae* and *Ugandax coryndonae* have intermediate values similar to one another, while Hippotragini and *Kobus* sp. cluster together with slightly, but still significantly, lower values.

*HAsfc*<sub>9</sub>- The Denen Dora bovids vary significantly in the *HAsfc*<sub>9</sub> variable, however, this difference is primarily accounted for by the significant difference between the values for Antilopini and *Gazella* sp. The taxa with the lowest recorded values for this variable are *Gazella* sp., *Aepyceros* sp., Hippotraginiand *Kobus* sp., while the taxa with the highest values are Antilopini, *Pelorovis* sp. and *Ugandax coryndonae* 

 $HAsfc_{81}$ - Tragelaphus kyaloae and Aepyceros sp. have similar  $HAsfc_{81}$  values that are significantly lower than those of the other taxa, while Alcelaphini and Antilopini cluster together with values that are significantly higher than those of any other bovid taxa.

<u>Kada Hadar Member</u>- Significant variation was reported among the ten bovid taxa from the Kada Hadar Member for all variables.

*Asfc*- Antilopini, *Gazella* sp. and Neotragini are similar to one another and have significantly higher values for surface complexity than the other seven taxa. Hippotragini, *Kobus* sp., *Oryx* sp. and *Pelorovis* sp.group together with significantly lower values than *Aepyceros* sp. and Alcelaphini using a Fisher's LSD test.

*epLsar-Aepyceros* sp., Antilopini, *Kobus* sp. and Neotraginihave significantly lower values than the other taxa, with *Kobus* sp. andNeotragini having the lowest recorded values for this variable. Alcelaphini, Bovini, *Oryx* sp. and *Pelorovis* sp. are not significantly different from one another and cluster together with significantly higher values for this variable than the other Kada Hadar taxa. *Gazella* sp. and Hippotragini have intermediate values for *epLsar* that are significantly different from *Kobus* sp., Neotragini, *Oryx* sp. and *Pelorovis* sp.

*Smc- Aepyceros* sp., Alcelaphini and Hippotragini cluster together with significantly lower values for this variable than any other taxa. Bovini, *Kobus* sp. and *Oryx* sp. have the highest recorded values, while Antilopini, *Gazella* sp., Neotragini and *Pelorovis* sp. have intermediate values.

*Tfv*- This variable divided the Kada Hadar bovids into two groups, with *Aepyceros* sp., Antilopini and Neotragini having significantly higher values for this variable than the other taxa. *Kobus* sp. and *Oryx* sp. have the lowest values for *Tfv*, while Antilopini and Neotragini have the highest.

*HAsfc*<sub>9</sub>- Antilopini, Neotragini and *Oryx* sp. have the highest recorded values for this variable and are significantly different from the remaining seven taxa. There are no significant differences in the *HAsfc*<sub>9</sub> variable among *Aepyceros* sp., Alcelaphini, Bovini, *Gazella* sp., Hippotragini, *Kobus* sp. and *Pelorovis* sp.

 $HAsfc_{8l}$ - Antilopini, Hippotragini, Neotragini and Oryx sp. group together and have the highest recorded values for this variable. This group is significantly different from all six other taxa in this member. *Gazella* sp. has an intermediate value for this variable and is significantly different from *Kobus* sp., Neotragini, Bovini, *Oryx* sp., *Pelorovis* sp. and Antilopini *Aepyceros* sp., Alcelaphini, Bovini, *Kobus* sp. and *Pelorovis* sp. are not significantly different from one another and have *HAsfc*<sub>8l</sub> values that are significantly lower than those of the other Kada Hadar bovids.

## **Results for taxa in multiple members**

*Aepyceros sp.*- This taxon was found in all three Hadar members used in this study, as well as the earlier Basal Member, and showed significant variation among members in all of the texture variables, with the exception of *HAsfc*<sub>81</sub>. The specimens from the Denen Dora and Kada Hadar members are similar in *Asfc, epLsar* and *Tfv*, while the Sidi Hakoma specimens are significantly different from the other two, with higher values for *Asfc, Tfv* and *HAsfc*<sub>9</sub>, and lower values for *epLsar*. The Kada Hadar and Sidi Hakoma specimens are not significantly different from one another in *Smc*, but both have significantly lower values than the specimens from Denen Dora.

Alcelaphini- This taxon has been recovered from all three hominin-bearing members at Hadar, as well as the Basal Member, and there are significant differences among the members for all of the texture variables. The specimens from Kada Hadar have significantly lower values for *Asfc*, *Smc*, *Tfv*, *HAsfc*<sub>9</sub> and *HAsfc*<sub>81</sub> and higher values for *epLsar* than those from Sidi Hakoma and Denen Dora. The specimens from Denen Dora have the highest values for *Smc* and *HAsfc*<sub>81</sub>. Antilopini- This taxon is reported for all four Hadar members. There were no significant differences among the Sidi Hakoma, Denen Dora and Kada Hadar specimens reported for any of the texture variables.

*Gazella sp.*- This taxon has been identified from the Sidi Hakoma, Denen Dora and Kada Hadar members. The only significant difference among *Gazella* specimens from these three members is in the *HAsfc*<sub>9</sub> variable. The specimens with the highest values for heterogeneity are those from the Sidi Hakoma Member, while those with the lowest values were recovered from the Denen Dora Member.

Hippotragini- This taxon is found in the all of the Hadar members included in this study; however only specimens from the Denen Dora and Kada Hadar members were suitable for microwear analysis. There are no significant differences between these members for any of the texture variables according to Tukey's HSD tests, although some suggestive differences were present according to Fisher's LSD test results. The Hippotragini specimens from the Kada Hadar Member have lower values for *epLsar* than those from Denen Dora, but again, this only significant when using a Fisher's LSD test.

*Kobus sp.*- This taxon has been recovered in the Sidi Hakoma, Denen Dora and Kada Hadar members. There were no significant differences among the members for any of the texture variables, although suggestive differences are present when analyzed with a Fisher's LSD test. This test suggests that the *Kobus* sp. specimens from Kada Hadar had lower values for *Tfv* than those from the Sidi Hakoma and Denen Dora members. Additionally, the *HAsfc*<sub>9</sub> values for *Kobus* sp.specimens from Sidi Hakoma were significantly lower than those of the Sidi Hakoma and Denen Dora specimens.

Neotragini- This taxon has been found in small quantities in the Sidi Hakoma, Denen Dora and Kada Hadar members. Only specimens from the Sidi Hakoma and Kada Hadar members were suitable for microwear analysis though there are suggestive differences between the members according to Fisher's LSD test. The *Asfc* values for the specimens from the Sidi Hakoma Member have marginally higher complexity values than those from Kada Hadar.

*Pelorovis sp.*- This taxon has been recovered from the Denen Dora and Kadar Hadar members. Significant differences are only present for the *HAsfc*<sub>9</sub>variable. The specimens from Denen Dora have higher values for this variable than those from Kada Hadar.

*Tragelaphus kyaloae-* This taxon has been recovered from all of the geologic members at Hadar, however only specimens from Sidi Hakoma and Denen Dora preserve unobscured microwear. Significant differences between specimens from these two members were identified in *Asfc, epLsar, Smc* and *HAsfc*<sub>81</sub>. The Sidi Hakoma specimens have significantly higher values for *Asfc* and *HAsfc*<sub>81</sub> and significantly lower values for *epLsar* and *Smc* than those from Denen Dora. A Fisher's LSD test also suggests significant differences in *HAsfc*<sub>9</sub> between the Sidi Hakoma and Denen Dora specimens, with individuals from Sidi Hakoma having higher values for this variable.

*Ugandax coryndonae*- This taxon has been recovered from all of the hominin-bearing members at Hadar, although this study only included specimens from the Sidi Hakoma and Denen Dora members. There are significant differences between the members reported for *Asfc*, *epLsar* and *Tfv*. The Sidi Hakoma specimens have significantly higher values for *Asfc* and *Tfv* and lower values for *epLsar* than those from Denen Dora.

# Discussion

The results presented here have important implications for the ability of dental microwear texture analysis to reconstruct the diets of fossil bovid taxa, and by extension, local paleoenvironments.

# **Extant Taxa**

The extant database used in this study has previously been used to demonstrate the presence of significant differences among the six dietary classifications identified by Gagnon and Chew (Scott, in review). Overall, grazing taxa had less complex, more anisotropic surfaces with smaller features than browsing taxa. In addition to clearly differentiating between obligate grazers and browsers, the data strongly indicate that dental microwear texture analysis can distinguish beyond the classic bovid dietary trichotomy and accurately separate variable grazers, generalists, browser-grazers intermediates and frugivores from obligate grazers and browsers, as well as from one another (Figure 4). Some differences among taxa within dietary categories were also identified, likely reflecting seasonal and/or geographic differences in diet.

#### Dietary reconstruction of the Hadar bovids by tribe

Overall, the Hadar bovids evince a range of microwear texture patterns comparable to the comparative extant taxa, suggesting similar levels of dietary variation in the taxa.

<u>Aepycerotini</u>- *Aepyceros* is the only genus within the tribe Aepycerotini and includes two extant subspecies: the common impala, *Aepyceros melampus melampus* and the black-faced impala, *Aepyceros melampus petersi*. Both of these modern taxa are classified as browser-grazer

intermediates, consuming equal proportions of monocots and dicots and including fruit as approximately 10% of their regular diet (Kingdon, 1997; Gagnon and Chew, 2000). Aepycerotine bovids typically inhabit a combination of thick bushveld and open savanna (Kingdon, 1992 a,b). Fossil specimens of *Aepyceros* sp. have been recovered from the Sidi Hakoma, Denen Dora and Kada Hadar members at Hadar and there is evidence of a change in the dietary signature over time at the site.

The *Aepyceros* sp. specimens from the Sidi Hakoma Member display microwear textures similar to modern dedicated browsing taxa, with high values for *Asfc* and *Tfv* and low values for *epLsar*. The overall microwear signature for these specimens is most similar to that of *Litocranius walleri*, the modern gerenuk, which consumes no monocots and only a small percentage of fruit. These specimens from Sidi Hakoma have significantly different microwear signatures from the specimens from the Denen Dora and Kada Hadar members. These later *Aepyceros* sp. specimens have texture values that fall clearly within the range of modern browser-grazer intermediates, a dietary category that includes the extant congener, *Aepyceros melampus*. Overall, the microwear evidence suggests that *Aepyceros* sp. consumed little to no grass during the Sidi Hakoma time period at Hadar, but began including grasses during the Denen Dora period, carrying that through the formation of the Kada Hadar Member.

<u>Alcelaphini</u>- The tribe Alcelaphini includes hartebeests, wildebeests, bonteboks and tsessebes. All modern alcelaphines are primarily classified as grazers and consume little to no fruit (Kingdon, 1997; Gagnon and Chew, 2000). The extant alcelaphine species included in the study are the tsessebe, *Damaliscuslunatus*, which occupies open savannas and floodplains and the bontebok, *Damaliscus pygargus*, commonly found in the coastal highveld (Kingdon, 1992a, b). These taxa are classified as an obligate grazer and variable grazer respectively. *Damaliscus*  *pygargus* is classified as an obligate grazer and consumes more dicots, but varies its diet on a seasonal basis. *Damaliscus lunatus* is categorized as a variable grazer, and while it consumes a wider variety of food types than the tsessebe, the bontebok has a diet that apparently does not vary seasonally or geographically.

Fossil alcelaphines are found in all three Hadar members and differences in microwear texture were found between the members. The fossil Alcelaphini specimens from the Sidi Hakoma and Denen Dora members all display indications of frequent browsing and are most similar to modern browser-grazer intermediates. The fossil specimens from these members have microwear signatures that cluster with the modern eland, a species with a diet consisting of nearly equal amounts of browse and graze, with some seasonal fruit consumption (Kingdon, 1982; Skinner and Smithers, 1990; Gagnon and Chew, 2000). Microwear evidence for more dedicated grazing by the Hadar alcelaphines doesn't occur until the Kada Hadar Member, for which the specimens have microwear signatures similar to those of extant variable grazers like *Damaliscus pygargus* and *Kobus ellipsiprymnus*. Browse makes up less than 15% of the diet for both of these modern taxa (Hofmann and Stewart, 1972; Skinner and Smithers, 1990; Gagnon and Chew, 2000).

<u>Antilopini</u>- The tribe Antilopini is composed of gazelles, gerenuks and springboks. Modern taxa subscribe to a wide range of diets, from the browsing *Litocranius walleri*, to the dedicated grazer, *Gazella granti* (Kingdon, 1997; Gagnon and Chew, 2000). The habitats occupied by antilopines are mostly semi-arid scrubland, savanna and even desert (Kingdon, 1992a, b). The Hadar fauna includes fossil species of *Gazella* sp., as well as unclassified Antilopini. Both of these taxa have been recovered from the Sidi Hakoma and Kada Hadar members and *Gazella* specimens have also been found in the Denen Dora Member.

The fossil *Gazella* sp. specimens from the three Hadar members are not significantly different and share microwear signatures most closely resembling those of modern browsergrazer intermediates. The fossil specimens have texture values that group with modern *Raphicerus sharpei*, the grysbok, and *Antidorcas marsupialis*, the springbok. Both of these extant taxa consume a variable diet that includes more than 55% browse and seasonal fruit consumption (Kingdon, 1982; Skinner and Smithers, 1990). Some subtle differences in dietary signature among the Hadar members are present, but the only statistically significant difference is in the heterogeneity variable. Unfortunately, the sample sizes for *Gazella* sp. are simply not large enough to draw many conclusions about potential change in diet over time at the site.

The Antilopini fossil specimens from the Sidi Hakoma and Kada Hadar members all have microwear signatures closely resembling those of the modern browsers *Litocranius walleri* and *Neotragus batesi*. While no significant differences were present between the representatives from these two members, the Sidi Hakoma specimens had higher values overall for *Asfc* and *Tfv*, as well as lower values for *epLsar* than those from Kada Hadar. When these specimens are compared to modern taxa, they fall well within the uppermost ranges of browsing taxa and overlap with the frugivorous bovids.

<u>Bovini</u>- The tribe Boviniis composed of large bovids that are classified as variable or obligate grazers (Kingdon, 1997; Gagnon and Chew, 2000). This includes domestic cattle, bison and the African buffalo, *Syncerus caffer*, which is part of the comparative database used here.Modern members of this tribe are highly successful grazers and found in habitats ranging from open grassland to more closed woodland (Kingdon, 1992a, b). Bovini specimens from Hadar include *Pelorovis* sp. and *Ugandax coryndonae*, as well as a number of specimens that cannot be classified past the level of tribe and are therefore considered separately here.

Bovinifossils have been recovered from all three Hadar members- *Ugandax coryndonae* in the Sidi Hakoma and Denen Dora members, *Pelorovis* in the Denen Dora and Kada Hadar members, and Bovini in the Kada Hadar Member. Differences in microwear textures are present both among and within the Hadar members.

Ugandax coryndonaeis the only Bovini taxon present in the Sidi Hakoma Member, and its microwear signature is comparable to those of modern browser-grazer intermediates, like *Raphicerus sharpei*. This suggests that *Ugandax* engaged in a diet very different from its closest living relative, *Syncerus caffer*, which is classified as variable grazer. *Ugandax coryndonae* and *Pelorovis* sp.are both present in the Denen Dora Member, and although the microwear evidence suggests that both were variable grazers, the two taxa fall at the opposite extremes of the category, suggesting some subtle differences in diet. For example, *Ugandax coryndonae* has texture values for complexity and textural fill volume that are at the highest end of the range for extant variable grazers, and actually overlap with browser-grazer intermediates, while *Pelorovis* has complexity and anisotropy values similar to modern obligate grazers. These results suggest that while grass was probably the staple resource for both taxa, browse may have been a more important component of the diet of *Ugandax coryndonae*.

The Kada Hadar assemblage includes both *Pelorovis* sp. and specimens classified only as Bovini. The microwear signature of the Bovini falls well within the range of modern obligate grazing taxa and clusters closely with *Redunca arundinum*, a species that consumes a diet of nearly all grass with no seasonal variation. *Pelorovis* sp., on the other hand, has a microwear texture nearly identical to that of *Syncerus caffer* and similar to fossils of the same genus from Denen Dora. While the sample for Bovini specimens is not large, the evidence presented here suggests an increase in grazing over time by these taxa at Hadar.

<u>Hippotragini</u>- Examples of modern hippotragines include the grazing oryx, a desertdweller, and sable antelope, commonly found in savanna woodlands and grasslands (Kingdon, 1992a, b; Gagnon and Chew, 2000). Members of the genus *Oryx* have been found at Hadar, along with fossil bovids identified only as hippotragines. Both Hippotragini and *Oryx* sp. have been recovered from all three hominin-bearing members at Hadar; however, specimens with well-preserved microwear were only identified from Denen Dora and Kada Hadar. The *Oryx* sp. specimens used in this study were all recovered from the Kada Hadar Member and these specimens have significantly higher anisotropy values than the Hippotragini specimens. This could be indicative of more dedicated grazing by *Oryx*. The specimens classified as Hippotragini have similar texture values to both variable and obligate grazers, suggesting the possibility of a seasonal shift in preferred resource availability. The *Oryx* sp. specimens clearly fall into the same range as obligate grazers including *Damaliscus lunatus* and *Redunca fulvorufula*, two taxa that consume almost no browse or fruit (Hofmann and Stewart, 1972).

<u>Neotragini</u>- Modern neotragines include the African dwarf antelopes including dik-diks, grysboks, oribis, and klipspringers. These taxa are primarily browsers and dietary generalists, consuming a wide variety of food types (Kingdon, 1997; Gagnon and Chew, 2000). Neotragines also occupy a variety of habitats, from moist closed forests to open savanna (Kingdon, 1992a, b). Fossil specimens belonging to the tribe Neotragini have been identified from all three homininbearing members at Hadar, but only specimens from the Sidi Hakoma and Kada Hadar members were suitable for microwear analysis.

The Sidi Hakoma and Denen Dora neotragin specimens are significantly different from one another, although both have microwear texture signatures similar to those of modern browsers. The microwear texture values of the Sidi Hakoma specimens are nearly identical to

those of the modern neotragin, *Neotragus batesi*, while those from Kada Hadar are most similar to the browsing gerenuk, *Litocranius walleri*. These similarities with different modern taxa may indicate subtle differences in browsing strategy. For example, fruit accounts for more than 20% of the diet of *Neotragus batesi*, while only making up about 5% of the diet of *Litocranius*.

<u>Reduncini</u>- Modern reduncines primarily inhabit watery environments including marshes, floodplains and swamps (Kingdon 1992a, b). The comparative database used in this study includes data on four members of this tribe: the waterbuck, *Kobus ellipsiprymnus*,the lechwe, *Kobus leche*, the southern reedbuck, *Redunca arundinum*, and the mountain reedbuck, *Redunca fulvorufula*. Nearly all members of Reduncini are classified as obligate grazers, with only *Kobus ellipsiprymnus*engaging in more variable grazing (Gagnon and Chew, 2000). Both *Kobus* sp. and Reduncini fossils have been found in all three hominin-bearing members at Hadar, however the only Reduncini specimens preserving well-preserved microwear were those from the Denen Dora Member.

While the differences between the *Kobus* fossils from the three members are only marginally significant (Fishers test, p-value=0.048), there is subtle variation among the specimens from each member. This becomes more apparent when the values are plotted in a multivariate space alongside the modern taxa (Figure 5). The microwear textures of the *Kobus* sp. specimens from Sidi Hakoma most closely resemble those of modern browser-grazer intermediates like *Raphicerus sharpei*, which consumes mostly dicots, while the Denen Dora specimens generally fall into a different space, overlapping variable grazing taxa like *Gazella granti*. The Kada Hadar *Kobus* sp. specimens fall on the border of obligate and variable grazing taxa and have a microwear signature that most closely resembles that of *Damaliscus pygargus*, a species that consumes primarily grass, with seasonal inclusion on dicots. This suggests that

browse and/or fruit may have always been an important component to the diet of *Kobus* sp. throughout the Hadar Formation, but that it may have become a seasonal component rather than a year-round staple by the time of Denen Dora and Kada Hadar deposition.

The Reduncini specimens from the Denen Dora Member have microwear textures consistent with those of modern obligate grazers. In fact, the taxon clusters most closely with the modern reduncines, *Redunca fulvorufula* and *Kobus leche*. This suggests that, at least during the Denen Dora period, Reduncinifavored a somewhat catholic diet primarily composed of grasses, with very occasional consumption of fruit or other dicots.

<u>Tragelaphini</u>- The spiral-horned antelopes include the genera *Tragelaphus* and *Taurotragus*. The tragelaphines recovered from Hadar have all been identified as members of the genus *Tragelaphus*. This genus also includes modern bovids representing nearly all dietary specializations, ranging from *Tragelaphus spekii*, a variable grazer, to *Tragelaphus euryceros*, a dedicated browser (Gagnon and Chew, 2000). All modern tragelaphines occupy forest habitats, although within this category, they range dramatically from open woodland to swamp forest to rainforest (Kingdon, 1992a, b).

*Tragelaphus* sp. has been identified in all three Hadar members, but only specimens from the Sidi Hakoma and Denen Dora members preserve suitable microwear for analysis. The specimens from Sidi Hakoma have a microwear texture signature similar to modern browsers, specifically the congener, *Tragelaphus euryceros*; while results for later Denen Dora specimens suggest a diet like that of extant browser-grazer intermediates. In fact, the microwear signature of the Denen Dora *Tragelaphus* sp. specimens is nearly identical to that of *Tragelaphus imberbis*, a species for which grass makes up more than one-third of its diet (Hofmann and Stewart, 1972). This suggests that graze may have become more important as a staple resource

for *Tragelaphus* at Hadar over time. It is worth noting, however, that modern tragelaphines engage in a wide variety of dietary strategies and that since the fossil taxa cannot be identified past the level of genus, there is a possibility that we are combining multiple taxa into a single sample.

#### Implications for paleoenvironment at Hadar

Dental microwear is a nongenetic signal that provides direct evidence of what an animal ate during its lifetime, rather than a prediction of what an animal was adapted to eat based on morphology. This is an important step in reconstructing diet because resource availability can change rapidly, but it can take many generations for morphology to follow suit. Therefore, microwear can be an excellent indicator of the types of resources available to local fauna. Dental microwear texture analysis of the Hadar bovids is an important step toward better understanding the local paleoenvironment throughout the occupation of *Australopithecus afarensis*.

Sidi Hakoma Member- The Sidi Hakoma Member was deposited between 3.43-3.26 mya and is composed of four submembers known as SH-1 through SH-4 (Campisano, 2007). Previous analyses of local fauna (Reed, 2008), as well as oxygen and carbon isotope studies (Hailemichael, 2000), have suggested that Hadar was made up of woodland and shrubland during this time, with low seasonal variation. This may not have been the case consistently, however, as palynological studies at the site have indicated that the local environment was a frequently changing mosaic (Bonnefille et al., 2004) during the time of Sidi Hakoma deposition.

Overall, the dietary reconstructions of the Sidi Hakoma taxa included in this study suggest a relatively closed habitat, or at least, regular access to regular fruit and browse. The microwear evidence presented here suggests that no obligate or variable grazers were present in

the assemblage and that the bovid taxa found in the Sidi Hakoma deposits primarily consumed browse. Additionally, there appears to have been less overall variation in the diets of the Sidi Hakoma bovids than in the Denen Dora or Kada Hadar members, where the bovids evince both dedicated grazing and browsing signatures. The absence of wear signatures similar to open country adapted grazers, along with the presence of so many browsing taxa, strongly suggests that the Sidi Hakoma paleoenvironment was more closed than that of the later Hadar members.

Denen Dora Member- The Denen Dora Member dates from 3.26-3.2 mya (Campisano, 2007), and is divided into three submembers, DD-1 through DD-3. Analysis of the fauna has suggested a local paleoenvironment comparable to that of modern mosaic woodlands that include both gallery forest and bushland (Reed, 2008). An overall increase in the number of alcelaphine bovids also suggests the presence of, or proximity to, grassland during this time period and this is also consistent with high levels of grass pollen collected in the lower strata of the member by Bonnefille et al. (2004). However, evidence of other habitat types are also present in the Denen Dora Member, including sedge and reed pollen (Bonnefille et al., 2004) and the presence of reduncin and aepycerotin bovids, all of which is suggestive of a browse-rich woodland environment.

The microwear textures of the Denen Dora bovids suggest a variety of dietary strategies, and are most similar to modern taxa with diets that depend on a substantial graze component. Most of the Denen Dora fossil taxa cluster with modern browser-grazer intermediates, variable grazers and/or obligate grazers. This is consistent with reconstructions of Hadar as a mosaic woodland habitat where both grasses and woody browse were available. The microwear data also suggest a shift in paleoenvironment between the deposition of Sidi Hakoma and Denen Dora. In contrast to the Sidi Hakoma Member, none of the Denen Dora bovids have microwear signatures

similar to those of modern dedicated browsers or frugivores. Additionally, the dietary signatures of several of the bovid taxa change between the two members. For example, Sidi Hakoma specimens of *Aepyceros* sp. and *Tragelaphus* sp. display microwear consistent with browsing, while Denen Dora members of the same taxa resemble modern browser-grazer intermediates that consume higher quantities of grass. *Kobus* sp. also shifts between these two members, going from a browser-grazer intermediate signature to one more consistent with variable grazing. All of these transitions suggest the availability of multiple resource types and access to both open and closed habitats.

<u>Kada Hadar Member</u>- The two collection units that make up the Kada Hadar Member were deposited between ~3.2-2.94 mya and are known as KH-1 and KH-2 (Campisano, 2007). Reed (2008) noted that the fauna from this member resembles the assemblages typical of modern open woodland/edaphic grassland environments such as in Amboseli National Park in Kenya. Antilopin and alcelaphin bovids are common in both of the submembers, indicating a drier and more open habitat (Vrba, 1980). However, pollen data suggests a humid woodland habitat (Bonnefille et al., 2004), although it has been suggested that this could be reflective of one of several types of closed physiognomies comprising the mosaic local environment and not an indicator of habitat for the whole site (Campisano and Reed, 2007).

The microwear evidence presented in the current study supports a more open habitat reconstruction for the Kada Hadar Member, although there is still evidence suggesting that local fauna had access to woodland resources. The majority of the bovid taxa have microwear textures consistent with an obligate or variable grazing strategy, although some browsing taxa are still present. This suggests the availability of a range of food types, including both significant graze and browse, and could indicate a mosaic environment including both grassland and woodland.
The difference in the paleoenvironmental signal between Denen Dora and Kada Hadar is not as pronounced as that between Sidi Hakoma and Denen Dora, but evidence of some change is present nonetheless. Bovids reconstructed as obligate grazers are more common in the Kada Hadar assemblages than those from Denen Dora, and while evidence of browse consumption is still present, it appears to have only been a primary resource for three of the ten taxa. There is also evidence to suggest that graze became more of a staple resource for some of the Kada Hadar bovids than it was for their predecessors. *Kobus* sp. specimens appear to transition from a variable grazing signature in the Denen Dora Member to an obligate grazing signature in the Kada Hadar Member, and both Alcelaphini and Hippotragini shift from browser-grazer intermediate to variable grazer. These changes in diet suggest that the Kada Hadar bovids had access to open habitats for grazing, although the presence of a few browsing taxa support mosaic reconstructions for this member.

In general, the microwear evidence is consistent with a gradual aridification trend at Hadar during the occupation of *A. afarensis*. This suggests that the local environment transitioned from closed woodland dominated by browsing bovids during the Sidi Hakoma Member, to more open, mosaic habitats with both woodland and grassland available to the local bovid populations by the time of the Kada Hadar deposition. It is necessary to note that because the sample sizes did not allow for comparison among submembers, changes in habitat availability within the members themselves cannot be commented on here. However, the differences in bovid diets reported between the Hadar members are sufficient to suggest that the overall environment changed considerably during the time in which *A. afarensis* inhabited the site.

#### Paleoenvironments of Australopithecus afarensis

The results of this study are generally consistent with previous reconstructions of Hadar as a mosaic of localized environments during the occupation of *Australopithecus afarensis*. In addition to Hadar, *A. afarensis* specimens have also been recovered from East African sites where the paleoenvironments appear to have ranged from open shrubland to closed woodland, with no evidence of specific habitat preference (Andrews, 1989, 2006; White et al., 1993; Reed, 1997). The fact that these early hominins have been found in such a variety of paleoenvironmental conditions has led to the suggestion that *A. afarensis* was ecologically plastic and could successfully inhabit a wide variety of habitat types (White et al., 1993).

# Conclusions

The dental microwear of the Hadar bovids indicates a wide range of dietary variation, and by extension, food availability. This suggests that local habitat at Hadar were not entirely open or closed, but instead included a variety of habitat types that changed during the occupation of *A*. *afarensis*. Dental microwear data agree with other lines of evidence that these may have ranged from intermediate cover physiognomies including bushland and woodland, to more open habitats like shrubland and edaphic grasslands. The dental microwear signatures of the fossil bovids indicate a gradual transition from the more closed habitats comprising the Sidi Hakoma Member to more open environments by the time of Kada Hadar deposition.

This research contributes to the large body of information on the paleoecology at Hadar and suggests that dental microwear texture analysis about a valuable additional proxy for the diet and habitat preferences of fossil mammals.

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# TABLES

**Table 1**. Descriptions of habitat categories discussed in this paper. (White, 1983; Reed, 1997, 1998).

Habitat Types	Characteristics
FORESTS	
Lowland rainforest	tall columnar with multi-structured canopies and a shrub layer and sparse ground cover
Montane forest	dominated by deciduous and evergreen trees, high annual rainfall
Dry or seasonal forest	single closed canopy forest, prolonged dry seasons
SAVANNAS	
Open woodland	deciduous trees comprise 20-25% of vegetation, grass and herb ground cover
Medium density woodland	deciduous trees comprise 30-45% of vegetation, grass and herb ground cover
Closed woodland	deciduous trees comprise more than 50% of vegetation, grass and herb ground cover
Shrubland or scrub woodland	shrubs constitute more than 20% of ground cover, poor quality grass, very few trees
Secondary or open grassland	dominated by grasses and herbs, with widely scattered trees and shrubs
Edaphic grassland or wetland	seasonally or permanently waterlogged soil, aquatic grasses
BUSHLAND	
Bushland	bushes constitute 40% of ground cover, also some trees, few grasses
DESERT	

TAXON	п	ASFC	epLSAR	SMC	TFV	HASFC9	HASFC81
SIDI			-				
HAKOMA							
Aepyceros	3	3.526147	0.0024	0.29761	8752.488	0.52981	0.94515
Alcelaphini	4	2.239167	0.003966	0.405793	6598.503	0.524273	0.884276
Antilopini	5	4.817109	0.003345	0.631563	11977.21	0.661216	1.155878
Gazella	2	2.288603	0.004937	0.6128	6071.115	0.498515	0.729166
Kobus	3	2.491008	0.002117	0.972458	4184.89	0.326508	0.837516
Neotragus	3	4.265469	0.001865	0.686543	9763.845	0.693515	0.953135
Tragelaphus	5	3.076213	0.003149	0.644873	8062.735	0.544011	0.855305
Ugandax	6	2.987175	0.003452	0.453171	5578.615	0.579619	0.840578
DENEN							
DORA							
Aepyceros	4	1.74854	0.00385	0.504261	5371.542	0.426949	0.750294
Alcelaphini	10	2.237113	0.004795	0.425972	6239.596	0.500645	0.924125
Gazella	4	2.791705	0.005235	0.505698	5333.485	0.4076	0.807513
Hippotragini	3	1.978451	0.005978	0.436846	3984.685	0.450178	0.813547
Kobus	9	1.598745	0.003698	1.187655	4462.355	0.419865	0.861324
Pelorovis	2	1.232126	0.006082	0.7799	2328.855	0.506156	0.855029
Reduncini	14	0.991558	0.006	1.215434	2593.004	0.444572	0.815354
Trgelaphus	10	1.963995	0.004993	0.855291	6539.146	0.482329	0.720312
Ugandax	9	2.068764	0.004933	0.502647	4267.688	0.513265	0.894055
KADA							
HADAR							
Aepyceros	5	1.484654	0.0041	0.26845	5972.351	0.386515	0.85441
Alcelaphini	4	1.485766	0.007602	0.322352	3331.107	0.400018	0.793935
Antilopini	5	3.587647	0.002977	0.618765	9597.235	0.698785	1.064988
Bovini	5	0.926677	0.00698	1.066665	3564.102	0.427961	0.724178
Gazella	4	2.281361	0.005181	0.534876	4276.6	0.451222	0.86058
Hippotragini	2	1.315717	0.00493	0.40046	2626.581	0.404886	0.921388
Kobus	3	0.987651	0.002976	1.135454	1103.832	0.432497	0.76133
Neotragus	4	2.698746	0.001985	0.624687	10268.15	0.626487	0.988435
Oryx	3	0.998747	0.007698	1.264877	1168.655	0.598465	0.975454
Pelorovis	4	1.315469	0.006076	0.753212	2677.005	0.427806	0.753905

**Table 2**.Sample sizes and median texture values for fossil taxa by member.

**Table 3**.Sample size, diet and habitat preferences of the extant taxa. The habitat categories are those of White, 1983 and preferences by taxon are summarized in Kingdon, 1982 and 1997. The diet categories are those assigned by Gagnon and Chew, 2000.

TAXON	n	PREFERRED HABITAT
BROWSERS		
Litocranius walleri	25	scrubland
Neotragus batesi	22	bushland and dry forest
Sylvicapra grimmia	25	scrubland and woodland
Tragelaphus euryceros	22	dense forest
OBLIGATE GRAZERS		
Damaliscus lunatus	22	open woodland and grassland
Redunca fulvorufula	24	forest and medium-closed woodland
Kobus leche	24	edaphic grassland
Redunca arundinum	25	edaphic grassland and woodland
VARIABLE GRAZERS		
Damaliscus pygargus	22	open grassland
Gazella granti	25	shrubland and open grassland
Hippotragus niger	22	open-medium density woodland
Kobus ellipsiprymnus	23	scrubland and edaphic grassland
Syncerus caffer	22	edaphic grassland and dry forest
Tragelaphus spekii	22	edaphic grassland
BROWSER-GRAZER		
INTERMEDIATES		
Aepyceros melampus	20	bushland and open woodland
Antidorcas marsupialis	21	open grassland, shrubland and desert
Raphicerus campestris	21	open woodland, shrubland and grassland
Taurotragus oryx	21	open woodland, shrubland and grassland
		bushland and open-medium density
Tragelaphus imberbis	23	woodland
Raphicerus sharpei	25	shrubland and open grassland
GENERALISTS	1	
Oreotragus oreotragus	24	open-medium density woodland
Tragelaphus angasi	22	dense forest
Tragelaphus strepsiceros	25	bushland and closed woodland
FRUGIVORES		
Cephalophus sylvicultor	25	rainforest
Philantomba monticola	21	rainforest

	SS	df	MS	F	р
Asfc	122927.657	27	4552.876	17.148	0.000
epLsar	110533.863	27	4093.847	10.267	0.000
Smc	136819.248	27	5067.380	43.633	0.000
Tfv	128649.096	27	4764.781	23.359	0.000
HAsfc9	88424.771	27	3274.992	5.145	0.000
HAsfc81	78937.456	27	2923.609	3.959	0.000

**Table 4**. Statistical results of the general linear estimate model.

 Table 5.Nested analysis of variance of fossil taxa within member.

SIDI					
HAKOMA					
	SS	df	MS	F	р
Asfc	5487.927	7	783.990	5.087	0.002
epLsar	7741.130	7	1105.876	5.760	0.001
Smc	16921.250	7	2417.321	44.103	0.000
Tfv	12481.102	7	1783.015	15.445	0.000
HAsfc9	13690.133	7	1821.029	5.460	0.003
HAsfc81	13914.774	7	1987.825	2.438	0.058
DENEN					
DORA					
Asfc	34597.524	9	3844.169	11.338	0.000
epLsar	23746.513	9	2638.501	4.710	0.000
Smc	64426.516	9	7158.502	44.785	0.00
Tfv	46997.090	9	5221.899	21.496	0.00
HAsfc9	19571.584	9	4597.251	9.784	0.007
HAsfc81	33479.190	9	3719.910	5.113	0.000
KADA					
HADAR					
Asfc	34658.500	9	3850.944	22.292	0.000
epLsar	41285.692	9	4587.299	28.828	0.000
Smc	44858.833	9	4984.315	88.385	0.000
Tfv	41061.588	9	4562.399	24.952	0.000
HAsfc9	34268.015	9	3948.264	16.264	0.004
HAsfc81	28289.500	9	3143.278	4.521	0.002

		d						d			
	SS	f	MS	F	р		SS	f	MS	F	р
AEPYCER											
OS						KOBUS					
	8326.8		4163.	30.4	0.0		3994.		1997.	3.10	0.0
Asfc	33	2	417	29	00	Asfc	861	2	431	4	94
	2831.3		1415.	8.07	0.0		2352.		1176.	2.42	0.1
epLsar	63	2	681	7	10	epLsar	250	2	125	2	44
	3111.4		1555.	23.3	0.0		111.1		55.55	0.32	0.7
Smc	50	2	725	05	00	Smc	11	2	6	5	31
	2698.6		1349.	14.5	0.0		1439.		719.6	3.38	0.0
Tfv	75	2	338	43	02	Tfv	361	2	81	6	80
	7029.6		3514.	19.6	0.0		2041.		1020.	1.98	0.1
HAsfc9	33	2	817	16	01	HAsfc9	944	2	972	5	93
	5408.1		2704.	2.59	0.1		1372.		686.4	0.64	0.5
HAsfc81	33	2	067	6	29	HAsfc81	861	2	31	0	50
ALCELAP						NEOTRAG					
HINI						US					
	4296.4		2148.	6.71	0.0		456.3		456.3	48.8	0.0
Asfc	44	2	222	2	08	Asfc	33	1	33	93	20
	11501.		5750.	9.39	0.0		16.33		16.33	0.12	0.7
epLsar	594	2	797	7	02	epLsar	3	1	3	2	61
	901.15		450.5	3.60	0.0				40.33	2.32	0.2
Smc	0	2	75	5	53	Smc	4.033	1	3	7	67
	6766.0		3383.	9.08	0.0					0.00	0.9
Tfv	65	2	033	6	03	Tfv	0.333	1	0.333	7	40
	7561.1		3780.	7.18	0.0					0.00	0.9
HAsfc9	94	2	597	5	06	HAsfc9	0.333	1	0.333	2	69
	5635.0		2817.	6.67	0.0		70.08		70.08	0.25	0.6
HAsfc81	00	2	500	1	08	HAsfc81	3	1	3	5	63
ANTILOPI						PELOROV					
NI						IS					
	138.37		69.18	1.85	0.2					0.00	1.0
Asfc	5	2	7	5	50	Asfc	0.000	1	0.000	0	00
			10.35	0.03	0.9		16.00		16.00	0.08	0.7
epLsar	20.700	2	0	0	70	epLsar	0	1	0	7	96
				0.01	0.9	_				0.01	0.9
Smc	1.800	2	0.900	3	87	Smc	1.000	0	0.000	1	26
	106.57		53.28	2.53	0.1		25.00		25.00	0.24	0.6
Tfv	5	2	7	0	74	Tfv	0	1	0	0	72
~	456.67		228.3	4.03	0.0		1681.		1981.	6.65	0.1
HAsfc9	5	2	38	1	91	HAsfc9	000	1	000	7	23
	258.80		129.4	0.30	0.7		1260.		1260.	2.79	0.2
HAsfc81	0	2	00	0	53	HAsfc81	250	1	250	3	37

 Table 6.Nested analysis of variance among members by taxon.

						TRAGELA					
GAZELLA						 PHUS					
	490.08		245.0	1.17	0.4		5713.		5713.	14.1	0.0
Asfc	3	2	42	0	21	Asfc	200	1	200	17	02
	112.33		56.16	0.07	0.9		7776.		7776.	12.8	0.0
epLsar	3	2	7	7	27	epLsar	300	1	300	19	03
	330.33		165.1	4.11	0.1		1598.		1598.	15.5	0.0
Smc	3	2	67	2	38	Smc	700	1	700	69	02
	1334.3		667.1	3.39	0.1		1203.		1203.	4.10	0.0
Tfv	33	2	67	0	70	Tfv	333	1	333	4	64
	1702.5		851.2	24.8	0.0		1360.		1360.	1.25	0.2
HAsfc9	83	2	92	86	14	HAsfc9	133	1	133	6	83
	2071.0		1035.	1.10	0.4		6720.		6720.	11.9	0.0
HAsfc81	00	2	500	1	38	HAsfc81	033	1	033	04	04
HIPPOTR						UGANDA					
AGINI						Х					
	952.03		952.0	8.47	0.0		3324.		3324.	15.7	0.0
Asfc	3	1	33	1	62	Asfc	544	1	544	94	02
	572.03		572.0	6.72	0.0		4579.		4579.	17.5	0.0
epLsar	3	1	33	5	81	epLsar	600	1	600	55	01
			38.53	0.33	0.6		208.5		208.5	2.09	0.1
Smc	38.533	1	3	2	05	Smc	44	1	44	8	71
	264.03		264.2	1.34	0.3		1232.		1232.	6.61	0.0
Tfv	3	1	33	9	29	Tfv	100	1	100	1	23
-	388.80		388.8	0.34	0.5		1020.		1020.	3.15	0.0
HAsfc9	0	1	00	7	97	HAsfc9	100	1	100	8	99
-	1642.8		1642.	5.38	0.1	-	298.8		298.8	0.53	0.4
HAsfc81	00	1	800	0	03	HAsfc81	44	1	44	2	79

# Figure Legends

Figure 1. Occlusal view of bovid M2, illustrating the shearing facets where microwear was examined (Illustration courtesy of Gildas Merceron). A = left  $m^2$ , B= right  $m_2$ 



Figure 2.Examples of dental microwear surfaces for extant bovids from different dietary categories. These images are photosimulations based on data collected using the confocal imaging profiler, and each represents an area of 276 x 204 μm. A) obligate grazer; B) variable grazer; C) browser-grazer intermediate; D) generalist; E) browser; and F) frugivore.



Figure 3.Examples of dental microwear surfaces for the Hadar bovids by member. These images are photosimulations based on data collected using the confocal imaging profiler, and each represents an area of 138 x 102 μm. Sidi Hakoma: A) *Aepyceros*; B) Alcelaphini; C) Antilopini;
D) *Gazella*; E) *Kobus*; F) *Neotragus*; G) *Tragelaphus*; H) *Ugandax*. Denen Dora: A) *Aepyceros*;
B) Alcelaphini; C) *Gazella*; D) Hippotragini; E) *Kobus*; F) *Pelorovis*; G) Reduncini; H) *Tragelaphus*; I) *Ugandax*. Kada Hadar: A) *Aepyceros*; B) Alcelaphini; C) Antilopini; D) Bovini;
E) *Gazella*; F) Hippotragini; G) *Kobus*; H) *Neotragus*; I) *Oryx*; J) *Pelorovis*.

Sidi Hakoma



Denen Dora



#### Kada Hadar



Figure 4.Bivariate plot of M<sub>1</sub> shear facet microwear texture anistropy and complexity for extant taxa by dietary category. Data are plotted for individuals with species as indicated by the markers.



Asfc



**Figure 5.** Bivariate plot of M<sub>1</sub> shear facet microwear texture anistropy and complexity for Hadar fossil taxa by member.

ASFC



# Figure 6. Box and whisker plots

Chapter Four: Paleoenvironmental change in Pliocene eastern Africa as inferred from dental microwear texture analysis of fossil Bovidae

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#### Abstract

Environmental hypotheses of Pliocene hominin evolution state that key ecological adaptations of early hominins are directly linked to shifts in local or regional climate. In order to determine relationships between climate change and human evolution, it is first necessary to reconstruct the habitats in which early hominins lived. This is especially important for Pliocene sites where increasing bipedality has been linked to a shift from closed woodland forests to more open, arid habitats. Many techniques have been used to refine our understanding of the Pliocene paleoenvironments of eastern Africa; however these have not led to consensus reconstructions. Here, I bring a new, independent dataset for the inference of diet, and by extension habitats, of individual bovids in the days before death. This study applies dental microwear texture analysis to reconstruct the diet, and therefore ecological contexts, of specimens from Kanapoi, Allia Bay and Laetoli. The microwear signatures of the fossils are interpreted using a comparative database of 25 extant species of African Bovidae and are compared to previously published data on the bovids from the Hadar hominin site in Ethiopia. Results indicate that the fossil bovid assemblages from all four sites included both browsing and grazing taxa, suggesting access to a wide variety of resources. The earlier sites include higher proportions of browsing taxa, suggesting more closed habitats on average. In general, the results presented here indicate that all of these sites associated with the early australopiths were mosaic and primarily composed of semi-closed habitats like woodland and bushland, with incursions of open savanna grassland.

# Introduction

Many researchers have argued that climate change was a motive force for human evolution and that key ecological adaptations of australopiths were directly linked to shifts in local or regional climate (for examples, see Dart, 1925; Robinson, 1963; Howell, 1978; Vrba, 1985, 1988, 1992, 1995a, 1995b, 2000; Stanley, 1992; deMenocal, 1995; Hill, 1995; Potts, 1996, 1998, 2007; Bobe et al., 2002; Trauth et al., 2005). The "savanna hypothesis" (Dart, 1925, 1953; Bartholomew and Birdsell, 1953; Robinson, 1954; Washburn, 1960; Jolly, 1970; Laporte and Zihlman, 1983) and "turnover pulse hypothesis" (Vrba et al., 1989; Vrba, 1995a,b) both suggest that a shift toward drier, more open settings led to adaptations for bipedality and the consumption of savanna resources, including large grazing mammals. However, recent studies of marine sediment cores suggest that widespread C4 grasslands did not develop in eastern Africa until after 3.0 mya and therefore, would not have influenced the habitats of australopiths or earlier probable hominins (deMenocal, 1995; Leakey et al., 1995; White et al., 1994, 2009). Recent paleoclimatological models instead suggest that a wide range of habitats were available to early hominins, including closed and open woodland, gallery forest, bushland, and wet and dry grassland habitats (Kingston et al., 1994; Kappelman et al., 1997; Leakey et al., 1995; Reed, 1997; Wynn, 2000; Kingston and Harrison, 2001; Schoeninger et al., 2003; Bonnefille et al., 2004; Campisano and Feibel, 2007).

The Pliocene australopiths from eastern Africa, *Australopithecus anamensis* and *A. afarensis*, form a putative single anagenetic lineage suggesting changing adaptations over time (Kimbel et al., 2006). *Australopithecus anamensis* is best known from the sites of Kanapoi and Allia Bay in Kenya, and the largest collections of *A. afarensis* fossils have been found at Laetoli in Tanzania and Hadar in Ethiopia. The middle Pliocene dates associated with these

chronologically distinct sites cover a span from 4.1-3.0 mya and represent a period of changing, variable habitats (deMenocal, 1995; Leakey et al., 1995). The proposed mosaic nature of these habitats has led to the suggestion that concomitant variation in available foods and diets might be considered a possible driving force behind evolutionary changes within the hominin lineage (Teaford and Ungar, 2000; Ungar, 2004; Reed and Fish, 2005; Kimbel et al., 2006). *Australopithecis anamensis* and *A. afarensis* are characterized by morphological adaptations for at least facultative bipedalism and more robust dentition than earlier taxa like *Ardipithecus ramidus* (White et al., 1994). The mosaic nature of reconstructed Pliocene sites accords well with the locomotor adaptations of the australopiths. *Australopithecus anamensis* and *A. afarensis* are characterized by postcranial adaptations allowing for both terrestrial bipedality and arboreal climbing (e.g., Rodman and McHenry, 1980; Johanson et al. 1987). This unique locomotor pattern has been hypothesized to be the result of an environmental shift towards mosaic habitats, where the ability to walk bipedally and yet retain climbing proficiency would have been beneficial.

The reconstruction of past environments often centers on the paleoecology of a single taxon or several taxonomic groups (e.g., Robinson, 1963; Vrba, 1974, 1975, 1980, 1988; Kay, 1975, 1978; Grine, 1981; Stern and Susman, 1983; Kappelman, 1988; Benefit and McCrossin, 1990, Ciochon, 1990; Spencer, 1995; Lewis, 1997). Paleoecological research on these fossil faunas has involved two principal approaches. Some have focused on the ecomorphology, or relationship between morphological adaptations and ecological preferences, of a single taxon (Vrba, 1974, 1975, 1980, 1988; 1995b; Grine, 1981; Kappelman, 1988; Spencer, 1995). Others have concentrated on analyses of fossil communities as a whole by using entire faunal assemblages from a fossil site to identify major locomotor and trophic adaptations, thus implying

local environmental conditions (Dodd and Stanton, 1990; Behrensmeyer et al., 1997; Reed and Rector, 2006; Reed, 1998, 2007, 2008). Fossil bovids are frequently used as paleoenvironmental indicator species because they are ubiquitous at fossil sites and modern taxa can be placed into discrete dietary categories that reflect habitat preference (Kappleman, 1984; Vrba, 1980, 1982, 1985; Shipman and Harris, 1988; Harris, 1991; Plummer and Bishop, 1994; Spencer, 1997; Sponheimer et al., 1999).

In this paper, I evaluate previous reconstructions of paleohabitats associated with *A*. *anamensis* and *A. afarensis* by reconstructing the diets of the fossil bovids from the Pliocene sites of Kanapoi, Allia Bay and Laetoli in Tanzania using dental microwear texture analysis. This technique provides a measurement-error-free method for studying microwear, incorporating automated data collection and three-dimensional surface analysis. The fossil bovid data are considered in relation to an established microwear texture database featuring 25 extant African bovid taxa with known diet and habitat preferences (Scott, in press), as well as previously described fossil bovid taxa from the Sidi Hakoma, Denen Dora and Kada Hadar members of the Hadar Formation in Ethiopia, associated with *A. afarensis* (Scott, in review). This paper also addresses implications of inferred fossil bovid diets for paleoenvironments at the sites. Finally, I discuss how the ecological settings at these important sites may have changed over the temporal span of *A. anamensis* and *A. afarensis*.

#### Bovid paleoecology and habitat

Studies of faunal assemblages as paleoenvironmental proxies for a fossil site typically apply the principle of taxonomic uniformitarianism, which is based on the assumption that fossil species had the same ecological preferences as their closest living relatives. Although this assumption is rarely questioned for recent faunas, its utility for reconstructing the paleoecology

of long extinct species is uncertain. The primary problems with taxonomic uniformitarianism are that it is not applicable to fossil taxa with few or no living relatives, and that it assumes that groups remain constant in their ecological preferences over long periods of time. While the concept of taxonomic uniformitarianism is relatively straightforward, critics have questioned many of its inherent assumptions (Solounias et al., 1988; Reed, 1996; Sponheimer et al., 1999; Schubert et al., 2006). For example, Sponheimer et al. (1999) demonstrated in a study of fossil bovid isotopes and ecomorphology that half of the included taxa had food preferences that differed from their assumed diets based on taxonomic uniformitarianism. While *Aepyceros* sp. and *Gazella vanhoepeni* were both assumed to be mixed/seasonal feeders like their closest living relatives, for example, isotopic signatures and ecomorphological data suggested that both were obligate browsers, and showed no evidence of C4 grass consumption. Another diet proxy that considers actual diets of individuals would certainly prove valuable for interpreting these discrepancies, and that is where dental microwear comes in.

Diet is the most direct way that an organism interacts with its environment. Attempts to reconstruct diets of fossil forms have included the use of tooth size, shape and structure. However these are all genetic signals that reflect only what a species capable of eating, not what individuals actually ate on a daily basis. The diet of an individual may vary over its lifetime, but it can take many generations for the morphology of species to change in response to selective pressures. Additionally, adaptation does not always imply food preference and less than optimal solutions given phylogenetic inertia may complicate interpretation. In order to reconstruct what individuals actually ate in the past, a non-genetic signal is needed. Dental microwear is direct line evidence of the foods eaten and, by extension, of the environments inhabited by extinct bovids. It can therefore be a valuable tool for habitat reconstruction.

Bovid diets have traditionally been classified as grazing, browsing and mixed feeding. These categories refer to the proportions of monocotyledons (grasses, sedges and roots) and dicotyledons (fruits, seeds, flowers, buds, leaves, tubers and shoots) included in their diets. Grazers consume more than 90% monocotyledons and inhabit open, arid environments (Fortelius and Solounias, 2000). Modern grazing bovids include Alcelaphini (wildebeest), Antilopini (gazelles), and Hippotragini (grazing antelopes) (Kingdon, 1982). Browsers eat more than 90% dicotyledons and live in closed, moist habitats like woodlands and forests (Fortelius and Solounias, 2000). Examples of modern browsing taxa include Aepycerotini (impala), Tragelaphini (kudu, eland, bongo, nyala and sitatunga), Reduncini (reedbucks and lechwe) and Bovini (buffalo and their relatives) (Kingdon, 1982). Finally, species classified as mixed feeders alternate between grazing and browsing and tend to inhabit seasonal and/or mosaic environments (Fortelius and Solounias, 2000). Extant mixed feeders include Cephalophini (duikers), Neotragini (dwarf antelope), Caprini (sheep and goats), and Boselaphini (four-horned antelope) (Kingdon, 1982). It should be noted, however, that actual bovid diets are often more complex than these categories suggest, and many intermediate classifications have been proposed (Gagnon and Chew, 2000). Even given the over-simplicity of the grazer-browser-mixed feeder categories, they have been proven useful in interpreting the diets and associated habitats of fossil bovids (Fortelius and Solounias, 2000; Merceron et al., 2004a, 2004b).

Differences in the dental microwear of grazing and browsing ungulates have been noted in previous studies of extant and fossil species (for examples, see Solounias et al., 1988; Solounias and Moelleken, 1993; Merceron and Ungar, 2005; Merceron et al., 2005; Schubert et al., 2006; Ungar et al., 2007), with browsers having more pits and grazers having more scratches. Mixed-feeding taxa tend to have intermediate microwear signatures, or patterns overlapping with

browsers and grazers. This has important implications for reconstructing paleoenvironments, as while the connection between diet and habitat are not exact, grazing and browsing microwear signatures do imply the availability of grasses and woody plants respectively.

### Hominin paleoenvironments in eastern Africa

Many techniques have been used to refine our understanding of the paleoenvironments of eastern Africa; however these have not led to consensus reconstructions. At Kanapoi, ecological diversity analysis indicates that at least part of the site was composed of closed woodland forest; however, taxonomic uniformitarianism of bovid taxa suggests a dry arid habitat. Similarly contradictory reconstructions exist for Allia Bay, with paleosol analysis and palynology suggesting a mosaic habitat dominated by savanna, and taxonomic uniformitarianism of faunal assemblages suggesting an environment composed of gallery forest, open woodland, floodplains and edaphic grasses. The Laetoli faunal assemblages have also led to varying reconstructions, with some suggesting habitats as disparate as open grassland and closed woodland. Here I test these opposing hypotheses using a new, independent dataset for the inference of diet by extension habitats of actual individuals in the days before death.

Although the link between dental microwear and paleoenvironment has been explored (Merceron et al., 2004a, 2005; Schubert et al., 2006; Ungar et al., 2007), few studies have used dental microwear to resolve conflicting interpretations of paleohabitat. This is surprising given the recognition of bovids as excellent paleoenvironmental indicators (Vrba, 1980, 1985; Kappelman, 1984; Plummer and Bishop, 1994; Spencer, 1997; Sponheimer et al., 1999). Due to the prevalence of fossil bovids at early australopith sites in eastern Africa, available bovid specimens from the sites of Kanapoi, Allia Bay and Laetoli are the focus of this study.

#### **Materials and Methods**

The fossil specimens used in this study are housed at the National Museums of Ethiopia (NME), Tanzania (NMT) and Kenya (NMK). All available maxillary and mandibular first and second molars were examined, as previous studies have not revealed significant differences in the microwear signals preserved on these teeth (Merceron et al., 2004a, 2004b). The teeth were assessed using the criteria of Teaford (1988) and King et al. (1999). The majority of fossil specimens examined were not suitable for microwear analysis, which is unsurprising given that bovid molars have thin enamel bands that are frequently damaged or broken. A total of 15 specimens from Allia Bay, 35 specimens from Kanapoi and 49 specimens from Laetoli possessed unobscured antemortem microwear and were included in the fossil sample, along with 124 specimens from Hadar that have been previously described (Scott, submitted). Table 2 details the fossil taxa and sample sizes. Only one tooth per individual was included to avoid bias in the sample.

This study uses a large microwear texture database of extant African bovids, previously published by Scott (in press) for comparison with the fossil taxa. This database includes 575 specimens representing 25 modern bovid taxa with well-understood diets and habitats. Original specimens are housed at the American Museum of Natural History (AMNH), New York; the Field Museum (FMNH), Chicago; the Smithsonian National Museum of Natural History (NMNH), Washington D.C.; and the Royal Museum of Central Africa (RMCA), Tervuren, Belgium. All are wild-shot specimens with known provenience. To ensure that the dietary reconstructions of the fossils are as precise as possible, extant bovids from all six diet categories of Gagnon and Chew (2000) were included in the comparative database. Particular emphasis was placed on the selection of taxa that engage in various levels of mixed feeding, as mosaic habitats

have been suggested for all four fossil sites included in this study. The modern taxa considered here are equally variable in habitat preference, ranging from open-country grazers (e.g., *Damaliscus lunatus, Redunca arundinum*) to closed-forest frugivores and browsers (eg. *Cephalophus sylvicultor, Neotragus batesi*).

The extant taxa included in the comparative database as listed in Table 1, along with their dietary category as assigned by Gagnon and Chew (2000). This system for classifying diet was selected because it expands on the traditional browser-grazer-mixed feeder trichotomy to include multiple levels of mixed feeding. As in traditional classification systems for bovids, the dietary categories used here are determined by the approximate percentages of monocotyledons and dicotyledons consumed during observation in the wild, although fruit is considered separately from other dicotyledons. When applied to fossil taxa, the incorporation of these various levels of mixed feeding improves the overall resolution of dietary reconstruction and the usefulness of microwear texture data as a proxy for habitat preference.

Gagnon and Chew (2000) recognized six dietary categories, defined by the amount of monocotyledons and dicotyledons included in the diet, with fruit considered separately. Obligate grazers have uniform diets consisting of at least 90% monocotyledons. These taxa typically occupy more open habitats like grasslands and savanna. Browsers consume more than 70% dicotyledons, while frugivores restrict at least 70% of their diet to fruit. Bovids belonging to these categories are usually found in closed or semi-closed habitats like forest or woodland. Variable grazing taxa can vary diets both seasonally and geographically, but always include 60-90% monocotyledons. Browser-grazer intermediates also have variable diets, but regularly consume 30-70% monocotyledons and dicotyledons, always including some fruit. Generalist bovids have more catholic diets that consist of more than 20% of each of the three food types.

Bovids that consume a wide variety of resources are frequently found in environments that consist of multiple habitat types.

The molding and casting procedures used in this study are those of Ungar (1996). As is standard in studies of bovid dental microwear, data were collected from the disto-buccal enamel band of the mesial cuspid of  $M1_{1-2}$  and the mesio-buccal enamel band of the mesial cusp of  $M^{1-2}$  (Janis, 1990; Merceron, 2005; Schubert, 2006; Ungar et al., 2007), as illustrated in Figure 1. The teeth were cleaned with acetone-soaked cotton swabs and then molded with President's Jet regular body polyvinylsiloxane dental impression material (Coltène-Whaledent Corp.). The casts were poured using Epotek 301 high-resolution epoxy resin and hardener (Epoxy Technologies Corp.). This process has been demonstrated to produce surface replicas that are accurate to a fraction of a micron (Beynon, 1987; Teaford and Oyen, 1989).

The casts were scanned using a Sensofar PL $\mu$  white-light confocal profiler (Solarius Development Inc., Sunnyvale, CA). This instrument uses 3-D point clouds to create digital elevation models of microwear surfaces. Each specimen was scanned with a 100x objective to generate a point cloud with a lateral sampling interval of 0.18  $\mu$ m, a vertical resolution of 0.005  $\mu$ m, and a field of view of 102 x 138  $\mu$ m. Four adjoining scans were collected for each specimen, for a total work envelope of 204 x 276  $\mu$ m. Solarmap Universal software (Solarius Development Inc., Sunnyvale, CA) was then used to level each scan. Dust particles and other adherents were removed from the scans using the erase and/or thresholding functions in Solarmap. The resulting point clouds were analyzed with Sfrax and Toothfrax scale-sensitive fractal analysis software (Surfract Corp, Worcester, MA.).

Scale-sensitive fractal analysis is based on the principle that apparent surface textures change with scale of observation. A surface that appears smooth when viewed at a coarse scale

may appear quite rough when examined at a fine scale. Apparent surface texture can be quantitatively characterized using several texture variables. Data were generated for five of these variables previously identified as informative for dietary reconstruction: area-scale fractal complexity (*Asfc*), length-scale anisotropy of relief (*epLsar*), scale of maximum complexity (*Smc*), textural fill volume (*Tfv*) and heterogeneity of area-scale fractal complexity (*HAsfc*). Specific details concerning these variables are well-documented in the literature (Ungar et al., 2003, 2007, 2008; Scott et al., 2005, 2006).

In general, taxa that consume hard or brittle foods tend to hard higher values for Asfc, Tfv and HAsfc as well as lower values for epLsar and Smc than do closely-related species that consume soft resources (Scott et al., 2006; Ungar et al., 2007, 2010; Prideaux et al., 2009; Schubert et al., 2010, Scott, in press; submitted). Complexity is a measure of surface roughness; and individuals with microwear surfaces dominated by pits will have high values for this variable. Anisotropy measures tendency toward directionality of surface features. Microwear surfaces dominated parallel striations should have high values for anisotropy. Previous studies have suggested that the range of scales at which complexity is calculated may be informative (Scott et al., 2005, 2006) and those scales yield the Smc variable. Higher Smc values correspond with fewer small features. Texture fill volumeis a three-dimensional measure of feature size, so surfaces with higher values for this variable tend to have many features in the 2-10 µm diameter range. Finally, *HAsfc* measures the overall heterogeneity of the surface by dividing each scan area into progressively smaller subsections and comparing the complexity of each. Two heterogeneity variables that have been used for distinguishing extant taxa with known differences in diet are *HAsfc*<sub>9</sub>, which divides in surface into a 3 x 3 gridand *HAsfc*<sub>81</sub>, which

divides the surface into a 9 x 9 grid. Individuals with features that vary across a surface typically have higher values for this variable than more surface homogeneity.

#### **Statistical Analyses**

Statistical analyses of the baseline sample have been previously published (Scott, in press), so the analyses here were focused on determining the extent of dietary variation found both within and among the fossil bovids at the four sites. The three Hadar submembers were considered independently since previous analysis has determined that the paleoenvironmental signals between them vary significantly (Scott, in preparation). Five variables showed significance in the overall model and included in the remaining statistical analyses: 1) *Asfc*,2) *epLsar*, 3) *Smc*, 4) *Tfv*, and 5) *HAsfc*<sub>9</sub>.

# **Differences among fossil sites**

The fossil bovid microwear data were analyzed using a general linear model to determine the extent of differences in microwear among the taxa within, as well as among, the sites. Data were rank transformed because raw microwear data typically violate assumptions inherent to parametric tests (Conover and Iman, 1981). I also employed analyses of variance for the individual texture attributes and pairwise comparisons tests to determine the sources of significant variation both among taxa within individual fossil assemblages and between sites. Tukey's Honestly Significant Difference (HSD) and Fisher's Least Significant Difference (LSD) tests were both used to balance the risks of Type I and Type II errors (Cook and Farewell, 1996). Where, p<0.05 for Fisher's but not Tukey's test, the result was considered of marginal significance and suggestive.
### Comparisons of extant and fossil taxa

Hierarchical cluster analysis was also used to cluster fossil groups with extant bovids, using each of the microwear variables that showed significant variation among taxa in the ANOVA models. Each of the texture variables was considered separately because they reflect different scales. Euclidean distance and complete linkage were used, following Fortelius and Solounias (2000). In each case, the extant taxa were grouped by dietary category and the fossil taxa were grouped by site and/or member.

# Results

Microwear texture summary statistics for individual taxa are presented in Table 2. Analytical statistics are presented in Tables 3 and 4. Examples of photosimulations and threedimensional reconstructions of microwear surfaces of the fossil taxa are pictured in Figure 2. Similar images for the comparative taxa are presented in Figures 3.

Statistical analyses of the comparative sample can be found in Scott (in press). The MANOVA results for the fossil bovid sample indicate significant variation in the overall model. Individual ANOVA tests reveal significant variation in all microwear variables except *HAsfc*<sub>81</sub>. Pairwise comparisons tests reveal that the Kanapoi, Allia Bay and Sidi Hakoma samples do not differ significantly from one another in any attributes, although there are suggestive differences between these sites. These samples generally have high values for *Asfc*, *Tfv* and *HAsfc*<sub>9</sub>. The samples from the Laeotil Beds and the Denen Dora and Kada Hadar Members also cluster together, with higher values for *epLsar* and *Smc*.

#### Significant differences among sites by microwear variable

<u>Asfc</u>- Bovids from the sites fall into two discrete complexity groups when analyzed with a Tukey's HSD test- Kanapoi, Allia Bay, the Sidi Hakoma Member and the Laeotil Beds in one group and the Denen Dora and Kada Hadar Members in the other. Kanapoi and Allia Bay have the highest values for this variable, while the Denen Dora and Kada Hadar Members of the Hadar Formation have the lowest value.

<u>Smc</u>- When analyzed with a Tukey's HSD test, only bovids from the Sidi Hakoma Member of the Hadar Formation, which has the lowest value for this variable, and from the Laeotil Beds, which has the highest value, vary significantly in the *Smc* variable. However, a Fisher's LSD test also suggests marginal differences between bovids in the Sidi Hakoma and Denen Dora Members, as well as between those from Kanapoi and the Laeotil Beds.

<u>epLsar</u>- The anisotropy variable also clearly differentiates bovids by site using ANOVA and Tukey's tests. The Kanapoi, Allia Bay and Sidi Hakoma Member of Hadar bovids group together with the lowest values for this variable and are significantly different those from the Denen Dora and Kada Hadar Members of Hadar. The Laeotil Bed bovids have mid-range anisotropy values and are significantly different from Allia Bay, Denen Dora and Kada Hadar.

<u>*Tfv*</u>- Tukey's HSD analyses of the textural fill volume variable suggests significant differences among the sites, resulting in two primary groups. The Kanapoi, Allia Bay and Sidi Hakoma Member bovids cluster together with the highest values for this variable, while the Denen Dora and Kada Hadar bovids have the lowest values. Additionally, the bovids from the Laeotil Beds are significantly different from the Allia Bay sample when analyzed with Tukey's tests, and are marginally different according to Fisher's LSD tests compared with those from Kanapoi and the Denen Dora Member.

<u>*HAsfc9*</u>- The bovids from Allia Bay have the highest value for this variable and are significantly different from those from all of the other included sites, with the exception of Sidi Hakoma. A Fisher's test also suggests differences among the Sidi Hakoma Member, Kanapoi and Denen Dora.

#### Hierarchical cluster analysis

The results of the hierarchical cluster analyses for each variable are depicted in Figure 4. In sum, the Kanapoi, Allia Bay and Sidi Hakoma Member samples cluster together for most variables, along with modern frugivorous, browsing and/or browser-grazer intermediate taxa. On the other hand, the fossil samples from the Laeotil Beds and the Denen Dora and Kada Hadar Members generally cluster with the extant obligate and/or variable grazers.

### **Results by variable**

<u>Asfc</u>-There are two distinct clusters for microwear complexity (*Asfc*). The taxa with the highest values for this variable, including the fossils from Kanapoi, Allia Bay and Sidi Hakoma, and the extant browsers and frugivores, are clearly separate from all other taxa. There is also a second-order division for this variable, with the bovids from the Laeotil Beds and the extant generalists and browser-grazer intermediates separating from the Kada Hadar and Denen Dora fossils and the modern obligate and variable grazers.

<u>Smc</u>- There are also two primary divisions for scale of maximum complexity (*Smc*) variable. The modern browser-grazer intermediates, generalists and frugivores are separated from the other taxa. The latter group contains two main subdivisions: the Allia Bay, Denen Dora and Laeotil Bed fossils with the modern browsers and variable grazers, and the Kanapoi, Kada Hadar and Sidi Hakoma bovids. None of the fossil bovids cluster with modern obligate grazers for this variable.

*epLsar*- The anisotropy variable (*epLsar*) divides the taxa into two primary clusters, with the Denen Dora and Kada Hadar fossils, along with the modern obligate and variable grazers, having higher values for this variable than the other groups. The taxa with the lower values further divide into two clusters. The Allia Bay fossils group with the modern browsers and frugivores, while the Kanapoi, Sidi Hakoma and Laetoli fossils are most similar to extant browser-grazer intermediates and generalists.

<u>*Tfv*</u>- There are also two primary clusters for textural fill volume (*Tfv*). The group with the highest values for this variable includes the Kanapoi and Allia Bay fossils and the modern generalists, browsers, and frugivorous taxa. The remaining group contains two secondary divisions. The first is comprised of the Denen Dora and Kada Hadar fossils and the extant obligate and variable grazers, while the second contains the bovids from Laetoli and Sidi Hakoma, along with modern browser-grazer intermediates.

<u>*HAsfc9*</u>- Finally, the heterogeneity variable (*HAsfc9*) separates the taxa into two primary groups. The fossils from Kanapoi and Allia Bay cluster with the extant browsers, frugivores and generalists. The second cluster is further divided. The Sidi Hakoma and Laetoli fossils group with the modern browser-grazer intermediates while the Denen Dora and Kada Hadar fossils cluster with the extant obligate and variable grazers.

## Discussion

The dental microwear data presented here provide a strong basis for the reconstruction of fossil bovid paleoecology, including habitat preference. These data offer an opportunity to evaluate previous paleoenvironmental reconstructions for Kanapoi, Allia Bay and Laetoli. When

they are considered alongside the data from Hadar, these data can contribute to the larger discussion of the ecological context of hominin evolution.

#### **Paleoenvironmental reconstructions**

*Kanapoi*- The site of Kanapoi is located in the Kerio River Valley of northern Kenya and has been dated to 4.2-4.17 mya (Leakey et al., 1995). It is composed of fluvial and deltaic sediments accumulated during a major lacustrine phase (Harris and Leakey, 2003). The dating and stratigraphy of Kanapoi are well established and provide solid temporal constraints for the *A*. *anamensis* fossil material found there (Leakey et al., 1995, 1998). While Pickford's (2001) study of the hominins from the site has called into question the integrity of the deposit (Pickford, 2001), Leakey et al.'s (1998) descriptions of specimen preservation as carnivore-damaged, fragmented and weathered, indicates that the assemblages are the result of attritional accumulation rather than mixing by taphonomic processes.

The Kanapoi paleosols provide an excellent opportunity for paleoenvironmental reconstruction, because the majority of the early hominin material is derived from two distinct strata, the Dite paleosols (Wynn, 2000). Stable carbon isotope analysis of these paleosols suggests a semi-arid and seasonal climate (Cerling, 1992). The Dite paleosols have high concentrations of carbonates that are only formed during arid periods, and this combined with fossilized seasonal vegetation has led to the hypothesis that *A. anamensis* at Kanapoi inhabited a mosaic environment that ranged from gallery woodlands to forb-dominated edaphic grasslands (Cerling, 1992; Kingston, 1994). Pollen data also suggest that Kanapoi was a highly mosaic habitat. Bonnefille (1995) found large percentages of Gramineae pollen, usually associated with clearings in forest or woodland environments, in the hominin-bearing strata at the site. Most recently, Cerling et al. (2011) reexamined <sup>13</sup>C/<sup>12</sup>C ratio data from hominin-bearing sites at

Kanapoi, as well as many other early hominin localities, and argued that open habitats like wooded grassland would have composed a significant portion of these sites during the late Pliocene.

Analyses of the fossil fauna have suggested that Kanapoi comprised variety of microhabitats. Leakey et al. (1995), categorized the majority of the assemblage from the Dite paleosols as adapted for dry, open bushland, based on taxonomic uniformitarianism, with some browse-adapted bovid taxa like reduncin and aepycerotin indicating the presence of gallery forest in the region, probably along river channels. Reed (1997), on the other hand, used ecomorphological analyses of the fauna and ecological structure analysis to conclude that early australopith habitats were fairly static and tightly constrained by annual rainfall and tree cover. In general, although all lines of evidence have led to the conclusion that a variety of habitats and food resources would have been available to early *A. anamensis*, disagreement remains about the degree of ecological mosaicism at Kanapoi.

The microwear texture data support the previous reconstructions that indicate the presence of both closed forest and open gallery woodland at Kanapoi (Cerling, 1992; Kingston, 1994; Bonnefille, 1995; Cerling et al., 2011). The overall microwear values for the Kanapoi bovids, deposited between 4.2-4.17 mya, are similar to those from Allia Bay at 3.9 mya and the Sidi Hakoma Member at Hadar, dating to 3.42~3.26 mya. This suggests that *A. anamensis* and early *A. afarensis* probably occupied, or at least had regular access to, somewhat closed habitats.

The overall dental microwear of the Kanapoi bovids most closely resembles that of modern browser-grazer intermediates and generalists. When broken down into individual fossil taxa, a range of dietary preferences is indicated. Three of the seven fossil taxa have reconstructed diets similar to those of their closest living relatives. *Madoqua* sp., for example, has a microwear

signature consistent with a browser-grazer intermediate diet, comparable to that of the modern antilopine, *Neotragus batesi*, a species whose diet consists of 80% dicots with some monocot and fruit consumption (Skinner and Smithers, 1990). This extant species is most commonly found in dry forest and thick bushland (Kingdon, 1997). The microwear of *Tragelaphus* sp.is not distinguishable from that of the extant *T. euryceros*, which consumes 75% dicots, along with some fruit and monocots and prefers dense forest habitats (Hofmann and Stewart, 1972). The remaining taxa, however, have wear patterns that suggest a different dietary strategy than that practiced by their modern congeners, suggesting that taxonomic uniformitarianism is not a reliable indicator of food preference for these bovids. For example, the microwear of the Alcelaphini sp. specimens from Kanapoi most closely resembles that of modern browser-grazer intermediates, but most modern alcelaphines are variable grazers. Likewise, *Raphicerus* sp. has a similar microwear pattern to modern browsers, but the three modern species are all classified as either browser-grazer intermediates or generalists.

When considered as a group, the Kanapoi bovids present a picture of a local environment with considerable forest habitat, or at least one with consistent access to nearby forest resources. The results of this study are consistent with the preliminary paleoenvironmental reconstruction of Kanapoi by Leakey et al. (1995). These authors suggested that australopiths at the site occupied a mosaic habitat composed of dry bushland and gallery forest. The microwear texture values for all but two of the bovid taxa from the site fall into the range of browser or browser-grazer intermediate species and are most similar to modern taxa that prefer forest or woodland habitats. However, there are indications of more open habitats at Kanapoi. Wynn (2000) utilized paleosols analysis to suggest that the environment at Kanapoi was more open than previously thought, including open low tree-shrub savanna. The presence of *Simatherium* sp. and *Hippotragus* 

sp.,two bovids with microwear signatures similar to modern open-country adapted grazers, suggests the presence of at least some grassland and supports previous reconstructions of Kanapoi as a mosaic habitat (Wynn, 2000; Cerling et al., 2011).

<u>Allia Bay</u>- The Allia Bay hominin site is located on the eastern shore of Lake Turkana in northern Kenya, and associated with the Omo River Drainage Basin. This site, which has been dated to 3.9 mya, is the last known locality associated with *A. anamensis* (Coffing et al., 1994; Leakey et al., 1995). Although the modern environment at Allia Bay is not riverine, five million years ago, this site was located on the banks of the Omo River (Coffing et al., 1994). The modern habitat is arid grassland, and previous reconstructions have suggested that temperatures during the occupation of *A. anamensis* were probably similar to the present (Dowsett et al., 1996). Reconstructions of ancient sea level, however, suggest that precipitation would have been higher during the Pliocene, making more closed habitats possible (Dowsett et al., 1996).

Most reconstructions of the Allia Bay paleoenvironment suggest a mosaic habitat of open woodland forest with expanses of grassland. The fossil plant composition at the site is primarily C3 trees and shrubs, with only 20-40% C4 grasses associated with open savanna (Cerling et al., 1988). The fossil fauna from the sites also suggests a mosaic paleoenvironment. These assemblages contain mammalian taxa that have been associated with habitats as diverse as gallery forest, bushland and open floodplain (Coffing et al., 1994; Feibel et al., 1991). Feibel et al. (1991) combined both floral and faunal data to suggest that the Turkana Basin was dominated by riparian woodland and gallery forest during the Pliocene, and that grassland probably existed on floodplains and on the borders of the extensive seasonal stream systems. Carbon and oxygen isotope analysis of fossil herbivore tooth enamel also indicates a variety of habitat types and

eurytopic adaptations for all documented taxa (Schoeninger and Reeser, 1999; Schoeninger et al., 2003).

While most studies have suggested a mosaic habitat dominated by woodland, some have opined that savanna was a more important component of the overall environment than previously thought. Using a study of pedogenic carbonates, Wynn (2000) suggested that open savanna was the predominant habitat type at Allia Bay during the occupation of *A. anamensis*. Carbon isotope ratios of marine samples also hint at a drier, more arid climate, similar to that at the site today (Raymo et al., 1996). In sum, while most studies agree that the early hominins had access to a variety of resources at Allia Bay, there is a lack of consensus on the importance of open versus closed habitats at the site.

Studies of stable isotopes (Cerling et al., 1988), palynology and faunal composition (Feibel et al., 1991; Coffing et al., 1994) suggested that Allia Bay was dominated by more closed habitat; and the bovid microwear data presented here are consistent with this hypothesis. The alcelaphin and *Tragelaphus* sp. specimens have dental microwear signatures that overlap with modern browsers and browser-grazer intermediates. While these data do not support previous reconstructions of Allia Bay as composed entirely of dry savanna, it does leave open the possibility of a mosaic habitat that included open grassland.

The alcelaphin sample from Allia Bay, while small, has a microwear signature that is very different from that of their modern relatives, *Damaliscus lunatus* and *D. pygargus*, which are classified as obligate and variable grazers respectively (Gagnon and Chew, 2000). In fact, the microwear textures of alcelaphins from this site are most similar to the extant steenbok, *Raphicerus campestris*, which consumes mostly browse and fruit, and grazes less than 30% of the time (Skinner and Smithers, 1990). This diverse diet is supported by preferred habitats that

include woodland and acacia savanna. Kingdon (1997) suggested that these bovids prefer transitional or even ecologically unstable habitats, and this may indicate that the Allia Bay paleohabitat had similar mosaic qualities.

Reconstructions based on taxonomic uniformitarianism are more consistent with microwear textures for other bovids from Allia Bay. The *Tragelaphus* sp. specimens from Allia Bay have a microwear signature that is indistinguishable from that of the modern tragelaphine, *T. euryceros*. The modern bongo occupies dense forests and is typified by a diet that consists primarily of browse from shrubs and low-hanging trees (Kingdon, 1982, 1997). The similarity between the microwear textures of these congeners strongly implies the presence of some forest coverage in or near the Allia Bay site.

While the previous reconstructions for Allia Bay have suggested a relatively closed habitat, there are also indicators of a highly seasonal environment. Macho et al. (2003) reported dental stress lines in the teeth of fauna from the site, including bovids, and that these strongly suggested seasonal fluctuations similar to those found in parts of Africa today (Feibel, 1999). Research on the relationship between dental microwear and seasonality are limited for bovids (Merceron et al., 2010), but study to date does suggest that seasonal changes in diet are reflected in the dental microwear texture. The results of the current study suggest that seasonal variation was present at Allia Bay, as all of the fossil bovid taxa have microwear signatures that are most similar to modern taxa with highly seasonal diets.

*Laetoli*- The Laetoli hominin site is located in northern Tanzania and is composed of an extensive series of outcrops that date from 4.2 mya-120 kya (Leakey and Harris, 1987). The strata associated with *A. afarensis*, the Upper Laetolil Beds, are dated to 3.7-3.5 mya. These beds are composed of several localities separated by volcanic tuffs that have been reliably dated to 3.6

mya (Hay, 1987). In general, reconstructions of Laetoli during the occupation of *A. afarensis* have been inconsistent, with inferred habitat ranging from open savanna to closed wetland (Leakey and Harris, 1987; Radosevich et al., 1992).

Early ecological studies hypothesized that the environment inhabited by the australopiths at Laetoli was similar to the modern savanna grassland habitat at the site (Hay, 1980; Harris, 1985; Bonnefille and Riollet, 1987; Leakey and Harris, 1987). This conclusion was supported by sediment cores that revealed extensive aeolian deposits of sand-sized ash particles, indicating a habitat with xeric vegetation (Hay, 1980). Paleosol analysis by Hay (1987) also revealed high frequencies of ash and phillipsite, commonly formed in dry, arid climates. Finally, pollen analysis and assemblages of preserved leaves and twigs from the hominin-bearing strata were reported to contain floral taxa closely related to modern species found in savanna grassland habitats (Bonnefille and Riollet, 1987; Hay, 1987).

Faunal assemblages have also been cited as support a savanna grassland reconstruction. Gentry (1981) observed that the Laetoli bovid assemblage is composed primarily of alcelaphin, antilopin, and neotragin taxa whose modern congeners inhabit more open habitats. Other taxa have also been utilized for paleoenvironmental reconstructions at Laetoli. Suid species associated with dry, open conditions, like *Nyanzachoerus euilus*, are among the more common fauna recovered from the site (Harris, 1987). Additionally, two arid-adapted species of tortoise (*Geochelone laetoliensis* and *Geochelone brachygularis*) are also found in the assemblages, as well as several genera of gastropods not common outside of dry environments (Meylan and Auffenberg, 1987; Verdcourt, 1987). All of these lines of evidence have led to the conclusion that dry savanna grassland and open woodland were dominant habitat types at Laetoli during the occupation of *A. afarensis*.

However, other analyses have suggested the Laetoli was a more complex environment than previously thought and might have included closed, deciduous woodland forest as an important part of the ecosystem (Cerling, 1992; Reed, 1997; Musiba, 1999; Harrison, 2005, Musiba et al., 2007; Su and Harrison, 2007). Andrews' (1989) study of ecological diversity at the site was the first to suggest a forested habitat, concluding that the frequency of arboreal fauna at Laetoli was best explained by a heavily wooded setting. He did note, however, that the presence of these taxa could also be the result of faunal mixing from a mosaic ecosystem or habitat changes during the late Pliocene. However, more recent studies of the Laetoli fauna have also supported the closed woodland/mosaic hypothesis. For example, Reed (1997) argued that the diverse assemblages of frugivorous and arboreal mammals are strong indicators of the presence of closed woodland habitats at the site.

Studies of bovid ecomorphology have also suggested that the initial reconstructions of Pliocene Laetoli as similar to the modern savanna grassland might be too simplistic. In fact, modern congeners of many of the bovid taxa represented in the fossil assemblage are not currently found in the Serengeti region. Although some taxa are associated with open habitats, others are more often found in wooded or mosaic habitats. These include Tragelaphini, Bovini and Cephalophini (Gentry, 1987; Bobe et al., 2002). Indicator taxa common in modern savanna ecosystems, like Hippotragini, are not abundant at Laetoli, while the bushland-adapted *Madoqua* is among the most common taxa at the site (Gentry, 1987; Kingdon, 1997). Additionally, the presence of forest adapted pythons and elephant shrews contradict the savanna model and suggest the presence of mosaic woodland habitats (Meylan and Auffenberg, 1987; Butler, 1987). The emerging picture of Laetoli as a complex environment is also supported by evidence from isotopes and palynological studies. Cerling (1992) conducted stable isotope analysis of the

paleosols from the hominin-bearing strata and found evidence for substantial quantities of C3 plants, suggesting that wooded habitats were common at Laetoli during the time of *A. afarensis*. Palynological comparisons of the modern and fossil pollen assemblages at the site are also consistent with more closed forests in the Pliocene than today (Bonnefille and Riollet, 1987).

The conflicting results from various studies of floral and faunal assemblages at Laetoli have been interpreted as indicators of a complex environment composed of a wide range of open and closed microhabitats. The dramatically different interpretations of the site may reflect a sampling bias or misinterpretation of the stratigraphically indistinguishable tuffs that make up the hominin-bearing strata (Gifford, 1981; Musiba et al., 2007). Another possible explanation for the confusing signal is taphonomic bias, and previous studies have suggested both sediment mixing and extensive bioturbation as potential contributing factors (Andrews, 1989; Harrison, 2005).

The reconstructions of paleoenvironment of the Upper Laetolil Beds have been more conflicting than those for any of the other sites addressed in this study, with inferred habitat ranging from closed woodland forest to open, arid savanna. If we assume that the faunal assemblages collected are not the result of taphonomic bias (Andrews, 1989; Harrison, 2005) or ill-defined stratigraphy (Gifford, 1981), the picture that emerges for the Upper Laetolil Beds is one of a mosaic environment even more complex than the one found at the modern site. The microwear evidence supports this hypothesis with dietary reconstructions for local bovid fauna including dedicated browsers/frugivores and variable grazers.

The Laetoli bovids evince a wide variety of microwear signals, suggesting a myriad of available resource types. Most of the fossil bovids from Laetoli have microwear signatures that overlap with modern browser-grazer intermediates. The alcelaphin and *Simatherium* sp.

specimens are most similar to the extant eland, *Taurotragus oryx*. This taxon inhabits grassland, bushland and open woodland and is known for a highly seasonal diet alternating between grazing and browsing (Skinner and Smithers, 1990; Kingdon, 1997). The microwear of the bovinspecimens overlaps with *Raphicerus sharpei*, the modern grysbok. This taxon lives in shrub and tall grasslands, consuming approximately 70% browse/fruit and 30% grasses (Kingdon, 1982). Finally, the specimens of *Gazella* sp. from Laetoli have a texture signature similar to that of the modern springbok, *Antidorcas marsupialis*, which occupies shrub and grasslands and consumes a variable diet (Skinner and Smithers, 1990).

Although many of the Laetoli bovids have been reconstructed as consuming a wide variety of resources, others appear to have had narrower diets. For example, *Antidorcas* sp., *Madoqua* sp., and *Tragelaphus* sp. all have microwear signatures that overlap with modern dedicated browsers and/or frugivores, suggesting regular access to forest resources. *Antidorcas* sp. most closely resembles the modern pygmy antelope, *Neotragus batesi*, which occupy dense forest and thick bushland habitats (Kingdon, 1982). The *Tragelaphus* sp. sample has a texture signal similar to that of their modern congener, *T. euryceros*, the bongo. This species lives in dense forest habitats and consumes more than 85% browse and fruit resources (Hofmann and Stewart, 1973; Kingdon, 1982). The *Madoqua* sp. specimens from Laetoli cluster with extant frugivorous duikers, *Philantomba monticola*, that inhabit rainforest (Kingdon, 1982).

Other fossil taxa appear to have focused on grazing. The *Hippotragus* sp. sample has a microwear signature nearly identical to that of the modern variable grazer, *Damaliscus pygargus*, which occupies open highveld grasslands. The fossil *Parmularius* sp. also has a grazing signature, most similar to *Kobus ellipsiprymnus*. These modern waterbucks prefer open savanna and scrub habitats, which provide year-round access to grasses (Kingdon, 1997). The fact that

these likely grazers are common in the faunal assemblage at Laetoli suggests the presence of, or at least proximity to, more open grassland environments.

The presence of bovids exhibiting a wide variety of dietary strategies suggests that the Laetoli hominin site was highly mosaic in nature. The dental microwear evidence presented here indicates that the local bovid taxa had regular access to resources found in both open and closed microhabitats. This supports and strengthens the previous reconstructions of Laetoli as containing closed forest (Andrews, 1989; Cerling, 1992; Reed, 1997; Musiba, 1999; Harrison, 2005, Musiba et al., 2007; Su and Harrison, 2007), but does not rule out the presence of grasslands, given to the presence of at least two grazing species. The predominance of browse-adapted taxa suggests, however, that closed habitats made up a significant component of the paleoenvironment at the site.

<u>Hadar</u>- The Hadar hominin locality in the Afar Region of Ethiopia dates to 3.42-2.9 mya and is the last known site associated with *A. afarensis* (Walter et al., 1984; Walter and Aronson, 1993; Walter, 1994; Campisano and Feibel, 2007). Approximately 370 *A. afarensis* individuals have been recovered from the site, along with more than 7500 other vertebrate specimens (Reed, 2008). As at Laetoli, paleoenvironmental reconstructions based on the associated fauna, sedimentology and paleosols at Hadar have generally indicated the presence of a mosaic of habitats that fluctuated between open and closed and woodland and grassland (Johanson et al., 1982; Radosevich et al., 1992).

The Hadar Formation was occupied by *A. afarensis* for more than 400 ky, and remains of these fossil hominins have been recovered from the Sidi Hakoma (~3.42-3.26 mya), Denen Dora (~3.26-3.2 mya), and Kada Hadar (<~3.2 mya) members (Taieb et al., 1976; Campisano, 2007). Studies of local geology have revealed that these formations were deposited during periods of

high fluvial activity of a large river system associated with the site, as well as incursions of a reoccurring paleolake (Aronson and Taieb, 1981). Comparison of preserved floral and faunal assemblages from the various members provides a unique opportunity to assess habitat change over time at Hadar during the occupation of *A. afarensis*.

Details of the dental microwear of the bovids from the Hadar hominin locality are detailed elsewhere (Scott, in review), and will be considered here in relation to results from the other sites. In general, significant differences were found between the microwear textures of the bovid samples from the three hominin-bearing strata: the Sidi Hakoma Member, the Denen Dora Member and the Kada Hadar Member. While all three faunal assemblages evidently contain both grazers and browsers, there is a general trend toward increasing numbers of variable and obligate grazers in the more recent strata.

#### Dental microwear texture analysis as a test of taxonomic uniformitarianism

One of the advantages of dental microwear analysis is that it provides direct evidence of what an animal actually ate during its lifetime. Many other forms of dietary reconstruction rely on morphology alone, resulting in conclusions based on what the animal may have been adapted to eat. However, diet can change quickly due to shifts in available resources and it can take many generations for morphology to change accordingly. Sponheimer et al. (1999) used stable isotope analysis to compare dietary reconstructions of fossil bovids from the South African site of Makapansgat based on taxonomic uniformitarianism to those resulting from the isotopes. Their results indicated inconsistencies between the reconstructions for nearly half of the taxa they studied. If bovids are going to be used as paleoenvironmental indicators, accurate reconstructions of diet must be available.

The current study offers similar results for the fossil bovids of Allia Bay, Kanapoi, Laetoli and Hadar. Only three of the seven taxa from Kanapoi, two of the three at Allia Bay, and five of the nine at Laetoli have reconstructed diets similar to their modern counterparts. Morphology is an important factor in how we understand and interpret paleoecology and should therefore continue to be a major component in dietary reconstruction. However, as this study and that of Sponheimer et al. (1999) have demonstrated, a combined approach including direct evidence of diet like microwear or isotopes, as well as morphology should be applied whenever possible.

#### African Pliocene Climate Dynamics and Human Evolution

Early studies of hominin paleoenvironments have suggested that the changes in locomotion and dentition associated with *A. anamensis* and *A. afarensis* were adaptations to receding forests and spreading savannas. However, marine sediment cores have revealed that increases in aridity and the spread of extensive grasslands did not occur in East Africa until after 3.0 mya and would not have influenced the habitats of the gracile australopiths (Tiedemann et al., 1994; deMenocal, 1995; deMenocal and Bloemendal, 1995). Thus, other explanations for how the morphological adaptations of early hominins correlate to environmental change at fossil sites must be sought.

Although disagreement over the interpretation of paleoenvironmental data from the hominin sites in eastern Africa exists, previous reconstructions of the Kanapoi-Allia Bay-Laetoli-Hadar chronology have generally suggested a predominance of mosaic environments and a possible increase over time in the frequency of open habitats. Both Kanapoi and Allia Bay have paleoenvironmental signals that suggest the presence of seasonal habitats during the Pliocene occupation of *A. anamensis*. These habitats were likely composed of mosaic woodland with

some incursions of grassland, although disagreement remains on the degree to which open habitats were present. Reconstructions of the paleoenvironment at Laetoli have not reached a consensus, with interpretations varying from open grassland and closed woodland. However, most recent studies have suggested that the site was composed of several microhabitats during the occupation of *A. afarensis* and that both forest and savanna biomes were important parts of the local ecology. Finally, the Pliocene environment at Hadar has also been reconstructed as mosaic, including closed-forest, bushland, open-woodland and edaphic-grassland habitats. Open habitats appear to have been more common at Hadar than at other sites.

These reconstructions have been used to suggest that the morphological adaptations of *A*. *anamensis* and *A*. *afarensis* were not responses to directional increases in the prevalence of open, dry environments, but could instead reflect increasingly fluctuating seasonal conditions across a range of locally available habitats, where the ability to occasionally utilize lower-quality foods could be advantageous (Teaford and Ungar, 2000; Ungar, 2004; Reed and Fish, 2005; Kimbel et al., 2006). If this is the case, other local species should also be affected. This suggests that the use of faunal assemblages may be essential in understanding the relationship between human evolution and environmental change in East Africa.

The results presented here support the presence of these highly mosaic habitats, as well as an increase in the amount of available grassland at the sites associated with *A. afarensis*. This suggests that the habitats associated with *A. anamensis* were more closed than those of the later hominins. The microwear signatures of fossil taxa from Kanapoi and Allia Bay are most similar to those of modern taxa that inhabit closed forest, open woodland and bushland. The presence of a few taxa with grazing signatures suggests that some edaphic grassland may have been present either within or near the fossil sites. However, these grazing taxa have been recovered in small

quantities and may have been migratory or seasonal members of the local faunal assemblages. The microwear evidence also suggests that *A. afarensis* occupied highly mosaic environments that included both open and closed habitat types. This would account for the presence of both grazing and browsing adapted bovid taxa. Even with some evidence of graze-adapted taxa at australopith sites, open and closed woodlands appear to have important components to the habitats. While habitat reconstructions alone cannot conclusively demonstrate that the adaptations of the australopiths were in response to specific environmental conditions, anatomical studies of these hominins suggest morphological adaptations for terrestrial bipedality, but that they also retained attributes associated with arboreal locomotion (Stern and Susman, 1983; Susman et al., 1984, Ward et al., 2001).

While there are some indications that the Laetoli and later Hadar members included drier, more open habitats, the overall results of this study suggest that both of these early australopith species occupied highly mosaic environments. The results presented here generally agree with Stanley (1992) that Pliocene hominin environments did not shift dramatically; however, there is some evidence for an increase in the number of grazing taxa at Laetoli and the later Hadar member. This supports the observation by Reed (1997) that the overall composition of the mammalian communities at these sites indicates a gradual aridification trend in eastern Africa that may have resulted in drier, more open habitats and the presence of a more pronounced dry season. This could explain why these early australopith taxa disappear around 2.8-2.5 mya and are replaced by members of the more ecologically adaptable genus *Homo*.

# Conclusions

The ability of dental microwear texture analysis to separate modern bovids with known diets and to assign dietary categories to fossil taxa has been demonstrated in previous papers (Ungar et al., 2007; Scott, in press, in review). This study applies the technique to fossil taxa from four of the sites occupied by *A. anamensis* and *A. afarensis*, with the goal of inferring paleoenvironment at the sites. In general, the dental microwear signatures of the fossil bovids from Kanapoi, Allia Bay, Laetoli and Hadar all reflect an array of dietary preferences and suggest access to a variety of resource types. Specifically, all of these sites have bovid taxa with microwear signatures similar to modern species that occupy wooded or thick bushland habitats, but grazers are still present in small but increasing numbers. This suggests that the early australopiths may have preferred mosaic environments that would have provided both protection from predation and access to an assortment of food types.

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# Figures

Figure 1. Occlusal view of bovid M2, illustrating the shearing facets where microwear was examined (Illustration courtesy of Gildas Merceron). A = left  $m^2$ , B= right  $m_2$ 



Figure 2.Examples of dental microwear surfaces of taxa from (a) Kanapoi, (b) Allia Bay, (c)
Laetoli. These images are photosimulations based on data collected using the confocal imaging profiler, and each represents an area of 276 x 204 µm. Kanapoi: A) *Aepyceros*;
B)Alcelaphini; C) Hippotragini; D) *Madoqua*; E) Raphicerus; F) *Simatherium*; G) *Tragelaphus*. Allia Bay: A) Alcelaphini; B) *Tragelaphus*. Laetoli: A) Alcelaphini; B) *Antidorcas*; C) Bovini; D) *Gazella*; E) Hippotragini; F) *Madoqua*; G) *Parmularius*; H) *Simatherium*; I) *Tragelaphus*.

#### Kanapoi



# Allia Bay



#### Lactoli





Figure 3.Examples of dental microwear surfaces for extant bovids from different dietary categories. These images are 3-D photosimulations based on data collected using the confocal imaging profiler, and each represents an area of 138 x 102 μm. a) obligate grazer; b) variable grazer; c) browser-grazer intermediate; d) generalist; e) browser; and f) frugivore.



Figure 4.Hierarchical cluster analysis of fossil and extant taxa by site or dietary category. These results are organized by microwear texture variable: (a) *Asfc*,surface complexity; (b) *epLsar*, anisotropy; (c) *Smc*, scale of maximum complexity; (d) *Tfv*, fill volume; *HAsfc9*, heterogeneity.


# Tables

**TABLE 1.**Extant bovid taxa included in this study, along with associated diets

Taxon	Common Name	Diet
<b>Obligate Grazers</b>		
Damaliscus lunatus Kobus leche Redunca arundinum Redunca fulvorufula	Tsessebe Lechwe SouthernReedbuck Mountain Reedbuck	Savanna and floodplain grasses Aquatic marsh grasses Tall grass specialist Seasonal grasses
Variable Grazers		
Damaliscus pygargus	Bontebok	Short Highveld grasses, some fynbos
<i>Gazella granti</i> shrub foliage	Grant's Gazelle	Short grasses, seasonal herbs and
<i>Hippotragus niger</i> foliage in dry season	Sable Antelope	Grasses in wet season and forbs and
<i>Kobus ellipsiprymnus</i> some rushes	Waterbuck	Medium to short grasses, reeds and
<i>Synceos caffer</i> swamp vegetation dur	African Buffalo ring dry season	Pasture grasses year-round and
<i>Tragelaphus spekii</i> low-level vegetation	Sitatunga	Swamp grasses, sedges, leaves and
Browser-Grazer Inte	ermediates	
Aepyceros melampus	Impala on seedpods, foliage, forbs a	Grazes on grasses when green, also browses nd shoots
Antidorcas marsupial young grasses during	<i>is</i> Springbok wet season	Shrubs and succulents in dry season,
Raphicerus campestri pods during dry seaso	<i>is</i> Steenbok on, young grass in wet season	Low herbaceous plants, seeds and
Raphicerus sharpei Taurotragus oryx	Sharpe's Grysbok Eland	Mature plants, fruit pods and roots Seasonal young grasses, small herbaceous
plants, roots, and tube <i>Tragelaphus imberbis</i> grasses, pronounced s	ers, highly seasonal Lesser Kudu easonal variation	Succulents, buds, leaves, pods and
<b>Browsers</b> <i>Litocranius walleri</i> and leaves of shrubs	Gerenuk not known to eat grass or here	Pure browsers, focuses on flowers
Neotragus batesi	Bate's Pygmy Antelope	Ground-level herbaceous vegetation

*Sylvicapra grimmia* Common Duiker tubers, insects, rarely known to consume grasses *Tragelaphus euryceros* Bongo including shrubs and creeping plants

## Generalists

*Oreotragus oreotragus* Klipspringer grasses and some fruit *Tragelaphus angasi* Nyala *Tragelaphus strepsiceros* Greater Kudu some grass

## Frugivores

Cephalophus sylvicultor Yellowback Duiker Philantomba monticola Blue Duiker mushrooms Leaves, shoots, fruit pods and seeds,

High-concentrate green herbage,

Leaves, buds, pods, bark, short

Mixed grasses and leaves Herbs, leaves, vines, fruit, tubers and

> Forbs and shrubs Fallen fruit, leaves, buds, seeds and

Diet information based on Walker, 1975; Kingdon, 1982, 1997; Grzimek, 1990; Nowak, 1991.

# Table 2. Median values by fossil taxon.

ΚΔΝΔΡΟΙ	
KANAI UI	

ICI II UII OI								
	Alcelaphini	5	1.4196	0.0037	0.7667	5740.80	0.4381	0.6433
	Aepyceros sp.	1	2.3279	0.0030	0.4502	8649.56	0.6444	1.0553
	Simatherium sp.	3	1.4262	0.0064	0.5015	1485.66	0.5266	0.9517
	Hippotragus sp.	2	2.0592	0.0054	0.3283	2636.06	0.3148	0.7553
	<i>Madoqua</i> sp.	9	3.9542	0.0035	0.9171	16220.87	0.4999	0.7307
	Raphicerus sp.	3	3.6907	0.0021	0.3645	5799.58	0.6612	1.1978
	Tragelaphus sp.	12	3.9042	0.0018	0.1502	13486.87	0.4030	0.7444
ALLIA BAY	Y							
	Alcelaphini	1	2.4609	0.0020	0.2692	4930.21	0.5960	0.6962
	Tragelaphus sp.	14	3.2013	0.0025	0.7820	10888.61	0.6248	0.9839
LAETOLI								
	Alcelaphini	10	2.1934	0.0048	1.0958	6551.74	0.4062	0.8868
	Antidorcas sp.	3	4.0596	0.0013	0.5302	8598.90	0.5264	1.0024
	Bovini	10	2.2624	0.0040	0.6885	4602.28	0.4606	1.2403
	<i>Gazella</i> sp.	4	2.0844	0.0030	0.5366	4287.92	0.3683	0.7620
	Hippotragus sp.	2	1.2170	0.0050	0.3109	6871.15	0.6293	0.7855
	<i>Madoqua</i> sp.	3	3.9078	0.0016	0.8505	14011.05	0.6154	1.2027
	Parmularius sp.	10	1.2937	0.0051	1.1150	3627.816	0.3761	0.7902
	Simatherium sp.	3	2.4788	0.0030	0.5208	7784.08	0.4644	0.7690
	Tragelaphus sp.	4	3.8115	0.0024	0.7204	13683.39	0.6426	0.8986
SIDI HAKO	MA							
	Aepyceros	3	3.5261	0.0024	0.2976	8752.488	0.5298	0.9452
	Alcelaphini	4	2.2392	0.004	0.4058	6598.503	0.5243	0.8843
	Antilopini	5	4.8171	0.0033	0.6316	11977.21	0.6612	1.1559
	Gazella	2	2.2886	0.0049	0.6128	6071.115	0.4985	0.7292
	Kobus	2	2.491	0.0021	0.9725	4184.89	0.3265	0.8375
	Neotragus	3	4.2655	0.0019	0.6865	9763.845	0.6935	0.9531
	Tragelaphus	5	3.0762	0.0031	0.6449	8062.735	0.544	0.8553
	Ugandax	6	2.9872	0.0035	0.4532	5578.615	0.5796	0.8406
DENEN DO	DRA							
	Aepyceros	4	1.7485	0.0039	0.5043	5371.542	0.4269	0.7503
	Alcelaphini	10	2.2371	0.0048	0.426	6239.596	0.5006	0.9241
	Gazella	2	2.7917	0.0052	0.5057	5333.485	0.4076	0.8075
	Hippotragini	3	1.9785	0.006	0.4368	3984.685	0.4502	0.8135
	Kobus	9	1.5987	0.0037	1.1877	4462.355	0.4199	0.8613
	Pelorovis	2	1.2321	0.0061	0.7799	2328.855	0.5062	0.855
	Reduncini	14	0.9916	0.006	1.2154	2593.004	0.4446	0.8154

	Tragelaphus	10	1.964	0.005	0.8553	6539.146	0.4823	0.7203
	Ugandax	9	2.0688	0.0049	0.5026	4267.688	0.5133	0.8941
KADA HAI	DAR							
	Aepyceros	5	1.4847	0.0041	0.2685	5972.351	0.3865	0.8544
	Alcelaphini	4	1.4858	0.0076	0.3224	3331.107	0.4	0.7939
	Antilopini	5	3.5876	0.003	0.6188	9597.235	0.6988	1.065
	Bovini	5	0.9267	0.007	1.0667	3564.102	0.428	0.7242
	Gazella	2	2.2814	0.0052	0.5349	4276.6	0.4512	0.8606
	Hippotragini	2	1.3157	0.0049	0.4005	2626.581	0.4049	0.9214
	Kobus	1	0.9877	0.003	1.1355	1103.832	0.4325	0.7613
	Neotragus	1	2.6987	0.002	0.6247	10268.15	0.6265	0.9884
	Oryx	3	0.9987	0.0077	1.2649	1168.655	0.5985	0.9755
	Pelorovis	2	1.3155	0.0061	0.7532	2677.005	0.4278	0.7539

# Table 3.Nested analysis of variance.

Source	SS	df		MS		F		р	
Complexity (A.	sfc ranked data)								
Asfc	255292.5	5 5		51058	3.52	17.28		0.00	
Error	6.	32021.41	214		2953	.37			
Anisotropy (ep	Lsar ranked data	ι)							
<i>epLsar</i>	238143.19 5	Ś	47628	8.64	15.70	1	0.00		
Error	64	49166.81	214		3033	49			
Maximum Con	nplexity (Smc rai	nked data)							
Smc	83613.58	3 5 <sup>°</sup>		16722	2.72	4.453		0.001	
Error	8	03701.42	214		3755.	.52			
Fill Volume (T	<i>fv</i> ranked data)								
Diet	226891.1	8 5		45378	3.24	14.704	Ļ	0.00	
Error	6	60422.82	214		3086	.09			
Heterogeneity	3x3 ( <i>HAsfc₀</i> rank	(ed data)							
Diet	121338.4	1 5		24267	7 69	6 780		0.00	
Error	7	65975.09	214	21201	3579	.32		0.00	
Heterogeneity	9x9 ( <i>HAsfcor</i> ran	ked data)							
Diet	34692 59	) 5		6938	52		1 742		0 1 2 6
Error	8	526219	214	0750.	3984	22	1.7 12		0.120

Table 4. Pairwise comparisons of bovids by fossil site/member.*- significant with Fisher's LSE
test, **- significant with Tukey's HSD test.

MEMBER	MEMBER	Asfc	epLsar	Smc	Tfv	HAsfc9
Allia Bay	Denen Dora	88.477**	-97.244**	3.004	94.46**	81.024**
Allia Bay	Kada Hadar	93.067**	-104.617**	26.933	95.683**	81.45**
Allia Bay	Kanapoi	23.6	-37.676*	34.181	24.219	83.819**
Allia Bay	Laetoli	55.51**	-57.166**	-9.178	64.631**	79.725**
Allia Bay	Sidi Hakoma	7.759	-30.57	45.844*	36.326*	39.633*
Denen Dora	Kada Hadar	4.59	-7.372	23.929	1.223	0.426
Denen Dora	Kanapoi	-64.877**	59.568**	31.177*	-70.241**	2.795
Denen Dora	Laetoli	-32.966**	40.078**	-12.182	-29.829*	-1.299
Denen Dora	Sidi Hakoma	-80.717**	66.674**	42.84*	-58.134**	-41.391**
Kada Hadar	Kanapoi	-69.467**	66.94**	7.248	-71.464**	2.369
Kada Hadar	Laetoli	-37.556**	47.451**	-36.112*	-31.052*	-1.725
Kada Hadar	Sidi Hakoma	-85.307**	74.046**	18.911	-59.357**	-41.817*
Kanapoi	Laetoli	31.91	-19.49	-43.359**	40.412**	-4.094
Kanapoi	Sidi Hakoma	-15.841	7.106	11.663	12.107	-44.186**
Laetoli	Sidi Hakoma	-47.751**	26.596*	55.023**	-28.305*	-40.092*

## **Chapter Five: Summary and Future Directions**

Some of the most important events in early human evolution occurred during the Pliocene epoch in eastern Africa, including major taxonomic diversification of early hominins and the introduction of bipedalism, our diagnostic form of locomotion. Previous hypotheses have linked these events to environmental pressures caused by global and regional climatic shifts (Vrba, 1985, 1988, 1995a, 2000; Stanley, 1992; deMenocal, 1995), and various lines of evidence have indicated that while the earliest known hominins inhabited closed woodland habitats (WoldeGabriel et al., 1994), middle Pliocene species witnessed a shift towards more open and mosaic environments that would have made changes in locomotion and diet advantageous (Kingston et al., 1994; Kappelman et al., 1997; Reed, 1997; Schoeninger et al., 2003).

*Australopithecus anamensis* and *Australopithecus afarensis* are two of the earlier identified hominins in eastern Africa and have been suggested to be part of an anagenetic lineage based on subtle changes in morphology over time (Leakey et al., 1995; Wolpoff, 1999; Ward et al., 2001; Kimbel et al., 2006). These two hominins have been primarily recovered from four fossil sites that span from 4.2-3.0 mya. Two of the primary sites associated with *A. anamensis* are Kanapoi and Allia Bay, both located in the Turkana Basin of Kenya (Leakey et al., 1995, 1998). Laetoli in Tanzania is the earliest known habitat associated with *A. afarensis* and Hadar in Ethiopia is the latest known site occupied by this species (Leakey and Harris, 1987; Lockwood et al., 2000; White et al., 2000). These hominins are characterized by morphological adaptations for habitual bipedalism and more robust dentition than earlier taxa like *Ardipithecus ramidus* (White et al., 1994). Additionally, Johanson et al. (1987) observed that these middle Pliocene hominins also exhibit short legs and long arms in relation to trunk length, which indicates combined terrestrial bipedality and arboreal climbing. This adaptation has been hypothesized to be the result of environmental changes in eastern Africa that would have made such a unique locomotor pattern advantageous (Stanley, 1992).

Establishing a direct relationship between climate change and evolution presents a unique challenge because climate is generally preserved in one context and evolution in another. Proxy climate records for the Pliocene have been recovered primarily from deep-sea sediments and often represent more than one million years of continuous environmental history (Dowsett et al., 1996; deMenocal and Brown, 1999). These extensive records provide an opportunity to analyze the tempo of climatic change and compare overarching trends to major events in human evolution. Records of evolution, however, are not continuous time series, but snapshots over a wide geographic range, usually based solely on fossil remains of hominins and other fauna.

This dissertation approaches the question of Pliocene hominin paleoenvironments in eastern Africa by reconstructing the diets of the associated bovid fauna. These taxa are considered by many to be reliable indicators of paleoenvironment due to predictable relationships between diet and habitat, and this dissertation explores this relationship for both extant and fossil species. The study also serves as a test of taxonomic uniformitarianism, comparing evidence of actual diet during the lifetime of individuals to the inferred diet based on morphology alone. Finally, the reconstructed bovid diets are used to evaluate previous hypotheses for the paleoenvironments at the sites of Kanapoi, Allia Bay, Laetoli and Hadar, as well as the implications of local habitats for the adaptations of the early australopiths.

## **Project Objectives**

From its conception, this project has had three primary objectives. First, I aimed to confirm the utility of dental microwear texture analysis as a powerful tool for bovid dietary reconstruction. The only study to date to apply this technique to bovids suggested that the method could be used reliably to reconstruct the diets of fossil taxa, and that those results had implications for local paleoenvironment (Ungar et al., 2007). However, in order for the technique to have wider applicability, it was first necessary to develop a large extant comparative database of bovid microwear textures that could be applied to the fossil samples included in this dissertation, as well as samples from other sites and time periods in Africa. In order to accomplish this, texture data were collected for 25 extant African bovid species with known dietary preferences, representing all of the dietary categories previously described by Gagnon and Chew (2000) and resulting in the largest database of microwear textures yet assembled.

The second objective was to reconstruct the diets of fossil taxa from Kanapoi, Allia Bay, Laetoli and Hadar. Microwear texture data were collected for 223 specimens from the four early hominin sites. These include 35 specimens from Kanapoi, 15 specimens from Allia Bay, 49 specimens from Laetoli, and 124 specimens from Hadar. The diets of these fossil taxa were determined by comparing their microwear signatures to those from the modern comparative sample.

In addition to dietary reconstruction, an important component of this dissertation was to explore the efficacy of dental microwear texture analysis as an indicator of paleohabitat. The sites included in this study are four of the key localities associated with the early australopiths and because of this, an accurate understanding the local environment is important. However, there has been considerable disagreement over the paleoenvironments at these sites during the

occupation of the hominins, even after the application of various methods of habitat reconstructions. In an effort to evaluate these competing hypotheses on paleoenvironment, the paleoenvironmental implications of the reconstructed bovid diets were used to evaluate the previous habitat reconstructions for the four sites.

The final dissertation objective was to use the bovid dietary reconstructions as a test of the principle of taxonomic uniformitarianism. This principle assumes that fossil taxa share the ecological preferences of their modern counterparts and while this is usually reliable for recent faunas, its applicability to the distant past is questionable. While many authors have questioned the utility of taxonomic uniformitarianism, only a few have sought to test its accuracy. An examination of the fossil bovids of Makapansgat, South Africa by Sponheimer et al. (1999) used carbon stable isotopes to demonstrate that nearly half of included taxa had actual diets that were different from those of their closest living relatives. In order to assess the accuracy of taxonomic uniformitarianism at the sites included in this dissertation, the reconstructed diets for the fossil bovids from Kanapoi, Allia Bay, Laetoli and Hadar were compared to previous dietary assumptions based on this principle.

Brief summaries and concluding remarks for each goal are presented individual below.

#### **Objective 1: Extant bovid microwear texture database**

It is assumed that a relationship exists between environmental change and evolution, however these connections are difficult to discern because of the challenges involved in accurately reconstructing paleoenvironments. One of the primary ways that past habitats are determined is through the reconstruction of the paleoecology of fossil fauna. Bovids are most frequently used indicator taxa for the paleoenvironments associated with human evolution. This is because they are ubiquitous at hominin fossil sites and modern taxa have can be placed into dietary categories that reliably predict habitat preference. However, since critiques of taxonomic uniformitarianism suggest that morphology alone is not always a reliable predictor of diet, direct line evidence must be sought from other methods. Dental microwear, or the microscopic pits and scratches on the enamel surface that result from the interaction of food and teeth during chewing, is one such line of evidence.

#### Use of dental microwear texture analysis

The first objective was to develop a database of extant African bovid dental microwear that could be used to interpret the texture signatures of fossil bovids. This goal could be subdivided into two further objectives. It was first necessary to confirm pilot data collected by Ungar et al. (2007), which suggested that texture analysis could successfully separate extant bovids with diets from different ends of the spectrum, i.e., obligate grazers and browsers. Previous work also suggested that microwear texture analysis has the potential for finer dietary distinctions (Ungar et al., 2007), so an important component of this study was to explore the resolution of the technique. In order to accomplish this, extant taxa classified as 'mixed-feeders' (i.e., variable grazers, browser-grazer intermediates, generalists) were statistically compared to identify significant differences.

The initial study of bovid microwear texture by Ungar et al. (2007) utilized two microwear texture variables, complexity (*Asfc*) and anisotropy (*epLsar*). Their data indicated that browser taxa have more complex surfaces with lower values for anisotropy than grazing bovids. These differences in complexity and anisotropy were confirmed with the increased sample. Additionally, since the original bovid microwear texture analysis study was conducted, new

variables have been added to the microwear texture suite, including scale of maximum complexity (*Smc*), textural fill volume (*Tfv*), and heterogeneity 3x3 (*HAsfc*<sub>9</sub>) and 9x9 (*HAsfc*<sub>81</sub>). While these variables have proven to be strong indicators of diet in primates and hominins, this relationship had to be explored for bovids of differing diets (for examples, see Ungar et al., 2008; Scott et al., 2009; Grine et al., 2010; Merceron et al., 2010; Ungar et al., 2010; Pontzer et al., in press). All of these variables proved to be significant indicators of bovid diet, although not for all dietary categories.

## Comparative sample selection

The extant taxa included in this study were selected for analysis because of their wideranging differences in diet and habitat preference. Gagnon and Chew (2000) identified six dietary categories for extant African bovids that are based on the percentages of monocotyledons and dicotyledons included in the diet, with separate consideration of fruit consumption. When selecting comparative taxa for this study, representative species from all six dietary categories were chosen, with specific emphasis on bovids that evince various levels of mixed feeding. The included taxa for each dietary category, along with habitat preferences are reviewed below. Dietary categories are from Gagnon and Chew (2000) and habitat preferences are from Kingdon (1982, 1997).

The browsing taxa selected were *Litocranius walleri*, *Neotragus batesi*, *Sylvicapra grimmia*, and *Tragelaphus euryceros*. Dedicated browsers focus primarily on dicots, consuming more than 70% of these resources and including small quantities of grass (Gagnon and Chew, 2000). Because of their preference for forest resources, browsing taxa tend to inhabit more closed habitats, like forest and woodland. The species included here represent a range of closed

habitats, from the rainforest preferred by the common duiker, *Sylvicapra grimmia*, to the scrub forest occupied by the gerenuk, *Litocranius walleri*. Bongos, *Tragelaphus euryceros*, are found in dense forest with considerable undergrowth, and pygmy antelope, *Neotragus batesi*, inhabit woodland forest and bushland.

The grazing taxa include four obligate grazers: *Damaliscus lunatus*, *Redunca fulvorufula*, *Kobus leche*, and *Redunca arundinum*. These bovids have diets that consist almost exclusively of monocotyledonous grasses, usually consuming less than 10% of any other type of resource (Gagnon and Chew, 2000). All four of these taxa inhabit open habitats, although some variation in habitat preference is represented. *Damaliscus lunatus*, the tsessebe, prefers open savanna and floodplains. The lechwe, *Kobus leche*, is typically found in wet marshes. The two *Redunca* species, the southern and mountain reedbuck, both prefer tall, seasonal grassland.

The mixed feeders included six variable grazers: *Damaliscus pygargus*, *Gazella granti*, *Hippotragus niger*, *Kobus ellipsiprymnus*, *Syncerus caffer*, and *Tragelaphus spekii*. Taxa classified as variable grazers consume 60-90% monocots and usually have diets that vary seasonally (Gagnon and Chew, 2000). Variable grazers typically occupy mosaic habitats where they have access to a variety of resource types. In order to include a wide representation of variable grazer environments, the extant sample includes taxa that favor a range of habitat types. This includes species as diverse as *Damaliscus pygargus*, the bontebok, which occupies highveld grassland and *Hippotragus niger*, the sable antelope, which inhabits wooded savanna. Other preferred habitats include edaphic grassland (*Tragelaphus spekii*), closed forest (*Syncerus caffer*), open grassland (*Gazella granti*) and scrub forest (*Kobus ellipsiprymnus*).

The mixed-feeding taxa also included six browser-grazer intermediates: *Aepyceros melampus*, *Antidorcas marsupialis*, *Raphicerus campestris*, *Taurotragus oryx*, *Tragelphus* 

*imberbis*, and *Raphicerus sharpei*. Browser-grazer intermediates include bovids that eat 30-70% of dicots and monocots, and always include some fruit (Gagnon and Chew, 2000). Like variable grazers, browser-grazer intermediates prefer habitats that allow them access to a variety of resource types. Overall, these bovids are more likely to be found in highly mosaic habitats and the extant sample presented here includes taxa that a representative of the environmental preferences of the category. The taxa that prefer more open habitats include the impala, *Aepyceros melampus*, which is found in bushland and open savanna, the springbok, *Antidorcas marsupialis*, usually found in grassland and the common eland, *Taurotragus oryx*, which inhabits savannas and plains. More closed habitat taxa include the lesser kudu, *Tragelaphus imberbis*, which occupies dry bushland and closed forest, and Sharpe's grysbok, *Raphicerus sharpei*, usually found in scrubland. Some browser-grazers intermediates are also highly variable and can be found in both woodland and grassland, like the steenbok, *Raphicerus campestris*.

Three taxa classified as generalists were included in the sample. These are *Oreotragus oreotragus*, *Tragelaphus angasi*, and *Tragelaphus strepsiceros*. Generalists consume a wide variety of foods and typically include more than 20% of the primary resource types (Gagnon and Chew, 2000). Like most mixed feeders, these taxa occupy a broad range of habitats. This sample includes taxa with a range of habitat preferences. The klipspringer, *Oreotragus oreotragus*, is found in open woodland and savanna, while the nyala (*Tragelaphus angasi*) prefers dense forests. The greater kudu, *Tragelaphus strepsiceros*, is intermediate to the other two, typically inhabiting woodland forest and thick bushland.

Finally, two frugivorous bovids were sampled in order to have microwear textures from all possible extant dietary categories. The two taxa, *Cephalophus sylvicultor* and *Philantomba monticola*, both consume more than 70% fruit, with little to no inclusion of grasses (Gagnon and

Chew, 2000). These two taxa were selected because neither varies its diet seasonally and reliable dietary information based on wild studies is available (Emmons et al., 1983; Dubost, 1984). Both species occupy dense forest habitats where fruit is widespread.

## Results

The second chapter of this dissertation, "Dental microwear texture analysis of extant African Bovidae" addresses the extant microwear texture database collected for this project. This comparative sample is the largest ever collected for bovids and includes 575 individual specimens representing 25 taxa. The technique accurately classified all 25 extant bovid taxa by dietary category, indicating that this dataset can be used to reliably assign dietary preference to the fossil taxa. Clear statistical differences were present among all dietary categories, with the exception of generalists and browser-grazer intermediates, both of which consume a wide variety of graze and browser resources. The results confirm the conclusions of Ungar et al. (2007) that grazers and browsers can be easily identified using dental microwear textures. In fact, there was no overlap between obligate grazers and dedicated browsers for any of the texture variables. In addition to examining the difference among the dietary categories, I also looked at variation within diets. Some significant differences among taxa within the dietary categories were identified. These may relate to seasonal or geographic differences in diet, and points to future research.

In sum, the construction of the extant comparative database for African bovids serves two important purposes. First, it confirms the ability of dental microwear texture analysis to distinguish among the dietary categories, including various levels of mixed feeding. Second, it provides a comprehensive dataset of bovid microwear textures that can be used to interpret the

microwear signatures of fossil taxa from sites across Africa, including the eastern localities addressed in this study.

#### **Objective 2: Bovid paleodiets and paleoenvironments**

The second objective of this dissertation was to reconstruct the diets of the fossil bovids from Kanapoi, Allia Bay, Laetoli using dental microwear texture analysis and Hadar and to use these data to test competing hypotheses regarding the paleoenvironments at these important early hominin sites. Microwear texture data were collected for each of the fossil bovid taxa and the resulting wear signatures were compared to the extant comparative database. Finally, the apparent resource preferences of the fossil bovids were used to infer likely habitat types at the various sites. Previous paleoenvironmental reconstructions were then evaluated based on the microwear texture data and current paleoclimatological models.

#### Fossil sample selection

Inferences of diet are frequently used as proxies for paleoenvironment, so testing the ability of dental microwear texture analysis to accurately classify taxa by dietary preference was a priority (e.g., Robinson, 1963; Vrba, 1974, 1975, 1980, 1988; Kay, 1975, 1978; Grine, 1981; Stern and Susman, 1983; Kappelman, 1988; Benefit and McCrossin, 1990, Ciochon, 1990; Spencer, 1995; Lewis, 1997). Although there is undoubtedly a relationship between bovid diet and habitat, reliable predictions of paleoenvironment depend on the ability to reconstruct diet accurately. Thus, the aim of this project was to develop a large database of fossil microwear textures to insure that the dietary inferences were as well-supported as possible.

The large collections of fossil bovids from the four sites allowed for a large volume of data to be collected. This made it possible to draw inferences not only for diet, but also habitat preference. All available bovid specimens from Kanapoi, Allia Bay, Laetoli and Hadar were examined for microwear preservation. As is typical for fossils, only a small percentage of specimens preserved unobscured antemortem microwear that could be analyzed. The Kanapoi and Allia Bay samples are the smallest, with 35 total specimens from Kanapoi, representing 8 taxa, and 15 from Allia Bay, representing 2 taxa. The Laetoli sample consists of 49 individuals, representing 9 taxa. Finally, the Hadar sample includes 124 specimens: 30 from the Sidi Hakoma Member, representing 8 taxa; 63 from the Denen Dora Member, representing 9 taxa; and 31 from the Kada Hadar Member, representing 10 taxa.

#### Bovids as proxies for paleoenvironment

Previous studies have shown that dental microwear patterns reflect diet in extant bovids (for examples, see Solounias et al., 1988; Solounias and Moelleken, 1993; Merceron and Ungar, 2005; Merceron et al., 2005; Schubert et al., 2006, Ungar et al., 2007). Several studies have explored the implications of dietary reconstructions based on microwear for understanding paleoenvironments (Merceron et al., 2005, Merceron and Ungar, 2005; Ungar et al., 2007). The results of these studies have been promising and this dissertation furthers the exploration of the diet-habitat relationship. The Pliocene hominin sites in eastern Africa are good candidates for testing previous paleoenvironmental hypotheses, as previous research has often led to conflicting reconstructions. The microwear data presented here are consistent with some of the previous reconstructions for Kanapoi, Allia Bay, Laetoli and Hadar, particularly those that suggest the presence of mosaic habitats.

Dental microwear analysis has been used previously to evaluate competing hypotheses for paleoenvironmental reconstruction (for examples, see Ungar et al., 2007; Merceron et al., 2009). Although there is not a one-to-one relationship between habitat type and the diet of local fauna, evidence of consumption of a particular resource type indicates its local availability. For example, the presence of obligate grazers doesn't confirm that the habitat consisted solely of open savanna, but it does strongly suggest the availability of grassland resources. The dietary variability among the local fauna can indicate the different habitat types found at and near a given site, which is particularly useful for sites thought to be highly mosaic.

### Results

In the third chapter, "*Dental microwear texture analysis of fossil bovids from Hadar*, *Ethiopia: implications for the paleoenvironment ofAustralopithecus afarensis*", the diets of the fossil bovids from Hadar were reconstructed using the new comparative database. The dietary categories assigned to the fossil taxa from the Sidi Hakoma, Denen Dora and Kada Hadar Members were then used to address implications for changing paleoenvironments at the site during the occupation of A. afarensis.

The reconstruction of local habitats at Hadar, Ethiopia has been the topic of considerable recent research. *Australopithecus afarensis* occupied this site from approximately 3.42-2.94 million years ago. Because the Hadar includes the last know habitat for the species, an understanding of local environment is important (Leakey and Harris, 1987; White et al., 2000). Although a wide variety of studies, including those utilizing palynological remains (Radosevich et al., 1992; Bonnefille et al., 2004), ecomorphological and diversity analyses of faunal assemblages (Bobe and Eck, 2001; Reed, 2008) and stable isotope data (Hailemichael, 2000),

have suggested a mosaic of microhabitats at Hadar throughout the temporal range of *A*. *afarensis*, some disagreement exists about the amount of fluctuation in habitat type, rainfall and humidity.

Clear differences in bovid microwear textures were found between the members. The Sidi Hakoma bovids are reconstructed as browsers or browser-grazer intermediates, suggesting the presence of woodland resources and more closed habitats during the earliest occupation of the australopiths. The Denen Dora bovids have microwear texture signatures ranging from obligate grazer to browser-grazer intermediate, indicating access to both browse and graze and a mosaic habitat. The Kada Hadar bovids have variable or obligate grazing signatures, although a few browsing taxa are still present. This suggests open grassland habitats would have been present during the end of the Hadar occupation by *A. afarensis*, although the presence of browsing taxa supports a more mosaic habitat reconstruction. The results presented here suggest increasing aridification over time at Hadar and highlight environmental changes that might have influenced the evolution and eventual extinction of *A. afarensis*.

The fourth chapter of the dissertation, "*Paleoenvironmental change in Pliocene eastern Africa as inferred from dental microwear texture analysis of fossil Bovidae*", takes a more regional approach to the paleoenvironments of the early australopiths. This paper adds dietary reconstructions of the fossil bovids from Kanapoi, Allia Bay and Laetoli to the Hadar sample to test previous hypotheses about changing environments at the sites. It also addresses the implications of these changes for early hominins.

The *Australopithecus anamensis* habitats at Kanapoi and Allia Bay have been reconstructed as seasonally fluctuating mosaics, probably composed of woodland forest with some incursions of grassland. The primary area of disagreement stems from the degree to which

grasslands contributed to the habitats. The bovids from these two sites generally have microwear signatures most similar to modern browsing taxa that inhabit closed forest, woodland and bushland habitats. However, a few taxa with grazing signatures do exist and this suggests that some edaphic grassland may have been present at or near the sites.

The Kanapoi bovids fall into two primary dietary categories. Five of the seven taxa closely resemble modern browser-grazer intermediate and generalist species. The prominence of browse-adapted bovids suggests a local environment with considerable forest habitat, or at least consistent access to forest resources. However, the remaining two taxa have microwear signatures that closely resemble those of modern variable grazers. The presence of these two grazing taxa indicates a more complex, mosaic paleoenvironment at Kanapoi, possibly including some grassland (Wynn, 2000; Cerling et al., 2011).

Although the fossil sample from Allia Bay is small, the microwear signatures are consistent with previous reconstructions of the bovids as forest-adapted browsers. The alcelaphin sample is most similar to modern browser-grazer intermediates and the tragelaphins are similar to dense forest adapted browsers. While the sample sizes are too small to draw definitive conclusions about this site, these results support previous reconstructions of Allia Bay as relatively closed forest, with the potential for seasonal fluctuations. Both of the fossil taxa have microwear signatures similar to modern taxa that inhabit seasonally variable environments.

Of all of the sites discussed in this dissertation, Laetoli has had the most inconsistent paleoenvironmental reconstructions, ranging from open savanna to closed forest. And the dental microwear of the fossil bovids reflects a wide variety of microwear signatures. As with Kanapoi and Allia Bay, most of the Laetoli fossil bovids have reconstructed diets similar to modern mixed feeding taxa, especially browser-grazer intermediates. However, there are also three taxa that

have microwear signatures similar to modern browsers and two that are similar to extant grazers. The increase in the ratio of grazing to browsing taxa from the earlier sites suggests more prominent incursions of open grassland habitats by the time of *A. afarensis*. All of these signals suggest that the Laetoli paleoenvironment was highly mosaic and offered a wide variety of available niches for the local fauna, including both open and closed microhabitats. This supports previous reconstructions of the site as containing multiple habitat types (Andrews, 1989; Cerling, 1992; Reed, 1997; Musiba, 1999; Harrison, 2005, Musiba et al., 2007; Su and Harrison, 2007).

Previous reconstructions have suggested that Hadar was highly mosaic, including edaphic grassland, open woodland, bushland and closed forest. Based on faunal assemblages, open habitats appear to have been more common at Hadar than at the other sites. The highly mosaic nature of the site is confirmed by the bovid microwear data. The bovids from the Sidi Hakoma and Denen Dora Members have microwear signatures like modern browsers and browser-grazers intermediates, similar to the assemblages from Kanapoi and Allia Bay. The Kada Hadar bovids include more grazing taxa, suggesting the presence of more open habitats. The largest percentage of the fossil taxa at Hadar overall are similar to mixed feeding taxa, including browser-grazer intermediates and generalists, however grazing taxa become increasingly common throughout the formation, with the largest numbers occurring in the Kada Hadar Member.

When viewing these sites over the occupation of *Australopithecus*, a general trend emerges. The dietary reconstructions reveal greater frequency of dedicated browser and/or browser-grazer intermediate taxa at the earlier sites, i.e. Kanapoi, Allia Bay and the Sidi Hakoma Member at Hadar. Later sites, particularly the Kada Hadar Member and Laetoli, have higher numbers of bovids that grouped with modern obligate or variable grazers. While none of the sites appear to have been dominated by open grassland, the results of this dissertation are consistent

with previous hypotheses that suggest a gradual aridification trend in eastern Africa during the occupation of the early australopiths.

In sum, the bovid microwear texture data agree with previous reconstructions of theaustralopith habitats as highly mosaic, with increasingly open habitats during the time of *A*. *afarensis*. This evidence of gradual aridification in eastern Africa supports previous observations by Reed (1997) that the fossil mammalian communities at these sites display increasing percentages of terrestrial species. In general, the results of this dissertation agree with the suggestion by Stanley (1992) that Pliocene hominin habitats did not change drastically over time, but rather that the morphological adaptations that characterize *Australopithecus afarensis* made them ecologically flexible enough to thrive in seasonally shifting mosaic habitats (Stern and Susman, 1983; Ward et al., 2001).

#### **Objective 3: Dental microwear as a test of taxonomic uniformitarianism**

Many studies of bovid paleoecology apply the principle of taxonomic uniformitarianism to fossil species, in an attempt to classify their dietary and habitat preferences by assuming that they are similar to closely related modern taxa (for examples, see Kappleman, 1984; Vrba, 1980, 1985, 2003; Shipman and Harris, 1988; Harris, 1991; Plummer and Bishop, 1994; Spencer, 1997; Sponheimer et al., 1999; Kovarovic et al., 2002; DeGusta and Vrba, 2003, 2005; Reed, 2008). Although this assumption is rarely questioned for recent faunas, its utility in reconstructing the paleoecology of long extinct species is uncertain. The primary problems with taxonomic uniformitarianism are that it is not applicable to fossil taxa with few or no living relatives, and that it assumes that groups remain constant in their ecological preferences over long periods of time. While taxonomic uniformitarianism is relatively straight-forward and requires minimal effort beyond a review of a faunal list, critics have questioned many of its inherent assumptions (Solounias et al., 1988; Reed, 1996; Sponheimer et al., 1999; Schubert et al., 2006). For example, in their study of fossil bovids from Makapansgat, Sponheimer et al. (1999) demonstrated that half of the included taxa had diets that differed from the assumed diets based on taxonomic uniformitarianism. For example, while *Aepyceros* sp. and *Gazella vanhoepeni* were both assumed to be mixed/seasonal feeders like their closest living relatives, carbon isotopic signatures and ecomorphological data suggested that both were obligate browsers, and showed no evidence of C4 grass consumption. Due to such discrepancies between inferred adaptations and inferred diet, it is critical that ecological similarities of extant and fossil taxa be tested in some reliable way in order to have confidence in these assumed relationships.

This study set out to test these previous dietary reconstructions for the fossil bovids of Kanapoi, Allia Bay, Laetoli and Hadar using dental microwear texture analysis. As in the Sponheimer et al. (1999) study, approximately half of the included taxa had microwear texture signatures that did not conform to expectations based on the dietary preferences of their modern counterparts. For both the Kanapoi sample, taxonomic uniformitarianism was an unreliable predictor of diet, with only three of the seven fossil taxa at Kanapoi having reconstructed diets similar to their closest living relatives. For the Allia Bay bovids, the *Tragelaphus* sp. specimens have a texture signature very similar to modern browsing tragelaphins. However, the microwear signatures of the fossil alcelaphin specimens are very different from those of modern taxa like *Damaliscus*. Taxonomic uniformitarianism was slightly more reliable at Laetoli, with five of the nine taxa having similar dietary preferences to their modern counterparts. Finally, the Hadar sample became more predictable over time. Only two of the eight fossil taxa from the Sidi Hakoma Member had microwear signatures similar to modern relatives.

more recent deposits, five of the nine Denen Dora taxa and seven of the ten Kada Hadar taxa had texture signatures that matched the dietary predictions made using taxonomic uniformitarianism.

In sum, this dissertation confirmed the opinion of Sponheimer et al. (1999), that taxonomic uniformitarianism is not always a reliable indicator of bovid diet, especially for ancient deposits. Given how frequently fossil bovid samples are used as paleoenvironmental proxies, it is critical that their diets be reconstructed accurately. This is especially important since diet can change rapidly, but it can take many generations for morphology to change. This suggests that morphology alone cannot be considered a reliable predictor of diet and should be confirmed using direct line evidence, like dental microwear or stable isotope analysis.

### **Future Goals**

My future goals related to this project are two-fold. First, I intend to expand the study to include additional and later sites in eastern Africa, such as Olduvai Gorge and Koobi Fora. The extant sample collected for this study has the potential to be applied to a many different sites and time periods in Africa. In fact, the comparative database presented here is already being used to interpret the microwear signatures of fossil tragulids from the Miocene sites of Rusinga Island and Songhor. Carbon isotope analyses are not an option for these sites, because C4 plants had not yet spread through Africa. The results suggest that ruminant dental microwear may be able to serve as a paleoenvironmental proxy at sites where stable carbon isotope data are not an option.

In the near future, I also hope to collaborate with other researchers to better integrate microwear with isotope and ecomorphological data to strengthen interpretations of diet and paleoecology. Stable isotopes and dental mesowear can provide complementary information on diet reflecting a longer time interval than microwear, which only reflects the last few days or

weeks of an individual's life. Carbon, oxygen, strontium and other isotopes hold the potential to provide a broad range of information about diets and mesowear informs on diet over the lifetime of the individual. Ecomorphology, depends on relationships between morphology and adaptation, and is another widely accepted method of dietary reconstruction that has been successful in distinguishing taxa. This provides a much longer time frame for diet, as it results from selective pressures over generations. These approaches taken together can complement one another in both the resolution of diet categorization, and provide important time-scale information for foods available and eaten, and by extension habitat setting.

The first of these projects is already underway. As part of a collaborative project with Dr. Francis Kirera, I am comparing dental microwear data of fossil bovids from the site of lleret, Kenya with stable isotope data for the same specimens. While previous studies have compared these types of data for the same taxa, none have utilized the same specimens; an important step given the short lifespan of microwear data. We hope that this project will be an important step forward in understanding the relationship between isotope and microwear data and how they can be used to complement and enhance one another.

Finally, I hope to include more diverse taxa into future studies of paleoecology and habitat. Environments are complex and not perfectly represented by any single type of inhabitant. Therefore, reconstructing the paleoecological preferences of multiple taxa should provide a more complete picture of how a past ecosystem operated. Our research group has begun applying this principle with the faunal assemblage from Langebaanweg, South Africa and to date we have analyzed the dental microwear of fossil bovids, carnivores, giraffes and elephants. In future studies, I would like to apply this multi-taxon approach to further analyze the faunal assemblages of the Hadar Formation. Hopefully, this detailed analysis would provide even

more complete information about changes in habitat availability during the occupation of

Australopithecus afarensis.

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Appendix 1: Specimen numbers and raw data for extant ta
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FMNH3447	Aepyceros	B-G	1.8399	0.0053	0.4165	5984.1	0.4788	0.75702
1	melampus	INT	825	97	16	35487	79642	9596
FMNH3447	Aepyceros	B-G	1.6287	0.0062	0.6471	6234.1	0.3267	0.51455
2	melampus	INT	54	915	08	5487	3844	5649
FMNH3447	Aepyceros	B-G	1.4031	0.0071	45.613	8126.0	0.4406	0.67371
3	melampus	INT	86	69	882	3154	46851	7485
FMNH3447	Aepyceros	B-G	1.3207	0.0052	0.4348	5648.3	0.5457	0.95544
4	melampus	INT	38	07	75	2154	98068	3019
FMNH3447	Aepyceros	B-G	2.1271	0.0030	0.3787	7328.6	0.3683	0.68968
5	melampus	INT	67	54	1	46879	47142	8248
FMNH3447	Aepyceros	B-G	1.5395	0.0023	0.8843	6139.6	0.3189	0.89709
6	melampus	INT	735	12	845	87685	01991	9476
FMNH3447	Aepyceros	B-G	2.1760	0.0033	1.5093	4926.8	0.5489	1.00883
7	melampus	INT	725	91	715	78647	25398	7187
FMNH3447	Aepyceros	B-G	1.2853	0.0010	0.8405	5306.6	0.4628	1.16809
8	melampus	INT	225	635	915	13203	94505	8188
FMNH3447	Aepyceros	B-G	1.9424	0.0023	0.5152	6280.4	0.6045	1.17598
9	melampus	INT	38	595	29	65787	46055	6342
FMNH3477	Aepyceros	B-G	2.2679	0.0014	0.5082	4872.0	0.3780	0.49113
0	melampus	INT	12	42	065	31647	07143	0727
FMNH9583	Aepyceros	B-G	2.1872	0.0049	0.4173	5139.3	0.5305	0.69106
8	melampus	INT	94	35	365	54867	61553	9594
FMNH9584	Aepyceros	B-G	1.5278	0.0037	0.6785	5791.3	0.9108	1.28970
1	melampus	INT	015	535	885	03566	72564	8096
FMNH9853	Aepyceros	B-G	2.8781	0.0050	0.2707	6138.4	0.6548	1.04727
4	melampus	INT	33	54	935	57553	25144	3161
FMNH3479	Aepyceros	B-G	1.7786	0.0043	0.5101	8304.0	0.4536	0.57588
0	melampus	INT	945	815	575	31547	60527	3342
FMNH3478	Aepyceros	B-G	2.0699	0.0027		5595.1	0.4565	0.64850
2	melampus	INT	555	76	0.5084	03155	7353	3386
FMNH1343	Aepyceros	B-G		0.0035	0.7620	6065.8	0.7561	1.04137
59	melampus	INT	2.9253	93	73	43164	73167	1792
FMNH3448	Aepyceros	B-G	1.5490	0.0023	0.4738	5603.5	0.4324	0.59261
0	melampus	INT	95	385	425	06157	13068	4069
FMNH3478	Aepyceros	B-G	1.2995	0.0080	0.5336	4408.6	0.4207	0.69779
1	melampus	INT	065	38	365	73501	12906	3045
FMNH3479	Aepyceros	B-G	1.4725	0.0043	0.4315	4291.2	0.4338	0.74081
1	melampus	INT	72	445	78	42405	26096	9063
FMNH3478	Aepyceros	B-G	2.1379	0.0037	0.5142	5381.3	0.4405	0.95813
4	melampus	INT	25	38	555	7551	50298	7737
FMNH1408	Antidorcas	B-G	2.6145	0.0038	0.1501	6806.2	0.3114	0.63875
61	marsupialis	INT	42	05	285	24023	4188	8604
FMNH3448	Antidorcas	B-G	2.2922	0.0056	0.2664	4390.5	0.8500	0.79875
2	marsupialis	INT	365	925	93	64865	87885	1564

FMNH3448	Antidorcas	B-G	1.2622	0.0039	0.7082	5708.6	0.6419	0.79837
3	marsupialis	INT	145	055	44	87808	29846	0595
FMNH3448	Antidorcas	B-G	4.8701	0.0007	0.1498	4417.1	0.4081	0.72949
8	marsupialis	INT	95	03	82	01144	87068	8909
FMNH3449	Antidorcas	B-G	2.7712	0.0044	0.8160	6470.2	0.3459	0.52225
0	marsupialis	INT	995	28	825	51084	10161	0395
FMNH3449	Antidorcas	B-G	3.2476	0.0040	0.1502	8317.6	0.4967	0.78356
4	marsupialis	INT	38	81	405	77593	67238	1875
FMNH3449	Antidorcas	B-G	1.6767	0.0038	0.4346	5376.1	0.5828	0.98158
6	marsupialis	INT	29	185	72	19861	61919	8347
FMNH3448	Antidorcas	B-G	1.5656	0.0035	1.2767	4520.8	0.7151	1.07597
7	marsupialis	INT	895	23	19	24627	57933	8167
FMNH3442	Antidorcas	B-G	2.9499	0.0038	0.1505	7106.8	0.4801	0.81314
8	marsupialis	INT	6	455	92	5564	86234	2231
FMNH3432	Antidorcas	B-G	1.2848	0.0024	0.3421	4428.7	0.6063	0.82048
8	marsupialis	INT	905	41	295	43964	44308	3526
FMNH3438	Antidorcas	B-G	1.6940	0.0055	0.5080	5390.3	0.4654	0.57060
6	marsupialis	INT	775	705	52	66566	19015	634
FMNH3439	Antidorcas	B-G	2.0641	0.0031	0.1503	6286.4	0.4071	0.69973
2	marsupialis	INT	17	465	71	63817	86675	2226
FMNH3447	Antidorcas	B-G	2.5660	0.0028	0.3421	5922.6	0.4095	0.80009
1	marsupialis	INT	63	545	1	63362	79459	7844
FMNH3447	Antidorcas	B-G	2.2329	0.0035	78.635	7836.6	0.6032	0.87806
6	marsupialis	INT	51	21	5245	07059	02071	7745
FMNH3448	Antidorcas	B-G	1.9542	0.0050	0.2666	5733.7	0.3551	0.57235
1	marsupialis	INT	65	97	325	02048	3256	5562
FMNH1297	Antidorcas	B-G	1.7942	0.0036	0.2676	4838.8	0.4371	0.69386
30	marsupialis	INT	415	325	775	3536	43884	1922
FMNH1408	Antidorcas	B-G	2.6113	0.0033	0.1504	7169.3	0.5817	0.76606
82	marsupialis	INT	54	205	445	01855	56163	9542
FMNH1408	Antidorcas	B-G	1.5701	0.0038	0.2667	8428.4	0.5176	0.90634
88	marsupialis	INT	145	145	14	13034	45294	9943
FMNH3448	Antidorcas	B-G	5.8033	0.0020	0.1502	7008.9	0.4720	0.87916
4	marsupialis	INT	59	42	85	57918	07532	9333
FMNH3449	Antidorcas	B-G	1.6578	0.0020	0.2679	5561.4	0.6459	0.89290
3	marsupialis	INT	24	04	78	37955	17378	5452
FMNH3449	Antidorcas	B-G	1.5370	0.0031	0.2673	5094.4	0.3759	0.88482
5	marsupialis	INT	595	835	78	67467	30171	7079
RMCA 83-	Cephalophus	FRU	6.6702	0.0023	0.2452	13817.	0.5530	0.81111
006M492	sylvicultor	G	57	195	39	38681	43323	9908
RMCA101	Cephalophus	FRU	3.9389	0.0013	0.2691	10142.	0.5174	0.79224
50	sylvicultor	G	43	985	015	07688	85365	9506
RMCA195	Cephalophus	FRU	5.1483	0.0040	0.2160	10881.	0.4103	0.83768
69	sylvicultor	G	26	68	635	05492	2014	8514
RMCA270	Cephalophus	FRU	5.0926	0.0037	0.2665	14158.	0.5194	0.74805
0	sylvicultor	G	73	985	385	30005	63937	9783

RMCA277	Cephalophus	FRU	4.1138	0.0048	0.2602	11272.	0.4486	0.80037
69	sylvicultor	G	935	46	28	66039	45227	1031
RMCA376	Cephalophus	FRU	4.8662	0.0023	0.2928	13820.	0.5721	0.74282
96	sylvicultor	G	96	205	76	64053	21646	1816
RMCA408	Cephalophus	FRU	3.0750	0.0040	0.3090	13081.	0.5261	0.77552
7	sylvicultor	G	255	855	895	05878	33798	2407
RMCA83-	Cephalophus	FRU	2.8322	0.0038	0.3207	9258.9	0.5328	0.81336
006M490	sylvicultor	G	415	115	975	20862	09885	6481
RMCA830	Cephalophus	FRU	3.0884	0.0024	0.2988	9552.7	0.5125	0.71041
06-M491	sylvicultor	G	285	93	395	81267	00812	9302
RMCA83-	Cephalophus	FRU	6.9214	0.0016	0.2253	14916.	0.4966	0.83077
006M493	sylvicultor	G	07	295	655	55155	71328	9121
RMCA83-	Cephalophus	FRU	3.6105	0.0031	0.1708	11762.	0.5369	0.83752
006M494	sylvicultor	G	785	68	535	78912	20863	6439
RMCA83-	Cephalophus	FRU	4.7926	0.0041	0.2846	14861.	0.4939	0.69199
006M495	sylvicultor	G	345	46	16	91373	09534	8357
RMCA83-	Cephalophus	FRU	3.4302	0.0019	0.1518	12316.	0.4853	0.77784
006M496	sylvicultor	G	775	195	705	54968	50216	9413
RMCA890	Cephalophus	FRU	4.1684	0.0013	0.2740	13326.	0.5592	0.70764
24M12	sylvicultor	G	08	9	735	36803	48572	0429
RMCA890	Cephalophus	FRU	5.7891	0.0042	0.1503	13161.	0.4983	0.80728
24M15	sylvicultor	G	505	215	23	94976	96007	8018
RMCA83-	Cephalophus	FRU	3.1033	0.0013	0.2693	10326.	0.5647	0.87184
006M483	sylvicultor	G	33	235	415	52966	07569	8007
RMCA83-	Cephalophus	FRU	5.3990	0.0040	0.2680	13813.	0.4152	0.85778
006M484	sylvicultor	G	085	375	02	58029	27932	5098
RMCA83-	Cephalophus	FRU	6.7293	0.0020	0.2584	14438.	0.5188	0.76213
006M485	sylvicultor	G	805	585	15	21677	95033	315
RMCA83-	Cephalophus	FRU	3.6431	0.0035	0.2636	13246.	0.4645	0.87264
006M486	sylvicultor	G	485	77	82	7359	24787	1064
RMCA83-	Cephalophus	FRU	5.4295	0.0039	0.2091	10922.	0.4573	0.88053
006M488	sylvicultor	G	545	47	71	42045	79418	1061
RMCA83-	Cephalophus	FRU	4.4478	0.0019	0.2332	13527.	0.4277	0.85838
006M489	sylvicultor	G	045	24	58	90933	62244	7933
RMCA83-	Cephalophus	FRU	4.7594	0.0045	0.3184	12691.	0.5601	0.78096
006M497	sylvicultor	G	625	135	59	64647	14807	5401
RMCA83-	Cephalophus	FRU	4.6280	0.0018	0.2082	9497.0	0.4263	0.81639
006M498	sylvicultor	G	135	03	335	49822	97879	333
RMCA83-	Cephalophus	FRU	6.6580	0.0030	0.2943	14279.	0.4272	0.81232
006M499	sylvicultor	G	63	035	9	2165	10612	8065
RMCA93-	Cephalophus	FRU	3.0810	0.0036	0.2668	11061.	0.4265	0.82448
006M487	sylvicultor	G	795	02	355	10869	2359	1001
FMNH1044	Damiliscus		1.1886	0.0053	1.4171	2786.2	0.4348	0.53256
28	lunatus	OG	99	155	795	19049	44591	2317
FMNH1044	Damiliscus		0.6447	0.0076	0.9079	1998.0	0.4575	0.64017
29	lunatus	OG	21	775	52	75311	14252	9566

FMNH1279	Damiliscus		0.7465	0.0064	1.6000	3527.0	0.3384	0.60685
13	lunatus	OG	805	63	965	32211	81019	8185
FMNH1279	Damiliscus		1.1139	0.0069	2.2668	3993.1	0.4355	0.52274
15	lunatus	OG	89	655	41	422	44594	6891
FMNH1279	Damiliscus		0.9089	0.0067	1.3424	3453.4	0.4607	0.64079
17	lunatus	OG	91	065	585	83693	11355	327
FMNH1279	Damiliscus		0.7608	0.0066	1.4332	3603.3	0.5617	0.69170
18	lunatus	OG	595	06	235	97394	52956	4045
FMNH1279	Damiliscus		1.2752	0.0071	1.3431	1326.6	0.4442	0.60902
19	lunatus	OG	405	05	635	15354	13312	6122
FMNH1279	Damiliscus		0.8453	0.0086	1.0658	1547.0	0.3869	0.55579
20	lunatus	OG	665	39	96	50014	82595	0278
FMNH2747	Damiliscus		0.7264	0.0053	0.8140	1906.7	0.5079	0.67830
7	lunatus	OG	58	36	125	61451	56643	7565
FMNH2952	Damiliscus		0.7337	0.0059	1.0715	3673.8	0.3954	0.54717
9	lunatus	OG	01	315	81	34317	32687	9698
FMNH2953	Damiliscus		1.3501	0.0060	1.7327	1048.8	0.3956	0.67439
1	lunatus	OG	465	535	13	69303	80164	3446
FMNH2953	Damiliscus		0.7189	0.0063	2.2086	1917.3	0.3681	0.69301
2	lunatus	OG	515	725	705	02852	46457	3021
FMNH3452	Damiliscus		0.9094	0.0065	1.2665	1785.9	0.3142	0.67063
7	lunatus	OG	97	015	825	1323	71277	7222
FMNH3948	Damiliscus		1.2257	0.0056	1.5912	1123.1	0.3803	0.64740
3	lunatus	OG	62	235	955	77318	88644	9181
FMNH3948	Damiliscus		1.1579	0.0068	2.2666	1506.6	0.4584	0.71499
9	lunatus	OG	135	56	805	92188	15489	0577
FMNH1353	Damiliscus		1.3470	0.0062	1.4342	1248.7	0.4884	0.74819
26	lunatus	OG	53	21	825	82114	55744	6099
FMNH1353	Damiliscus		0.9341	0.0060	1.1504	1270.0	0.4768	0.67332
27	lunatus	OG	945	405	86	78704	01194	2721
FMNH2748	Damiliscus		1.0385	0.0073	2.1409	2771.0	0.5227	0.76688
1	lunatus	OG	74	685	975	00033	83777	8357
FMNH3452	Damiliscus		0.7329	0.0070	1.7422	1353.0	0.3559	0.79797
6	lunatus	OG	735	385	69	9745	71412	2943
FMNH3453	Damiliscus		0.5123	0.0068	1.2689	2395.8	0.5265	0.73239
0	lunatus	OG	225	015	295	48276	60028	1462
FMNH3459	Damiliscus		1.2105	0.0059	1.4173	1934.7	0.3995	0.63051
3	lunatus	OG	41	1	67	17118	89089	3305
FMNH3459	Damiliscus		0.8555	0.0072	0.9441	1672.9	0.4671	0.67841
7	lunatus	OG	065	13	255	7646	55886	6228
FMNH1326	Damiliscus		0.6203	0.0037	1.6011	5000.8	0.3725	0.61690
5	pygargus	VG	4175	72	3	94419	89569	2725
FMNH1326	Damiliscus		1.7364	0.0045	1.4467	4529.8	0.3063	0.50439
6	pygargus	VG	035	725	525	74261	76769	929
FMNH1352	Damiliscus		1.9064	0.0031	1.2094	5504.1	0.2407	0.55733
5	pygargus	VG	13	695	175	83468	10027	4983

FMNH3456	Damiliscus		1.0751	0.0062	1.4219	5243.3	0.3346	0.65208
5	pygargus	VG	36	31	84	00845	82697	4819
FMNH3456	Damiliscus		0.9488	0.0032	1.5999	3169.2	0.2116	0.52545
6	pygargus	VG	495	835	68	54588	78184	5951
FMNH3456	Damiliscus		1.1979	0.0067	1.3442	5417.1	0.4398	0.71787
7	pygargus	VG	345	43	47	55721	97202	9219
FMNH3460	Damiliscus		1.4067	0.0057	1.2085	8450.4	0.4625	0.68603
6	pygargus	VG	81	35	95	43965	10218	258
FMNH3761	Damiliscus		1.6304	0.0035	0.9195	3424.4	0.4668	0.65118
5	pygargus	VG	03	85	94	81287	47549	7287
FMNH3915	Damiliscus		1.2987	0.0066	1.4646	4456.4	0.4633	0.77987
4	pygargus	VG	285	97	565	54168	20476	9943
AMNH829	Damiliscus		0.8092	0.0042	1.2826	8959.3	0.3093	0.51225
84	pygargus	VG	9645	5	1875	98296	30231	6858
AMNH826	Damiliscus		1.6916	0.0064	1.4386	3470.3	0.3074	0.59738
48	pygargus	VG	24	31	29	29225	23142	2057
AMNH835	Damiliscus		1.5114	0.0034	1.7440	4156.6	0.2835	0.58500
94	pygargus	VG	77	37	655	87159	08085	9497
AMNH826	Damiliscus		0.9378	0.0062	1.3610	5923.8	0.3942	0.62692
49	pygargus	VG	26	885	975	71279	71213	8963
AMNH875	Damiliscus		0.9322	0.0040	1.3149	4411.8	0.4852	0.69027
94	pygargus	VG	015	425	735	68577	06935	5479
AMNH834	Damiliscus	10	1.2928	0.0066	1.2693	3590.9	0.4292	0.60107
59	pygargus	VG	46	495	455	99473	59446	0706
AMNH845	Damiliscus	10	1 6310	0.0035	1 3857	1690 8	0 3603	0.59014
97	nygargus	VG	68	49	49	89911	95185	3106
NMNH164	Damiliscus		0.8782	0.0058	1.1104	4002.6	0.4605	0.67193
83	pygargus	VG	8	655	68	31669	05675	488
NMNH184	Damiliscus	. –	1.3018	0.0032	1.2681	5068.6	0.4716	0.61902
95	nygargus	VG	03	295	495	06153	07888	1708
NMNH198	Damiliscus		1.2369	0.0066	0.9334	5986.7	0.4700	0.52880
42	pygargus	VG	52	2	155	9752	40061	6628
NMNH203	Damiliscus	. –	1.0245			4833.1	0.3842	0.51291
48	pygargus	VG	4	0.0061	1.2654	09736	82489	107
NMNH216	Damiliscus		2 1 3 6 4	0.0039	1 3704	6597 1	0 4953	0 64386
48	nygargus	VG	8	625	85	62828	22058	4474
NMNH216	Damiliscus	10	0.9846	0.0066	0.9999	6403.2	0.2275	0.50491
47	pygargus	VG	17	055	155	05833	77232	5139
FMNH1279	r <i>J 8</i> 8		2 3680	0 0052	0 1500	2699 6	0 4505	0 87641
32	Gazella granti	VG	835	79	785	57586	24852	4888
FMNH1343	Sulfina Brandi	10	1 1 3 9 3	0 0044	0 6643	6374 3	0 4869	0 74495
13	Gazella granti	VG	815	7	135	142	23834	4415
FMNH2959	Surgering Brandt	. 0	1.9548	0.0040	0.2665	5177.0	0.3403	0.68766
5	Gazella oranti	VG	485	635	07	31861	30289	3969
FMNH2959	Sucona Brand	, 5	1 3029	0 0037	0 2664	3052.5	0 3937	0 87893
6	Gazella granti	VG	47	38	865	66522	19771	847
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FMNH2959			1.9346	0.0043	0.2087	2340.8	0.2986	0.51622
8	Gazella granti	VG	49	16	325	64853	44873	4572
FMNH2959	C		1.8160	0.0033	0.6086	4743.6	0.4914	0.64015
9	Gazella granti	VG	155	985	515	94918	93389	0466
FMNH2960	C		1.4811	0.0041	0.5478	4345.8	0.2893	0.81407
0	Gazella granti	VG	555	355	41	94489	6413	5934
FMNH2960	U		1.0129	0.0055	1.0352	3496.9	0.4395	0.51918
4	Gazella granti	VG	6	87	85	84815	4105	365
FMNH2960	U		1.6476	0.0057	0.5322	2459.0	0.3773	0.73702
5	Gazella granti	VG	28	85	895	57563	6376	3683
FMNH3291	8		1.9297	0.0057	0.5499	1511.1	0.3421	0.57080
1	Gazella granti	VG	81	51	715	14275	08431	7613
FMNH3291	Surrena Brana	10	1 9977	0.0031	0 2083	3070.0	0 4146	0 63278
3	Gazella granti	VG	925	245	235	02093	68609	7295
FMNH3291	Surrena Brana	10	/	0.0033	0 5996	1889 1	0 3640	0 50811
4	Gazella oranti	VG	0 9071	925	475	93374	84873	037
RMCA83-	Guzenia granti	10	2 0317	120	0 1 5 0 0	24727	0 3048	0 57184
006M795	Gazella granti	VG	085	0.0039	675	39663	81122	0692
RMCA83-	Guzenia granti	10	2 1830	0.0050	1 2158	2073.6	0 3883	0 76971
006M496	Gazella oranti	VG	69	37	495	42176	05524	6113
RMCA83-	Guzena grann	٧U	2 2539	0.0033	775	1746.6	03397	0 73145
006M794	Gazella oranti	VG	2.2557	975	0.8312	95454	79332	8226
RMCA80-	Guzena grann	٧U	1 6367	0.0042	0.0512	33323	0 3976	0 51804
006-M487	Gazella oranti	VG	535	96	0.102)	9179	04125	7342
RMCA83-	Guzenia granti	10	1 1711	0.0055	0.8367	4866 3	0 2912	0 59153
006M846	Gazella oranti	VG	41	565	78	55263	21259	5544
FMNH8462	Guzenia granti	10	1 4771	0.0032	0 2085	2331.7	0 2233	0 79266
8	Gazella oranti	VG	245	56	625	25165	99203	5717
FMNH4978	Guzenia granti	10	2 0069	0.0035	0 4085	4253 3	0.2616	0 47627
1	Gazella oranti	VG	2.000)	81	115	64348	34677	8085
FMNH2648	Guzenia granti	10	1 1 3 7 6	0.0036	0 2668	3456 1	0 3698	0 73061
7	Gazella oranti	VG	83	585	21	33165	33977	7751
, FMNH4987	Guzenia granti	10	2 1452	0.0043	0 3424	3096.9	0 3004	0 75593
5	Gazella oranti	VG	78	335	35	314	69356	7048
FMNH8264	Guzenia granti	10	1 2626	0.0046	0 2084	3120.9	0 4353	0 78414
8	Gazella oranti	VG	03	34	21	16915	88721	9931
FMNH8462	Guzena grann	٧U	2 0055	0 0045	0 6054	4023.2	0.3157	0.63910
5	Gazella oranti	VG	2.0055 875	0.0015	7585	20215	93319	0273
FMNH2649	Guzenia granti	10	1 0659	0.0038	0.9627	1353.6	0 2438	0.69155
8	Gazella oranti	VG	675	18	325	36387	67804	1513
FMNH2684	Guzenia granti	10	2 1419	0.0037	0.9659	2540.5	0 3910	0 72947
5	Gazella granti	VG	11	285	0.9039	2040.5 96105	46442	2791
2	Hinnotrague	, U	1 7634	0 0055	0 4331	5013.9	0 4613	0 75580
FMNH1166	niger	VG	745	315	795	10017	77743	1343
FMNH1343	Hinnotrame	10	1 3902	0 0063	0 1522	4288 1	0 3 9 3 4	0 73983
60	niger	VG	385	385	105	08204	76197	5765
00	111601	, 0	505	505	105	00207	10174	5705
FMNH3451	Hippotragus		1.5387	0.0054	0.2082	6877.0	0.6520	0.85565
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4	niger	VG	23	19	55	24095	76392	9588
FMNH3451	Hippotragus		0.1923	0.0038	0.1505	3042.6	0.3852	0.76916
7	niger	VG	125	31	185	71161	65309	8815
FMNH3460	Hippotragus		1.8166	0.0049	0.1518	4929.2	0.5059	0.92958
5	niger	VG	1	575	45	76177	84396	7744
FMNH9811	Hippotragus		2.1091	0.0046	0.1507	3904.4	0.5357	0.82091
1	niger	VG	19	845	065	04539	95297	5995
FMNH5648	Hippotragus		1.5302	0.0079	0.4353	5534.0	0.4541	0.77957
2	niger	VG	15	47	99	58008	43633	1034
FMNH9874	Hippotragus		1.6811	0.0065	0.3422	4846.0	0.4711	0.58632
5	niger	VG	9	945	015	0692	74223	1932
FMNH6486	Hippotragus		1.1473	0.0047	0.2089	6227.5	0.5478	0.69794
3	niger	VG	055	82	555	37395	57128	3873
FMNH4978	Hippotragus		1.0514	0.0055	0.2673	3493.6	0.5232	0.62770
5	niger	VG	755	04	73	86877	54273	0764
FMNH1648	Hippotragus		1.7542	0.0075	0.2668	3921.2	0.5724	0.64091
6	niger	VG	775	72	95	88865	54879	0376
FMNH2648	Hippotragus		1.7893	0.0074	0.1504	4569.8	0.6446	0.88685
5	niger	VG	92	22	31	7071	74304	0159
FMNH9751	Hippotragus		1.1794	0.0062	0.3416	3998.1	0.5793	0.81375
6	niger	VG	79	485	71	87867	41123	6889
FMNH1648	Hippotragus		0.9151	0.0071	0.3431	4049.3	0.5108	0.83345
8	niger	VG	65	43	15	57266	73467	0651
AMNH978	Hippotragus		1.4955	0.0072	0.3416	6650.0	0.5704	0.71708
46	niger	VG	175	045	845	19712	41267	9761
AMNH364	Hippotragus		1.6825	0.0063	0.4193	4582.1	0.5996	0.65347
84	niger	VG	79	825	985	70104	19872	2897
AMNH648	Hippotragus		0.9196	0.0064	0.2712	5892.6	0.4200	0.74155
46	niger	VG	43	755	91	82569	55571	7875
AMNH795	Hippotragus		2.1532	0.0078	0.3418	4716.8	0.3838	0.67339
48	niger	VG	515	185	04	94046	04785	3367
AMNH154	Hippotragus		0.8257	0.0038	0.1506	5091.6	0.6219	0.73627
86	niger	VG	24	58	695	42845	29908	2299
AMNH648	Hippotragus		2.1544	0.0041	0.1503	6193.5	0.5800	0.75127
56	niger	VG	72	635	31	43239	93862	4757
AMNH849	Hippotragus		1.4657	0.0076	0.2671	5567.1	0.6706	0.73650
75	niger	VG	535	65	77	64694	85775	0097
AMNH294	Hippotragus		1.2814	0.0052	0.4184	5170.4	0.5499	0.59424
56	niger	VG	52	79	545	12253	95627	223
AMNH874	Hippotragus		1.8369	0.0066	0.2668	6479.3	0.4289	0.56335
96	niger	VG	86	77	59	2915	7573	2409
	Kobus							
FMNH1279	ellipsyprymn		1.5380	0.0058	0.8665	3113.3	0.5365	0.67308
46	us	VG	99	695	51	24437	30358	6554
FMNH1279	Kobus	VG	1.9052	0.0037	0.7007	4028.7	1.1780	2.71162

48	ellipsyprymn us		875	175	375	88714	31612	2225
	Kobus							
FMNH1279	ellipsyprymn		2.3850	0.0058	0.9429	4321.2	0.8569	1.30491
52	us	VG	805	37	4	08131	62042	7308
	Kobus							
FMNH1279	ellipsyprymn		1.3479	0.0060	0.7667	4330.6	0.3342	1.06947
53	us	VG	535	17	4	12514	24022	6172
	Kobus							
FMNH2066	ellipsyprymn		0.8044	0.0069	0.7244	3977.3	0.7077	1.17907
1	us	VG	825	25	03	4692	60679	8383
	Kobus							
FMNH2432	ellipsyprymn		1.1496	0.0052	0.8350	3081.4	0.8155	1.30365
1	us	VG	78	225	02	49793	07193	912
	Kobus							
FMNH2432	ellipsyprymn		0.6035	0.0058	0.9770	1459.4	0.4980	0.57978
2	us	VG	49	765	875	35166	97047	0556
	Kobus							
FMNH2719	ellipsyprymn		1.3979	0.0063	0.9437	5345.9	0.5280	0.76592
9	us	VG	21	98	72	06738	96761	3418
	Kobus							
FMNH2954	ellipsyprymn		1.6799	0.0059	0.8321	5389.4	1.3543	1.59748
2	us	VG	215	705	485	69843	17774	047
	Kobus							
FMNH8543	ellipsyprymn		0.6943	0.0068	0.6167	3099.9	0.7361	1.16550
3	us	VG	705	085	92	65436	76824	3274
	Kobus							
FMNH6485	ellipsyprymn		1.7401	0.0057	0.9738	3787.5	0.8875	3.82494
6	us	VG	065	36	16	82828	85653	9696
	Kobus							
NMNH874	ellipsyprymn		0.9931	0.0062	0.7617	2449.0	0.5107	0.82438
92	us	VG	97	265	59	34552	66054	7615
	Kobus							
NMNH216	ellipsyprymn		1.5075	0.0065	0.6417	2217.2	0.4652	0.92970
48	us	VG	095	175	585	41508	48596	31
	Kobus							
NMNH975	ellipsyprymn		1.7829	0.0060	0.8830	4881.9	0.3629	0.78171
16	us	VG	445	57	15	27964	44884	79
	Kobus							
NMNH784	ellipsyprymn		1.2048	0.0058	0.6011	2571.6	0.5685	0.76180
68	us	VG	78	055	48	33798	3676	3619
	Kobus							
	ellipsyprymn		1.3977	0.0057	0.7082	2837.5	0.4404	0.75490
FMNH1091	us	VG	085	775	09	07848	30078	4722
FMNH2720	Kobus		1.3746	0.0044	0.6172	5066.7	0.3340	0.70357
0	ellipsyprymn	VG	94	12	13	45343	18466	5632

	us							
	Kobus							
	ellipsyprymn		0.9831	0.0051	0.9419	3159.4	0.5760	0.69955
FMNH7225	us	VG	39	505	58	81996	158	5967
	Kobus		• •					• • • • •
FMNH8543	ellipsyprymp		0 8553	0.0065	0 8004	3073 5	0 5338	0 62778
2	us	VG	535	235	705	39046	53664	5919
<i>L</i>	Kobus	.0	555	255	105	57010	55001	5717
FMNH8544	ellinsvnrvmn		1 1898	0.0070	0 7083	4483.6	0 3468	0 62777
1	us	VG	405	0.0070 <u>/</u> 9	425	00501	25901	7283
1	Kobus	٧U	705	<b>ر</b> ۲	723	00501	25701	1205
NMNH487	allingunrumn		0 5001	0.0067	0.0178	25121	0.6012	0 83730
56	cmpsyprynni	VC	61	0.0007 59	0.9170	2342.1	0.0912	0.03739
50	us Vobus	VU	01	30	21	00079	92203	9019
	NOUUS		1 4222	0.0066	0 0000	4170 2	0 4174	0 60576
ΙΝΙVΙΙΝΠΟΙ Ζ 5 Δ	empsyprynni	VC	1.4522	0.0000	0.0990	41/0.2	0.41/4 52405	0.08570
54 ENDUL4697	us	٧G	0.02(5	803	82 1.5215	03041	52495	/008
FMINH468/	IZ 1 1 1	00	0.8265	0.0057	1.5315	3468.3	0.4215	0.83135
65 ENDU1254	Kobus leche	ŪĠ	48	0.005/	4	2153	4	4
FMNH1354	77 1 1 1	00	1.2648	0.0071	1.0321	16/8.3	0.3865	0./6545
6	Kobus leche	OG	4	0.0071	5	6	45	684
FMNH2368			0.9874		0.9654	3268.3	0.3587	0.83215
4	Kobus leche	OG	564	0.0046	1352	215	5	4
FMNH3498			0.6898		0.8984	4685.1	0.3105	0.63215
7	Kobus leche	OG	745	0.0051	521	32	48	4
FMNH6543			0.9874		1.5321	1735.3	0.4598	
5	Kobus leche	OG	51	0.0058	354	215	75	0.73154
FMNH5215			1.1032		1.6684	3594.1	0.4265	0.99876
4	Kobus leche	OG	5	0.0065	5	8	48	54
FMNH4875			0.7841		2.1345	1832.1	0.4688	1.03154
4	Kobus leche	OG	51	0.0049	4	3125	45	5
FMNH1656			0.8894		1.8645	2975.1	0.2913	0.64315
84	Kobus leche	OG	51	0.0058	6	6	545	4
FMNH5595			1.0254		1.3315	2384.2	0.3615	0.73213
5	Kobus leche	OG	86	0.0055	4	721	648	54
FMNH1648			0.8654		2.1354	1972.1	0.3246	0.69778
45	Kobus leche	OG	8	0.0047	54	4	87	45
FMNH4978			0.9751		0.9984	3942.3	0.2836	0.63213
4	Kobus leche	OG	54	0.0067	51	7815	45	5
FMNH1648			1.5687		0.9135	1267.3	0.4132	
5	Kobus leche	OG	8	0.0061	154	215	36	0.73245
FMNH1648			0.6987		1.0351	2975.1	0.4526	0.83546
76	Kobus leche	OG	465	0.0048	35	924	48	8
FMNH1984			0.9845		1.1684	4268.3	0.3668	0.79876
51	Kobus leche	OG	1	0.0055	6	215	78	4
FMNH7648			1.4321		1.7645	2781 2	0.4126	0.83215
13	Kobus leche	OG	5	0.0062	1	1	54	4
			-		-	-		•

FMNH6846			1.2654			4038.6	0.3031	0.68784
1	Kobus leche	OG	8	0.0059	2.1648	548	354	5
FMNH7846			0.8784		2.3684	3058.2	0.4265	
1	Kobus leche	OG	151	0.0051	51	15	48	1.0322
FMNH9784			0.9321		1.4687	1682.1	0.3313	0.72134
6	Kobus leche	OG	65	0.0064	751	824	54	5
AMNH716			0.7654		2.8775	3451.1	0.3946	
4	Kobus leche	OG	8	0.0049	1	214	545	0.73648
AMNH164			1.5321		2.0654	2854.3	0.4031	0.83654
8	Kobus leche	OG	54	0.0072	84	215	5	51
AMNH795			0.7321			1753.2		0.73213
3	Kobus leche	OG	534	0.0049	0.9874	15	0.3487	54
AMNH487			0.5987		1.6984	4038.1	0.4568	0.96545
8	Kobus leche	OG	845	0.0058	51	654	786	1
AMNH197					2.1684	3951.3	0.3687	0.83215
84	Kobus leche	OG	1.1548	0.0066	5	21	454	3
AMNH768			0.7321		1.5841	2267.1	0.3265	0.68784
48	Kobus leche	OG	54	0.0052	151	324	48	53
RMCA83-	Litocranius	BRO			1.1345	13594.	0.7651	
008M485	walleri	W	4.7548	0.0019	4	32156	5	1.1548
RMCA83-	Litocranius	BRO			1.3684	9857.3	0.6215	0.96354
008M468	walleri	W	3.2648	0.003	52	2156	4	32
RMCA83-	Litocranius	BRO	2.2651		0.5874	12684.	1.1654	
008M483	walleri	W	68	0.0028	51	1654	8	1.2548
RMCA83-	Litocranius	BRO	3.6548		0.8645	14875.	1.0654	1.53213
008M451	walleri	W	78	0.0034	21	32156	8	5
RMCA83-	Litocranius	BRO	, .		0 4684	10685	0 5874	0 96543
008M489	walleri	W	1.9874	0.0028	51	15468	51	2
RMCA83-	Litocranius	BRO	2 1654		0 7645	8627 3	0 8857	0 86846
008M462	walleri	W	8	0.0017	1	5468	451	5
RMCA83-	Litocranius	BRO	4 3654	010017	1 1 3 5 4	11975	0 7684	C
008M493	walleri	W	8	0 0022	5	1654	5	0 9315
RMCA83-	Litocranius	BRO	3 8484	0.00	0 9874	13294	0 5684	0.7010
008M459	walleri	W	65	0.0018	51	6548	5	1 1 5 4 8
RMCA83-	Litocranius	BRO	5 4987	0.0010	0.8132	10584	0 8645	1.10 10
008M454	walleri	W	8	0.0029	12	3156	3	1 33548
RMCA83-	Litocranius	BRO	3 0321	0.002)	0 5897	12394	0 5874	0.83203
008M491	walleri	W	68	0.0021	451	32156	451	25
RMCA83-	Litocranius	BRO	4 3165	0.0021	1 0354	9751 3	0.8354	20
008M492	walleri	W	48	0.0018	5	215	8	0 96545
RMC 483-	Litocranius	BRO	3 6548	0.0010	0 6845	14952	0 5845	0.97845
008M456	walleri	W	5.0540 7	0.0016	1	1354	31	1
BMC 483-	Litocranius	BRO	1 3878	0.0010	0 9645	1168/	0.8684	1
006M472	wallari	W	л.3070 Л5	0.0025	12	32156	Q/	1 065/19
RMC A 83	Litocranius	BBO	тл Л 265Л	0.0025	0.887/	082130	0 0125	1.00040
006M407	wallori		4.2034	0.002	0.00/4 51	7024.J 7165	0.9155 15	1 22510
00010149/	wallell	vv	70	0.005	51	2103	40	1.23348

FMNH6487	Litocranius	BRO	1.8784		1.1354	8734.1		1.43213
6	walleri	W	5	0.0029	8	354	1.0325	5
FMNH1354	Litocranius	BRO	3.1265		1.4987	12846.	0.6321	
8	walleri	W	468	0.0027	456	1654	354	0.96543
FMNH4687	Litocranius	BRO	2.2654		1.3654	8736.6		0.99842
7	walleri	W	8	0.002	8	548	0.7315	3
FMNH1446	Litocranius	BRO	3.7654		0.5315	9521.3	0.5984	0.89845
5	walleri	W	65	0.0018	4	215	51	1
AMNH164	Litocranius	BRO	2.6548		0.8968	10684.	0.6215	
854	walleri	W	6	0.0032	452	1654	485	1.0326
AMNH164	Litocranius	BRO	5.8798		1.2254	9824.6	1.0321	
859	walleri	W	4	0.0016	86	5468	53	1.36848
AMNH164	Litocranius	BRO	2.2654		2.0354	10834.	0.6321	
853	walleri	W	8	0.0021	35	6548	57	1.26548
AMNH164	Litocranius	BRO	3.6876		0.6984	12168.	0.5832	0.96545
785	walleri	W	845	0.0031	5	65465	435	2
AMNH164	Litocranius	BRO	3.0265		0.9684	11762.	0.5987	0.99423
798	walleri	W	4	0.0024	521	1354	84	15
AMNH164	Litocranius	BRO	2.0265		0.8654	8762.6	0.6987	0.89874
856	walleri	W	48	0.0028	1321	5846	4512	5
AMNH164	Litocranius	BRO	3.2135		1.2354	9762.1	0.5987	0.83321
795	walleri	W	46	0.0014	5	354	854	5
NMNH454	Neotragus	BRO	4.2651		1.1236	9587.2	0.5931	
86	batesi	W	54	0.0019	5	315	5	0.96545
NMNH467	Neotragus	BRO				11268.	0.4318	
84	batesi	W	2.6568	0.0026	0.7645	26548	54	1.0654
NMNH164	Neotragus	BRO	5.2154		0.6874	9123.3	0.5214	
856	batesi	W	6	0.0021	51	48	8	0.86542
NMNH164	Neotragus	BRO	4.1315		0.8642	8395.2	0.6842	
875	batesi	W	64	0.0017	1	354	15	0.98451
NMNH464	Neotragus	BRO	2.1254		0.9154	12759.	0.4325	0.81354
856	batesi	W	68	0.002	8	2156	48	6
NMNH497	Neotragus	BRO			1.0354	12157.	0.5214	
846	batesi	W	5.2356	0.0026	4	1654	6468	1.16548
NMNH264	Neotragus	BRO			1.3875	10268.	0.5468	0.95132
856	batesi	W	4.1345	0.0027	1	35468	78	15
NMNH164	Neotragus	BRO	2.6548		0.5321	8628.3	0.6254	
887	batesi	W	7	0.0021	51	215	8	0.8545
NMNH458	Neotragus	BRO	5.1321		0.8354	9258.3	0.6865	
16	batesi	W	54	0.0019	12	215	421	1.26548
NMNH464	Neotragus	BRO	3.6468		0.6874	10284.		
857	batesi	W	7	0.003	51	1324	0.7315	1.0654
NMNH459	Neotragus	BRO	4.9875		0.9584	9685.5	0.6315	0.94561
78	batesi	W	4	0.0024	1	468	4	23
NMNH164	Neotragus	BRO	4.1213		0.5845	12584.		
872	batesi	W	54	0.0017	1	16547	0.8657	0.93245

NMNH164	Neotragus	BRO	2.3213		0.6845	9851.3	0.4562	0.86543
978	batesi	W	54	0.0014	1	215	132	2
NMNH487	Neotragus	BRO	5.1321		0.4874	8627.3	0.4987	0.91324
26	batesi	W	56	0.0028	51	2156	561	15
NMNH597	Neotragus	BRO	3.8768		0.6684	7925.6	0.7135	
82	batesi	W	4	0.0025	521	548	48	0.93215
NMNH465	Neotragus	BRO	2.2326		0.8351	12958.	0.5987	
87	batesi	W	54	0.0022	321	1546	5	0.79451
NMNH132	Neotragus	BRO	4.3215		0.6845	10285.	0.6987	0.83215
648	batesi	W	4	0.0019	132	6548	51	34
NMNH134	Neotragus	BRO	2.1248		1.0654	8637.1	0.6154	
856	batesi	W	74	0.0021	65	65465	8	0.93215
NMNH487	Neotragus	BRO	3.6548		1.6321	9875.3	0.5432	1.23548
92	batesi	W	7	0.0028	354	21564	49	5
NMNH465	Neotragus	BRO	3.2354		0.5687	10975.	0.6984	
97	batesi	W	8	0.0031	45	1354	5	1.06548
NMNH164	Neotragus	BRO	5.1354		0.4987	12752.	0.5315	
875	batesi	W	87	0.0027	451	3215	4	0.92154
NMNH164	Neotragus	BRO	2.9848		0.9654	11695.	0.7122	0.93215
972	batesi	W	7	0.0016	121	11354	36	4
NMNH164	Neotragus	BRO	4.8321		0.6845	8267.2	0.4725	
978	batesi	W	564	0.0012	21	6468	45	0.86545
AMNH808	Oreotragus		2.0907	0.0045	0.5370	9315.1	0.5101	0.82124
84	oreotragus	GEN	745	775	62	35714	36769	964
AMNH810	Oreotragus	0 DI V	1 8636	0.0031	0 2668	7670.8	0 5814	0 90694
23	oreotragus	GEN	5	67	755	69766	45087	9342
AMNH810	Oreotragus	0 DI V	2 1900	0.0030	0 2666	5792.2	0.5177	0 76822
28	oreotragus	GEN	26	12	695	69424	71683	6966
FMNH1109	Oreotragus	0 DI V	2 8989	0.0038	0 2124	7681.5	0 2504	0 62498
45	oreotragus	GEN	<u>-</u> .0909 7	735	255	82914	04112	469
FMNH1350	Oreotragus	GEI	,	0.0036	0 4522	9452.8	0 4720	0 67589
81	oreotragus	GEN	2 481	4	755	94277	90453	0933
FMNH1958	Oreotragus	GER	1 9664	0.0032	0 1 5 0 5	9152.0	0 4896	1 28368
2	oreotragus	GEN	89	975	96	92558	34071	7617
FMNH1958	Oreotragus	OLIV	1 4100	0 0047	0 3080	12976	0 4482	0.91202
3	oreotragus	GEN	885	75	55	39999	24865	8384
FMNH1961	Oreotragus	ULIV	2 2550	0 0029	0 4014	7073 1	0 4180	0 52139
3	oreotragus	GEN	055	0.002) 47	51	45954	0118	9613
5 FMNH2695	Oreotragus	ULIV	1 6806	0 0024	0 2385	8840.6	0 4379	0 64033
7	oreotragus	GEN	515	52	345	34245	158	1447
, FMNH2606	Oreotragus	ULIV	2 7360	0.0032	0 5427	7815.6	0 / 0 3 5	0.61748
0	oreotragus	GEN	385	305	57	64	30145	1222
FMNH2716	Oreotragus	ULIN	1 7027	0.00/1	0 2456	11614	0 2608	0 67768
6	oreotragus	GEN	005	0.0041	285	3/505	0.3090	6525
0	Oreotragus	ULIN	1 2125	000	20J 0 5577	12/05	07550	1 06050
EMNILI7470	orootrogus	CEN	1.2133 905	0.0020 07	0.5522	12403.	0.4393 1220	7050
$\Gamma_{\rm WIIN} \Pi / 4 / 0$	oreotragus	UEIN	000	0/	233	07/30	4337	1030

FMNH8100	Oreotragus		2.3979	0.0046	0.2103	10781.	0.3158	0.67652
8	oreotragus	GEN	205	645	54	72619	6269	6952
AMNH978	Oreotragus		2.0907	0.0036	0.3470	8315.1	0.3101	0.83624
47	oreotragus	GEN	745	775	62	35714	36769	964
AMNH948	Oreotragus		1.9873	0.0030	0.5768	8270.7	0.4914	0.92149
78	oreotragus	GEN	65	67	755	89766	45087	3423
AMNH134	Oreotragus		2.3900	0.0051	0.2766	6892.2	0.4377	0.77522
86	oreotragus	GEN	26	12	695	79424	71683	6966
AMNH648	Oreotragus		1.9989	0.0037	0.2324	8681.6	0.3104	0.63519
51	oreotragus	GEN	7	735	255	32914	04112	8469
AMNH648	Oreotragus			0.0029	0.4822	11552.	0.4930	0.69528
53	oreotragus	GEN	2.4681	4	755	91428	90453	9093
AMNH216	Oreotragus		3.8664	0.0030	0.1905	8162.9	0.5196	1.13368
48	oreotragus	GEN	89	975	96	02558	34071	7617
AMNH978	Oreotragus		2.3200	0.0028	0.3180	10976.	0.4434	0.93602
16	oreotragus	GEN	885	75	55	44999	24865	8384
AMNH214	Oreotragus		2.2550	0.0028	0.4114	7513.1	0.4283	0.68599
78	oreotragus	GEN	055	47	51	65954	0118	6135
AMNH164	Oreotragus		3.1806	0.0045	0.2485	8652.7	0.4429	0.67893
83	oreotragus	GEN	515	52	345	54245	158	3145
AMNH648	Oreotragus		2.6169	0.0033	0.3327	8521.6	0.2985	0.71658
56	oreotragus	GEN	385	395	52	74	30145	1233
AMNH784	Oreotragus		1.6837	0.0046	0.4556	7295.3	0.5028	0.82368
26	oreotragus	GEN	905	035	285	35047	07556	6525
	Philantomba	FRU	4.1423	0.0019	0.2667	15935.	0.6308	0.60257
FMNH1289	monticola	G	88	835	615	25756	27058	6061
FMNH1343	Philantomba	FRU	3.6635	0.0012	0.2679	12983.	0.6227	0.85129
11	monticola	G	14	18	59	95727	48968	3954
FMNH1353	Philantomba	FRU	5.8825	0.0022	0.2561	14533.	0.4676	0.79326
22	monticola	G	3	345	075	073	42749	2144
FMNH1866	Philantomba	FRU	6.1846	0.0019	0.1513	12214.	0.4416	0.91833
72	monticola	G	16	65	445	82153	89594	0539
FMNH2754	Philantomba	FRU	3.6726	0.0020	0.1501	14335.	0.4949	0.62915
3	monticola	G	5	345	08	81519	58047	91
FMNH2754	Philantomba	FRU	6.3192	0.0030	0.2097	16135.	0.3708	0.83630
8	monticola	G	125	955	585	71384	27978	1541
FMNH3428	Philantomba	FRU	5.2394	0.0016	0.1499	16378.	0.6500	0.71295
1	monticola	G	585	3	53	51354	6411	1645
FMNH3428	Philantomba	FRU	4.4326	0.0014	0.1530	12860.	0.5545	0.62039
5	monticola	G	42	58	12	79432	39415	423
FMNH8160	Philantomba	FRU	4.1137	0.0023	0.2358	14597.	0.6293	0.88700
3	monticola	G	22	68	625	5401	79945	564
FMNH8716	Philantomba	FRU	3.7500	0.0024	0.1366	15507.	0.4839	1.05677
4	monticola	G	27	105	405	25694	57163	5105
FMNH1648	Philantomba	FRU	3.6478	0.0020	0.1509	13611.	0.6347	0.69675
5	monticola	G	98	255	735	192	26461	5112

FMNH8716	Philantomba	FRU	5.2941	0.0022	0.2243	12182.	0.5940	0.77388
9	monticola	G	865	38	835	95993	02336	3095
FMNH8726	Philantomba	FRU	5.5493	0.0030	0.2101	16884.	0.4924	1.01918
4	monticola	G	825	005	255	98806	71527	007
RMCA83-	Philantomba	FRU	4.2144		0.1856	17852.	0.6325	0.85479
006M972	monticola	G	51	0.0018	32	3625	69	26
RMCA83-	Philantomba	FRU	3.8955		0.2035	16598.	0.5754	0.86592
006M267	monticola	G	4	0.0023	14	2654	48	1
RMCA83-	Philantomba	FRU	3.1215		0.1568	11658.	0.5695	0.76985
006M384	monticola	G	48	0.0011	47	21	87	2
RMCA83-	Philantomba	FRU	5.9845		0.1895	15358.	0.5148	0.91235
006M982	monticola	G	1	0.0024	62	0235	7	4
RMCA83-	Philantomba	FRU	3.2541		0.1654	14526.	0.6695	0.76582
006M458	monticola	G	7	0.0012	78	3258	81	14
RMCA83-	Philantomba	FRU	5.1235		0.2256	15265.	0.4985	0.87562
006M973	monticola	G	48	0.0023	8	2565	21	4
RMCA83-	Philantomba	FRU	4.3658		0.2685	11256.	0.6526	0.84526
006M977	monticola	G	9	0.0016	47	36547	17	3
RMCA83-	Philantomba	FRU	3.2036		0.2541	16547.	0.3958	0.78652
006984	monticola	G	5	0.0017	665	2658	71	14
FMNH1109	Raphicerus	B-G	2.8182	0.0021	0.5664	6915.3	0.2709	0.67158
46	campestris	INT	94	5	675	73691	64816	1814
FMNH1279	Raphicerus	B-G	2.5197	0.0014	0.5674	8420.8	0.5636	0.76712
77	campestris	INT	675	86	305	106	95494	3972
FMNH1343	Raphicerus	B-G	2.0438	0.0036	0.6518	7471.5	0.4049	1.05561
64	campestris	INT	16	335	245	82682	19932	9315
FMNH1343	Raphicerus	B-G	2.1216	0.0024	0.8501	5705.0	0.4319	0.85738
65	campestris	INT	66	325	715	33737	20484	1374
FMNH1408	Raphicerus	B-G	3.2663	0.0023	0.4174	10693.	0.4926	0.67081
59	campestris	INT	5415	175	165	41693	38011	4704
FMNH1782	Raphicerus	B-G	2.4764	0.0025	0.7091	10791.	0.3867	0.94093
7	campestris	INT	695	69	815	35374	28206	6359
FMNH3450	Raphicerus	B-G	2.0576	0.0023	0.7086	9944.2	0.5305	1.13919
9	campestris	INT	685	475	775	23036	96696	729
FMNH3451	Raphicerus	B-G	2.6467	0.0025	0.6679	7648.6	0.5266	0.95393
0	campestris	INT	335	515	62	28919	52519	5446
FMNH3451	Raphicerus	B-G	1.7586	0.0025	0.8086	8724.6	0.5280	0.52244
1	campestris	INT	98	365	46	45868	71125	3075
FMNH8599	Raphicerus	B-G	1.5751	0.0039	0.6082	9724.1	0.4108	0.70504
9	campestris	INT	215	285	135	58241	46428	4282
FMNH5495	Raphicerus	B-G	2.8500	0.0034	0.5689	8293.1	0.4160	0.95946
6	campestris	INT	27	35	075	78717	54306	7871
FMNH1648	Raphicerus	B-G	1.5574	0.0023	0.8571	10545.	0.5813	0.89787
5	campestris	INT	885	135	335	97653	85685	9669
AMNH978	Raphicerus	B-G	3.0671	0.0034	0.7085	9057.9	0.2980	0.82606
56	campestris	INT	515	715	795	05827	72632	6884

85 0	Capilicelus	B-G	2.3549	0.0035	0.6088	7620.5	0.5104	0.93251
	campestris	INT	065	535	735	44291	33992	4101
AMNH464 I	Raphicerus	B-G	1.6399	0.0015	0.5546	8854.7	0.5261	0.78397
85 0	campestris	INT	16	155	98	72553	37007	9284
AMNH978 I	Raphicerus	B-G	2.4254	0.0024	0.8508	9712.0	0.5325	0.94235
15 0	campestris	INT	48	56	17	76998	67154	6462
AMNH974 I	Raphicerus	B-G	1.6428	0.0040	0.5417	6736.3	0.5134	1.09466
85 0	campestris	INT	19	71	86	89633	06566	3089
AMNH978 I	Raphicerus	B-G	2.7726	0.0027	0.4165	7098.0	0.4076	0.87427
26	campestris	INT	77	29	24	74985	17579	2389
AMNH874 I	Raphicerus	B-G	1.9322	0.0040	0.7416	6829.7	0.4879	0.97083
56	campestris	INT	285	115	6	69029	74367	8616
AMNH164 I	Raphicerus	B-G	2.9914	0.0025	1.2164	5292.4	0.4360	0.87208
85 0	campestris	INT	665	465	535	12487	57002	2124
AMNH978 I	Raphicerus	B-G	1.6972	0.0011	0.8669	9824.3	0.3193	0.67777
58	campestris	INT	945	02	665	33663	00296	702
NMNH485 I	Raphicerus	B-G	2.3654		0.6984	5103.1	0.5132	0.76131
446	sharpei	INT	8	0.0045	56	324	135	54
NMNH494 I	Raphicerus	B-G	1.8321		0.3645	3625.1	0.3965	0.81315
515	sharpei	INT	35	0.009	61	313	41351	4
NMNH487 I	Raphicerus	B-G	2.7321		1.0351	5873.3	0.6731	0.93213
94	sharpei	INT	35	0.0037	564	8	354	54
NMNH475 I	Raphicerus	B-G	1.5321			4862.9	0.5006	0.73135
16	sharpei	INT	5	0.0041	0.8315	62	4	4
NMNH478 I	Raphicerus	B-G	2.9315		0.5687	3587.1	0.5365	0.83213
52	sharpei	INT	4	0.0035	4651	1	45	54
NMNH497 I	Raphicerus	B-G	2.5231		0.4326	2951.1	0.6231	0.93543
21	sharpei	INT	534	0.0051	56548	35	35	53
NMNH497 I	Raphicerus	B-G	1.9684		0.7645	4951.3	0.4913	0.86546
51	sharpei	INT	52	0.0028	4	2135	54	8
NMNH479 I	Raphicerus	B-G	2.8321		0 7000			
(1	1 .		2.0521		0.7323	6/56.8	0.5732	0.98784
61	sharpei	INT	5	0.0031	0.7323	6756.8 4	0.5732 135	0.98784 35
NMNH487 I	sharpei Raphicerus	INT B-G	5 1.9321	0.0031	0.7323 154 1.0645	6756.8 4 4761.3	0.5732 135 0.3865	0.98784 35 0.76546
61 NMNH487 I 58	sharpei Raphicerus sharpei	INT B-G INT	5 1.9321 5	0.0031 0.0046	0.7323 154 1.0645 4	6756.8 4 4761.3 5	0.5732 135 0.3865 456	0.98784 35 0.76546 8
61 NMNH487 H 58 NMNH497 H	sharpei Raphicerus sharpei Raphicerus	INT B-G INT B-G	5 1.9321 5 2.5354	0.0031 0.0046	0.7323 154 1.0645 4 0.3984	6756.8 4 4761.3 5 2542.4	0.5732 135 0.3865 456 0.4232	0.98784 35 0.76546 8 0.86486
61 NMNH487 I 58 NMNH497 I 81	sharpei Caphicerus sharpei Caphicerus sharpei	INT B-G INT B-G INT	5 1.9321 5 2.5354 68	0.0031 0.0046 0.0037	0.7323 154 1.0645 4 0.3984 5	6756.8 4 4761.3 5 2542.4 8	0.5732 135 0.3865 456 0.4232 135	0.98784 35 0.76546 8 0.86486 7
61 NMNH487 H 58 NMNH497 H 81 NMNH485 H	sharpei Raphicerus sharpei Raphicerus sharpei Raphicerus	INT B-G INT B-G INT B-G	5 1.9321 5 2.5354 68 2.8984	0.0031 0.0046 0.0037	0.7323 154 1.0645 4 0.3984 5 0.5989	6756.8 4 4761.3 5 2542.4 8 5435.6	0.5732 135 0.3865 456 0.4232 135 0.5365	0.98784 35 0.76546 8 0.86486 7 1.02541
61 NMNH487 H 58 NMNH497 H 81 NMNH485 H 61	sharpei Raphicerus sharpei Raphicerus sharpei Raphicerus sharpei	INT B-G INT B-G INT B-G INT	5 1.9321 5 2.5354 68 2.8984 5	0.0031 0.0046 0.0037 0.0041	$\begin{array}{c} 0.7323 \\ 154 \\ 1.0645 \\ 4 \\ 0.3984 \\ 5 \\ 0.5989 \\ 9 \end{array}$	6756.8 4 4761.3 5 2542.4 8 5435.6 5	0.5732 135 0.3865 456 0.4232 135 0.5365 4165	$\begin{array}{c} 0.98784\\ 35\\ 0.76546\\ 8\\ 0.86486\\ 7\\ 1.02541\\ 5\end{array}$
61 NMNH487 I 58 NMNH497 I 81 NMNH485 I 61 NMNH478 I	sharpei Raphicerus sharpei Raphicerus sharpei Raphicerus sharpei Raphicerus	INT B-G INT B-G INT B-G INT B-G	5 1.9321 5 2.5354 68 2.8984 5 2.8321	0.0031 0.0046 0.0037 0.0041	0.7323 154 1.0645 4 0.3984 5 0.5989 9 0.2984	6756.8 4 4761.3 5 2542.4 8 5435.6 5 4526.3	$\begin{array}{c} 0.5732 \\ 135 \\ 0.3865 \\ 456 \\ 0.4232 \\ 135 \\ 0.5365 \\ 4165 \\ 0.6798 \end{array}$	0.98784350.7654680.8648671.025415
61 NMNH487 H 58 NMNH497 H 81 NMNH485 H 61 NMNH478 H 46	sharpei Raphicerus sharpei Raphicerus sharpei Raphicerus sharpei Raphicerus sharpei	INT B-G INT B-G INT B-G INT B-G INT	5 1.9321 5 2.5354 68 2.8984 5 2.8321 35	0.0031 0.0046 0.0037 0.0041 0.0029	0.7323 154 1.0645 4 0.3984 5 0.5989 9 0.2984 8	6756.8 4 4761.3 5 2542.4 8 5435.6 5 4526.3 1	0.5732 135 0.3865 456 0.4232 135 0.5365 4165 0.6798 45	$0.98784 \\ 35 \\ 0.76546 \\ 8 \\ 0.86486 \\ 7 \\ 1.02541 \\ 5 \\ 1.3545$
61 NMNH487 H 58 NMNH497 H 81 NMNH485 H 61 NMNH478 H 46 NMNH476 H	sharpei Raphicerus sharpei Raphicerus sharpei Raphicerus sharpei Raphicerus sharpei Raphicerus	INT B-G INT B-G INT B-G INT B-G	5 1.9321 5 2.5354 68 2.8984 5 2.8321 35 2.6354	0.0031 0.0046 0.0037 0.0041 0.0029	$\begin{array}{c} 0.7323 \\ 154 \\ 1.0645 \\ 4 \\ 0.3984 \\ 5 \\ 0.5989 \\ 9 \\ 0.2984 \\ 8 \\ 0.3987 \end{array}$	6756.8 4 4761.3 5 2542.4 8 5435.6 5 4526.3 1 3129.2	0.5732 135 0.3865 456 0.4232 135 0.5365 4165 0.6798 45 0.5935	$\begin{array}{r} 0.98784\\ 35\\ 0.76546\\ 8\\ 0.86486\\ 7\\ 1.02541\\ 5\\ 1.3545\\ 0.95435\end{array}$
61 NMNH487 I 58 NMNH497 I 81 NMNH485 I 61 NMNH478 I 46 NMNH476 I 465	sharpei Raphicerus sharpei Raphicerus sharpei Raphicerus sharpei Raphicerus sharpei Raphicerus sharpei	INT B-G INT B-G INT B-G INT B-G INT	$ \begin{array}{r} 5\\ 1.9321\\ 5\\ 2.5354\\ 68\\ 2.8984\\ 5\\ 2.8321\\ 35\\ 2.6354\\ 68\\ \end{array} $	0.0031 0.0046 0.0037 0.0041 0.0029 0.0037	$\begin{array}{c} 0.7323 \\ 154 \\ 1.0645 \\ 4 \\ 0.3984 \\ 5 \\ 0.5989 \\ 9 \\ 0.2984 \\ 8 \\ 0.3987 \\ 484 \end{array}$	6756.8 4 4761.3 5 2542.4 8 5435.6 5 4526.3 1 3129.2 3251	0.5732 135 0.3865 456 0.4232 135 0.5365 4165 0.6798 45 0.5935 45	$\begin{array}{c} 0.98784\\ 35\\ 0.76546\\ 8\\ 0.86486\\ 7\\ 1.02541\\ 5\\ 1.3545\\ 0.95435\\ 1\end{array}$
61 NMNH487 H 58 NMNH497 H 81 NMNH485 H 61 NMNH478 H 46 NMNH476 H 465 NMNH478 H	sharpei Raphicerus sharpei Raphicerus sharpei Raphicerus sharpei Raphicerus sharpei Raphicerus sharpei Raphicerus sharpei Raphicerus	INT B-G INT B-G INT B-G INT B-G INT B-G	$\begin{array}{c} 2.8321\\ 5\\ 1.9321\\ 5\\ 2.5354\\ 68\\ 2.8984\\ 5\\ 2.8321\\ 35\\ 2.6354\\ 68\\ 2.3321\end{array}$	0.0031 0.0046 0.0037 0.0041 0.0029 0.0037	$\begin{array}{c} 0.7323 \\ 154 \\ 1.0645 \\ 4 \\ 0.3984 \\ 5 \\ 0.5989 \\ 9 \\ 0.2984 \\ 8 \\ 0.3987 \\ 484 \\ 0.4265 \end{array}$	6756.8 4 4761.3 5 2542.4 8 5435.6 5 4526.3 1 3129.2 3251 6534.3	$\begin{array}{c} 0.5732 \\ 135 \\ 0.3865 \\ 456 \\ 0.4232 \\ 135 \\ 0.5365 \\ 4165 \\ 0.6798 \\ 45 \\ 0.5935 \\ 45 \\ 0.7231 \end{array}$	$\begin{array}{r} 0.98784\\ 35\\ 0.76546\\ 8\\ 0.86486\\ 7\\ 1.02541\\ 5\\ 1.3545\\ 0.95435\\ 1\\ 0.96878\end{array}$
61 NMNH487 H 58 NMNH497 H 81 NMNH485 H 61 NMNH478 H 46 NMNH476 H 465 NMNH478 H 16	sharpei Raphicerus sharpei Raphicerus sharpei Raphicerus sharpei Raphicerus sharpei Raphicerus sharpei Raphicerus sharpei Raphicerus sharpei	INT B-G INT B-G INT B-G INT B-G INT B-G INT	$5 \\ 1.9321 \\ 5 \\ 2.5354 \\ 68 \\ 2.8984 \\ 5 \\ 2.8321 \\ 35 \\ 2.6354 \\ 68 \\ 2.3321 \\ 5 \\ $	0.0031 0.0046 0.0037 0.0041 0.0029 0.0037 0.0045	$\begin{array}{c} 0.7323 \\ 154 \\ 1.0645 \\ 4 \\ 0.3984 \\ 5 \\ 0.5989 \\ 9 \\ 0.2984 \\ 8 \\ 0.3987 \\ 484 \\ 0.4265 \\ 486 \end{array}$	6756.8 4 4761.3 5 2542.4 8 5435.6 5 4526.3 1 3129.2 3251 6534.3 215	$\begin{array}{c} 0.5732 \\ 135 \\ 0.3865 \\ 456 \\ 0.4232 \\ 135 \\ 0.5365 \\ 4165 \\ 0.6798 \\ 45 \\ 0.5935 \\ 45 \\ 0.7231 \\ 5 \end{array}$	$\begin{array}{c} 0.98784\\ 35\\ 0.76546\\ 8\\ 0.86486\\ 7\\ 1.02541\\ 5\\ 1.3545\\ 0.95435\\ 1\\ 0.96878\\ 78\end{array}$
61 NMNH487 I 58 NMNH497 I 81 NMNH485 I 61 NMNH478 I 46 NMNH478 I 16 NMNH478 I	sharpei Raphicerus sharpei Raphicerus sharpei Raphicerus sharpei Raphicerus sharpei Raphicerus sharpei Raphicerus sharpei Raphicerus sharpei Raphicerus	INT B-G INT B-G INT B-G INT B-G INT B-G INT B-G	$\begin{array}{c} 2.5321\\ 5\\ 1.9321\\ 5\\ 2.5354\\ 68\\ 2.8984\\ 5\\ 2.8321\\ 35\\ 2.6354\\ 68\\ 2.3321\\ 5\\ 2.5321\end{array}$	0.0031 0.0046 0.0037 0.0041 0.0029 0.0037 0.0045	$\begin{array}{c} 0.7323 \\ 154 \\ 1.0645 \\ 4 \\ 0.3984 \\ 5 \\ 0.5989 \\ 9 \\ 0.2984 \\ 8 \\ 0.3987 \\ 484 \\ 0.4265 \\ 486 \\ 0.4698 \end{array}$	6756.8 4 4761.3 5 2542.4 8 5435.6 5 4526.3 1 3129.2 3251 6534.3 215 5975.1	$\begin{array}{c} 0.5732 \\ 135 \\ 0.3865 \\ 456 \\ 0.4232 \\ 135 \\ 0.5365 \\ 4165 \\ 0.6798 \\ 45 \\ 0.5935 \\ 45 \\ 0.7231 \\ 5 \\ 0.5831 \end{array}$	$\begin{array}{c} 0.98784\\ 35\\ 0.76546\\ 8\\ 0.86486\\ 7\\ 1.02541\\ 5\\ 1.3545\\ 0.95435\\ 1\\ 0.96878\\ 78\\ 0.76465\end{array}$

NMNH487	Raphicerus	B-G	2.0321		0.5984	6429.6	0.5613	0.86545
59	sharpei	INT	35	0.0039	54	915	54	1
NMNH487	Raphicerus	B-G	2.6684		0.4164	3824.3	0.4731	0.96875
52	sharpei	INT	54	0.004	8	215	54	4
NMNH485	Raphicerus	B-G	1.5654		0.2987	4624.3	0.5316	0.85145
76	sharpei	INT	56	0.0029	84	215	54	4
NMNH487	Raphicerus	B-G	1.9684		0.3216	6948.3	0.5931	0.91321
82	sharpei	INT	5	0.0051	548	5	543	5
NMNH487	Raphicerus	B-G			0.3744	6248.3	0.6531	
59	sharpei	INT	2.5315	0.0038	6546	215	54	1.13545
NMNH428	Raphicerus	B-G	2.7321		0.3984	2842.1	0.7131	
46	sharpei	INT	35	0.0041	84	354	54	1.26845
NMNH428	Raphicerus	B-G	2.7651		0.4987	5361.6	0.7335	
61	sharpei	INT	32	0.0045	84	61	15	1.0648
NMNH428	Raphicerus	B-G	1.8351		0.5164	2762.3	0.6965	
75	sharpei	INT	32	0.0052	84	2135	4153	0.95645
NMNH425	Raphicerus	B-G	1.9651		0.2987	5934.1	0.5931	0.86543
97	sharpei	INT	3	0.0045	84	324	54	21
NMNH428	Raphicerus	B-G	2.2654		0.3648	4816.4	0.6431	
71	sharpei	INT	65	0.0048	48	4	54	0.96845
AMNH568	Redunca		0 6854	0.0064	1 7616	41327	0 41 31	0.86155
76	arundinum	OG	9	5	54	82	54	4
AMNH465	Redunca	00	0 9841	ç	2 6845	1975 1	0 3568	0 63213
46	arundinum	OG	32153	0.0053	1	231	754	534
AMNH759	Redunca	00	1 2326	0100000	2 1845	4458 3	0 3846	0 73135
48	arundinum	OG	54	0 0077	1	215	51	4
AMNH795	Redunca	00	0 7135	0.0077	0 9984	4658.6	0 4032	0 86514
36	arundinum	OG	4	0.0061	1521	521	546	51
AMNH794	Redunca	00	1 1654	0.0001	1 8987	31571	0 3684	0 86415
28	arundinum	OG	68	0.0057	451	124	5	64
AMNH792	Redunca	00	0.8351	0.0007	1 9135	3497.2	0 3832	01
85	arundinum	OG	35	0.0055	45	34	15	0 79451
AMNH792	Redunca	00	0 9351	0.00000	1 4987	3024 3	0 4265	0.86878
81	arundinum	OG	5	0.0078	541	24	468	4
AMNH795	Redunca	00	0 6487	0.0070	1 3516	4521.2	0 3648	0 79845
83	arundinum	OG	8654	0.0062	54	15	678	0.79045 4
AMNH795	Redunca	00	0.6878	0.0002	1 1845	3751.1	0 3864	0.86515
80	arundinum	OG	0.0070 45	0.0056	216	2/31.1 24	51	0.00515 A
4 MNH703	Redunca	00	1 2154	0.0050	210	2975 3	0 3621	0 76158
/8	arundinum	OG	1.215 <del>4</del> 8	0 0040	2.1040 54	2775.5	0.3021 54	186
40 A MNH400	Redunca	00	1 0354	0.0047	2 0876	1578.6	0 3532	0 60708
8/	arundinum	06	1.0554 8	0.0064	2.0070	1378.0 A	1354	0.09798
04 AMNH720	Redunce	00	0 1 /221	0.0004	1 86/18	+ 1168 2	0 2025	4.5 0 50979
6/	arundinum	00	251	0.0057	1.0040 7	+100.2 7	0.2955	0. <i>37</i> 0/0 /
04 A MATU721	Rodunce	υŪ	0 8651	0.0037	1 6001	27015	1.5	4 0 60101
$\frac{1}{04}$	Neuuiica	00	201	0 005 1	1.0704	J∠04.J 24	0.5052	0.00404 5
94	arunainum	UG	321	0.0051	3	34	15	3

AMNH734	Redunca		0.9876		1.1684	2751.6	0.3668	
85	arundinum	OG	54	0.0062	74	924	45	0.79845
AMNH729	Redunca		0.8321		1.0431	3975.5	0.3764	0.79265
45	arundinum	OG	35	0.0067	54	424	541	4
AMNH487	Redunca		0.7687		0.9843	1851.9	0.4254	0.86545
46	arundinum	OG	4351	0.0053	32165	54	8	61
AMNH471	Redunca		1.0321		0.8315	4861.8	0.3365	0.69874
59	arundinum	OG	5	0.0047	4	424	48	854
AMNH154	Redunca		0.6687		0.9545	2834.1	0.3468	0.76515
82	arundinum	OG	451	0.0071	31	324	45	64
AMNH465	Redunca		0.7984		1.3984	3752.4	0.3406	0.71351
86	arundinum	OG	35	0.0068	564	8	5468	45
AMNH164	Redunca		1.1321			1985.4	0.4164	0.86512
82	arundinum	OG	45	0.0059	2.1845	67	84	1
AMNH164	Redunca		0.8984		1.9654	4297.5	0.3268	
85	arundinum	OG	5	0.0073	6	8	451	0.7645
AMNH164	Redunca		0.7984		1.9325	3685.2	0.3865	0.98465
87	arundinum	OG	561	0.0075	4	48	121	13
AMNH426	Redunca		1.2321		2.0631	4397.5	0.3632	0.69878
48	arundinum	OG	5	0.0059	54	46	15	4
AMNH426	Redunca		0.7846		0.9654	1862.4	0.4731	0.79845
87	arundinum	OG	51	0.0068	5	586	235	41
AMNH428	Redunca		0.6874		0.8156	2306.8	0.3235	0.83154
94	arundinum	OG	51	0.007	48	5	15	5
FMNH1279	Redunca		1.3857	0.0066	1.3420	1892.2	0.2999	0.47331
89	fulvorfula	OG	345	565	845	24607	00988	8097
FMNH1279	Redunca		1.1321	0.0086	1.2666	1811.5	0.3104	0.60822
90	fulvorfula	OG	645	55	665	71008	96452	9883
FMNH1279	Redunca		1.0742	0.0060	1.4182	1163.4	0.6151	0.57386
91	fulvorfula	OG	785	585	7	40413	12236	4017
FMNH1279	Redunca		1.6332	0.0081	1.2664	1676.5	0.2646	0.37847
92	fulvorfula	OG	485	57	905	49445	48469	7259
FMNH1279	Redunca		0.8723	0.0095	0.9183	1715.4	0.5307	0.55376
93	fulvorfula	OG	305	93	97	11128	1263	7309
FMNH1897	Redunca		1.1828	0.0081	0.8414	1593.3	0.4170	0.65730
3	fulvorfula	OG	795	465	585	66169	47865	7249
FMNH1958	Redunca		1.4216	0.0062	1.3444	1531.2	0.4365	0.64185
9	fulvorfula	OG	295	94	85	58576	36172	3988
FMNH1959	Redunca		1.2960	0.0075	1.1346	2722.2	0.5022	0.61964
1	fulvorfula	OG	185	99	1525	94797	10664	115
FMNH1959	Redunca		0.9517	0.0081	1.2673	2088.8	0.5723	0.62099
3	fulvorfula	OG	48	8	59	32416	66611	5272
FMNH1959	Redunca		1.4381	0.0073	0.7348	1925.9	0.4480	0.63563
5	6 1 6 1	00	105	265	17	10551	07220	052
	Tulvortula	ŪĠ	405	303	1/	48331	07239	932
FMNH1959	Redunca	ĐĠ	405 1.4789	363 0.0080	0.8096	48551 2425.3	07239	9 <i>32</i> 0.67496

FMNH1959	Redunca		1.0972	0.0100	1.3457	1207.1	0.4433	0.67362
7	fulvorfula	OG	3	55	115	81232	59833	3089
FMNH3291	Redunca		0.8918	0.0070	0.9456	1202.2	0.4654	0.62020
6	fulvorfula	OG	34	9	12	73098	01769	9662
FMNH3291	Redunca		1.1852	0.0089	1.2689	1609.5	0.4118	0.56006
7	fulvorfula	OG	05	995	665	43905	22931	1449
	Redunca		1.1819	0.0069	1.2496	1243.2	0.3615	6.25196
FMNH7041	fulvorfula	OG	3	21	46	4669	21754	7753
NMNH316	Redunca		1.5677	0.0069	0.7960	1056.6	0.3987	0.72939
48	fulvorfula	OG	385	98	875	75015	75464	0302
NMNH846	Redunca		1.0479	0.0075	0.8453	1638.6	0.3471	0.67580
48	fulvorfula	OG	84	5	51	78558	04179	4481
NMNH846	Redunca		1.2017	0.0065	0.8684	2496.7	0.5629	0.85771
28	fulvorfula	OG	425	04	44	89124	2832	8576
NMNH849	Redunca		0.8282	0.0071	0.7671	1885.5	0.3235	0.58857
75	fulvorfula	OG	355	695	68	23072	75729	7267
NMNH848	Redunca		1.4033	0.0079	1.2677	1036.7	0.4165	0.60266
78	fulvorfula	OG	455	73	3	67889	55965	146
NMNH316	Redunca		1.3857	0.0069	0.8076	2836.4	0.3034	0.59592
74	fulvorfula	OG	85	115	14	37782	66008	6782
NMNH319	Redunca		1.0225	0.0059	1.2915	1144.5	0.3604	0.62007
78	fulvorfula	OG	31	54	685	02632	25238	4565
NMNH318	Redunca		1.3501	0.0063	0.9337	1920.5	0.3790	0.51083
54	fulvorfula	OG	8	395	24	23414	36989	4646
NMNH849	Redunca		0.7501	0.0069	1.2722	1402.8	0.3083	0.54873
25	fulvorfula	OG	465	8	9	85014	02719	4277
FMNH1279	Sylvicapra	BRO	2.0500	0.0022	0.6367	18006.	0.4077	0.88439
98	grimmia	W	505	82	12	07053	9242	2643
FMNH1279	Sylvicapra	BRO	2.5138	0.0038	0.8079	15213.	0.5232	0.73444
99	grimmia	W	545	56	27	06576	02301	8797
FMNH1778	Sylvicapra	BRO	3.6115	0.0014	1.1414	9428.8	0.6079	1.26587
9	grimmia	W	34	17	53	83912	5212	0932
FMNH1779	Sylvicapra	BRO	3.3431	0.0020	0.5012	16031.	0.9296	0.90598
0	grimmia	W	93	855	72	60848	58022	036
FMNH2713	Sylvicapra	BRO	4.1213	0.0036	0.7333	12368.	0.4401	0.89243
9	grimmia	W	2	585	985	77115	1526	9224
FMNH2848	Sylvicapra	BRO	1.6169	0.0032	0.3831	15816.	0.6148	1.04895
9	grimmia	W	195	955	16	13497	27255	9348
FMNH3454	Sylvicapra	BRO	3.1845	0.0021	0.4669	7839.5	0.5182	0.89666
0	grimmia	W	135	385	335	25332	21199	592
	Sylvicapra	BRO	2.0451	0.0020	0.5088	8257.8	0.4671	0.85808
FMNH8166	grimmia	W	96	35	34	21808	07338	9012
FMNH8400	Sylvicapra	BRO	3.9328	0.0016	1.1090	10112.	0.5558	0.71347
6	grimmia	W	925	83	75	42845	15357	1115
FMNH8542	Sylvicapra	BRO	3.3433	0.0023	0.4066	11580.	0.9859	0.73946
6	grimmia	W	23	455	035	44965	27184	5318

FMNH4987	Sylvicapra	BRO	2.4864	0.0030	0.7665	8367.8	1.2134	0.83678
2	grimmia	W	13	255	47	3358	25675	1444
FMNH4987	Sylvicapra	BRO	2.6319	0.0025	0.3414	13818.	0.7424	0.96179
6	grimmia	W	635	54	735	71182	12794	1543
FMNH9785	Sylvicapra	BRO	3.6134	0.0027	1.2000	15307.	0.4444	0.75973
9	grimmia	W	305	57	78	56705	3176	3748
FMNH7485	Sylvicapra	BRO	3.2113	0.0033	0.5997	10017.	0.8525	0.75893
9	grimmia	W	225	92	495	67539	18137	3868
FMNH4965	Sylvicapra	BRO	3.9620	0.0011	0.5996	7720.2	0.7702	1.26328
8	grimmia	W	34	885	265	14279	34269	2816
FMNH9810	Sylvicapra	BRO	3.0043	0.0025	0.3427	10070.	0.4979	0.73333
5	grimmia	W	925	325	425	19987	60378	6386
NMNH498	Sylvicapra	BRO	4.1754	0.0037	0.5094	11599.	1.3028	0.91339
46	grimmia	W	515	455	19	34859	60292	6438
NMNH498	Sylvicapra	BRO	4.5078	0.0031	0.4347	14949.	1.3241	0.73163
59	grimmia	W	655	925	71	67774	44045	748
NMNH497	Sylvicapra	BRO	3.3255	0.0020	0.7422	10964.	0.8908	0.88181
85	grimmia	W	36	095	115	57491	5846	5256
NMNH487	Sylvicapra	BRO	3.2013	0.0036	0.4423	14007.	0.6034	0.95633
59	grimmia	W	345	64	685	36695	7707	6849
NMNH497	Sylvicapra	BRO	2.0662	0.0026	0.6019	5890.8	0.4742	0.88944
87	grimmia	W	97	265	035	15983	99246	2923
NMNH498	Sylvicapra	BRO	1.9087	0.0030	0.8670	6858.0	0.9325	0.68469
52	grimmia	W	175	775	14	74502	44122	1927
NMNH491	Sylvicapra	BRO	2.7052	0.0020	0.5672	13329.	0.4852	1.02221
58	grimmia	W	375	59	295	67597	949	6532
NMNH498	Sylvicapra	BRO	3.0628	0.0036	0.5110	11620.	0.4348	0.71769
75	grimmia	W	085	8	535	74159	19833	9377
NMNH497	Sylvicapra	BRO	3.2707	0.0027	0.5098	13092.	1.0993	1.03963
87	grimmia	W	02	45	75	61391	02644	4919
FMNH1050	Synceros		1.4033	0.0041	0.7499	3165.2	0.3288	0.52162
33	caffer	VG	16	95	655	604	57912	925
FMNH1280	Synceros		1.4870	0.0039	0.5012	1788.3	0.3050	0.89784
01	caffer	VG	07	725	75	90473	15028	0656
FMNH1408	Synceros		1.4103	0.0047	0.9196	2105.7	0.5752	0.96963
54	caffer	VG	94	755	17	26392	20711	2053
FMNH1408	Synceros		2.3902	0.0056	0.6696	3649.5	0.3949	0.64771
55	caffer	VG	61	04	11	00183	92258	7299
FMNH2068	Synceros		1.5133	0.0048	0.7430	3287.3	0.2609	0.46189
0	caffer	VG	35	295	59	1197	85439	9613
FMNH2114	Synceros			0.0038	0.5084	1416.9	0.3957	0.75509
8	caffer	VG	2.3356	66	88	83726	49006	7684
FMNH2114	Synceros		1.6032	0.0069	1.2618	4913.1	0.3892	0.88476
9	caffer	VG	64	145	569	85286	92343	3339
FMNH2115	Synceros		1.8847	0.0036	0.8132	2913.2	0.4963	0.74380
0	caffer	VG	965	535	859	3353	40376	2786

FMNH2432	Synceros		0.9365	0.0060	0.6727	3113.7	0.4547	0.80447
4	caffer	VG	31	905	153	99584	78326	4952
FMNH3454	Synceros		1.9005	0.0042	0.7814	2832.1	0.4045	0.81701
7	caffer	VG	645	065	935	50291	76633	1565
FMNH4406	Synceros		1.1074	0.0041	0.8498	4272.4	0.3476	0.55428
3	caffer	VG	27	55	89	94192	93256	8879
AMNH465	Synceros		1.9153	0.0055	0.5996	3918.7	0.4269	0.80718
79	caffer	VG	605	33	345	76672	65578	5008
AMNH984	Synceros		1.9741	0.0042	1.1708	4135.0	0.5593	0.67563
68	caffer	VG	86	785	6665	14902	15767	3593
AMNH487	Synceros		0.9647	0.0038	0.5120	1980.5	0.4565	0.53261
59	caffer	VG	795	945	1	13799	38607	6662
AMNH479	Synceros		1.0304	0.0050	0.6020	4087.3	0.5580	0.66621
54	caffer	VG	045	24	075	19285	97736	0604
AMNH497	Synceros		1.0937	0.0041	0.6053	1171.2	0.4577	0.72738
85	caffer	VG	845	995	505	42875	24385	2528
AMNH487	Synceros		1.4785	0.0052	0.6885	1236.9	0.5412	0.58579
52	caffer	VG	19	795	28	46571	47712	8048
AMNH987	Synceros		1.7228	0.0073	0.7458	2609.2	0.4180	0.78052
49	caffer	VG	025	075	6	80039	70454	7277
AMNH978	Synceros		1.7422	0.0066	0.6034	4762.2	0.4527	0.93341
45	caffer	VG	615	125	795	14682	02044	5434
AMNH965	Synceros		0.9579	0.0066	1.5616	1316.5	0.3587	0.70981
84	caffer	VG	28	48	6085	44993	85519	4067
AMNH948	Synceros		0.9354	0.0069	0.7421	2440.3	0.5001	0.61608
26	caffer	VG	37	635	955	7152	46852	987
AMNH948	Synceros		1.0004	0.0052	0.7439	1532.8	0.4629	0.73521
54	caffer	VG	495	065	26	48436	39499	5211
FMNH1280	Taurotragus	B-G	2.2958	0.0056	0.1507	7943.3	0.5592	1.23563
03	oryx	INT	27	39	67	66344	67859	8355
FMNH1820	Taurotragus	B-G	1.7721	0.0056	0.3419	5601.1	0.5350	0.80102
02	oryx	INT	51	1	435	40831	91514	8739
FMNH3433	Taurotragus	B-G	1.8356	0.0037	0.2101	6829.6	0.7095	1.23792
5	oryx	INT	33	095	14	29325	22794	262
FMNH3453	Taurotragus	B-G	1.9845	0.0023	0.4164	6529.6	0.4370	0.81372
2	oryx	INT	96	58	155	22807	46572	5227
FMNH3453	Taurotragus	B-G	1.7447	0.0016	0.5081	5435.7	0.5158	0.94430
4	oryx	INT	845	965	015	96314	88932	3662
FMNH3459	Taurotragus	B-G	1.9524	0.0044	0.7062	8380.1	0.4327	0.84183
9	oryx	INT	355	625	475	47767	80249	5736
FMNH4978	Taurotragus	B-G	1.8341	0.0046	1.0659	9080.3	0.4491	0.74910
5	oryx	INT	905	81	74	36939	33542	0014
FMNH3849	Taurotragus	B-G	2.1406	0.0035	0.9414	8745.8	0.3302	0.79342
5	oryx	INT	635	52	15	88058	49835	7681
FMNH4294	Taurotragus	B-G	2.5016	0.0033	0.9409	9373.2	0.3980	0.83152
8	oryx	INT	2	42	89	15265	47689	4505

FMNH4879	Taurotragus	B-G	2.1329	0.0050	0.8162	6785.0	0.3885	0.69158
5	oryx	INT	575	695	54	17311	93546	8538
FMNH4987	Taurotragus	B-G	2.3549	0.0038	0.6348	8774.3	0.4775	0.79075
9	oryx	INT	9	52	9955	59309	19898	1365
FMNH1948	Taurotragus	B-G	1.4277	0.0051	0.2090	5609.2	0.7221	1.15800
9	oryx	INT	655	38	855	65711	62699	6383
FMNH4985	Taurotragus	B-G	2.3558	0.0021	0.4167	6149.7	0.5515	0.96883
9	oryx	INT	305	81	615	10887	40178	9296
FMNH4785	Taurotragus	B-G	1.8054	0.0029	0.3415	5898.5	0.8378	1.13997
8	oryx	INT	665	46	48	25379	61314	1889
FMNH2064	Taurotragus	B-G	2.3097	0.0047	0.2698	7895.3	0.3360	0.76436
9	orvx	INT	97	9	62	06196	73405	2765
FMNH2066	Taurotragus	B-G	2.3323	0.0030	0.1511	7824.2	0.8002	1.35677
4	orvx	INT	705	17	36	02539	21572	8142
FMNH2075	Taurotragus	B-G	3.1454	0.0029	0.1499	5847.0	0.6061	1.21459
0	orvx	INT	775	07	215	25776	46123	0387
FMNH2075	Taurotragus	B-G	1.3409	0.0034	0.2677	6615.3	0.5043	0.67647
5	orvx	INT	085	045	895	54493	95178	9054
FMNH3511	Taurotragus	B-G	1.8280	0.0048	0.4346	5306.4	0.4457	0.97000
4	orvx	INT	8	69	025	22416	39001	2623
FMNH4084	Taurotragus	B-G	2.4745	0.0024	0.1504	7317.7	0.4019	0.72158
9	orvx	INT	7	885	945	23911	39213	5179
FMNH2087	Taurotragus	B-G	3.0203	0.0041	0.4362	7199.3	0.8845	1.23945
6	orvx	INT	835	135	125	55896	19879	1013
FMNH1343	Tragelaphus		1 8672	0 0029	0 4504	9816.5	0 4866	1 18038
27	angasi	GEN	9	495	89	25136	00825	025
FMNH3023	Tragelaphus		3.6479	0.0060	0.1536	10481.	0.3609	0.72677
8	angasi	GEN	61	365	675	39136	6696	6338
FMNH2615	Tragelaphus		1.7943	0.0056	0.4218	7764.8	0.4254	0.82396
8	angasi	GEN	635	125	005	61537	42373	4274
FMNH2648	Tragelaphus		1.3734		0.1505	11019.	0.4942	0.98566
5	angasi	GEN	075	0.0037	2	58504	83153	1954
FMNH3615	Tragelaphus		1.2142	0.0024	0.3503	8250.8	0.3999	0.65074
4	angasi	GEN	66	47	735	93464	70737	3117
FMNH6152	Tragelaphus	CLI	1 2632	0.0021	0 2501	10351	0 3228	0 64702
6	angasi	GEN	575	965	35	87431	16279	1623
FMNH4215	Tragelaphus	CLI	1.6801	0.0027	0.4819	10577.	0.3764	0.82144
9	angasi	GEN	36	495	905	79867	02196	9841
AMNH945	Tragelaphus	0	2 4937	0 0046	0 2094	8129 5	0 5418	0 96381
19	angasi	GEN	105	885	42	29326	13501	297
AMNH926	Tragelaphus	CLI	2 6315	0.0065	0 4439	99296	0 4914	1 11861
48	angasi	GEN	985	97	645	72413	08025	9272
AMNH824	Tragelaphus	0211	2.0237	0.0040	0.2011	8821 5	0.5479	0.88166
96	angasi	GEN	565	045	355	49956	73105	2083
AMNH495	Tragelanhus	5211	2 5521	0 0060	0 2673	89364	0 3938	0 83859
36	angasi	GEN	36	1	63	17317	5178	9087
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AMNH849	Tragelaphus		1.1393	0.0028	0.1900	10437.	0.4002	0.64752
52	angasi	GEN	01	74	69	83267	54153	0823
AMNH182	Tragelaphus		1.8842	0.0035	0.1509	11545.	0.3253	0.73202
94	angasi	GEN	28	29	23	62743	74345	6767
AMNH294	Tragelaphus		1.5056	0.0028	0.4507	12636.	0.3168	0.56346
85	angasi	GEN	69	3	545	83523	0385	964
AMNH198	Tragelaphus		1.3168	0.0028	0.1503	11283.	0.5819	0.75774
25	angasi	GEN	59	93	845	29722	39945	3849
AMNH497	Tragelaphus		1.7612	0.0035	0.2669	7116.0	0.5845	0.79951
85	angasi	GEN	065	555	005	62027	71536	3515
AMNH154	Tragelaphus		1.5561	0.0038	0.3505	11499.	0.4570	0.75465
76	angasi	GEN	135	055	765	7955	55124	4006
RMCA83-	Tragelaphus		3.2654		0.3854	9835.1	0.3598	0.82658
006M475	angasi	GEN	8	0.0042	7	548	745	47
RMCA83-	Tragelaphus		3.2165		0.4523	6987.2	0.5287	0.89658
006M496	angasi	GEN	4	0.0039	01	6484	45	44
RMCA83-	Tragelaphus		2.4687		0.2412	10658.	0.5841	0.78512
006M685	angasi	GEN	8	0.0045	54	23651	25	54
RMCA83-	Tragelaphus		2.8654		0.1985	9563.1	0.3845	0.62548
006M985	angasi	GEN	87	0.0051	236	548	4	4
RMCA83-	Tragelaphus		2.4877		0.2987	8659.2	0.4568	
006M698	angasi	GEN	6	0.0035	54852	657	4	0.78954
FMNH3510	Tragelaphus	BRO	3.2615	0.0010	0.7081	10659.	0.5657	0.95194
6	euryceros	W	91	4	855	91743	13982	4315
FMNH3510	Tragelaphus	BRO	3.7387	0.0013	0.4114	9022.8	0.4163	0.80093
7	euryceros	W	87	595	88	74742	94806	9684
FMNH3510	Tragelaphus	BRO	4.1842	0.0020	0.5679	12424.	0.6653	0.99197
8	euryceros	W	795	34	14	7385	73682	2042
FMNH9784	Tragelaphus	BRO	4.5045	0.0022	1.0437	15965.	0.5647	0.73714
6	euryceros	W	545	19	985	16457	01118	9167
FMNH2154	Tragelaphus	BRO	4.2614	0.0017	0.9474	15666.	0.5462	1.14476
6	euryceros	W	865	35	605	39715	09313	5017
FMNH9781	Tragelaphus	BRO	3.6741	0.0022	0.8541	13793.	0.6745	0.73875
6	euryceros	W	795	6	4164	0638	83967	5144
RMCA83-	Tragelaphus	BRO	5.5199	0.0017	0.3586	10407.	0.8330	1.17669
006M978	euryceros	W	54	805	96377	30745	25423	0199
RMCA83-	Tragelaphus	BRO	4.3838	0.0008	1.1107	13508.	0.6852	1.31546
006M548	euryceros	W	03	025	59	74607	45594	0231
RMCA-	Tragelaphus	BRO	3.1264	0.0014	0.9852	11622.	0.5132	0.81333
006M956	euryceros	W	865	04	5404	47498	37876	4031
RMCA83-	Tragelaphus	BRO	3.0509	0.0020	1.0412	15701.	0.6438	0.99611
006M876	euryceros	W	475	82	8255	45271	81079	1473
RMCA83-	Tragelaphus	BRO	4.5702	0.0014	0.7086	13696.	0.7147	1.17290
006-M548	euryceros	W	33	895	9635	31189	39554	5911
RMCA83-	Tragelaphus	BRO	2.6771	0.0022	0.8954	9358.3	0.5101	0.96569
006M872	euryceros	W	01	285	7876	27995	10416	692

RMCA83-	Tragelaphus	BRO	3.7078	0.0020	0.5207	15149.	0.6271	1.27496
006-M648	euryceros	W	3	015	965	95463	56059	9762
RMCA83-	Tragelaphus	BRO		0.0021	0.4324	15251.	0.7132	1.20990
006M318	euryceros	W	3.8597	83	38	58376	72012	1026
RMCA83-	Tragelaphus	BRO	2.5054	0.0019	1.0398	13962.	0.6544	0.97253
006M264	euryceros	W	07	86	4015	39552	64974	6148
RMCA83-	Tragelaphus	BRO	2.1190	0.0017	0.9813	10148.	0.4901	0.79246
006M856	euryceros	W	26	71	485	64298	97946	7921
RMCA83-	Tragelaphus	BRO	4.2776	0.0015	0.8540	11887.	0.7576	1.20725
006M361	euryceros	W	81	51	77495	17106	61973	7637
RMCA83-	Tragelaphus	BRO	3.1288	0.0021	1.2817	9561.1	0.5327	0.80262
006M879	euryceros	W	36	965	2665	05741	0095	4041
AMNH164	Tragelaphus	BRO	4.3521	0.0021	0.5632	13056.	0.6211	0.95123
85	euryceros	W	57742	0325	514	2658	25468	548
AMNH498	Tragelaphus	BRO	3.8594	0.0018	0.7985	11215.	0.7598	1.02154
81	euryceros	W	5	5945	17	2648	421	84
AMNH164	Tragelaphus	BRO	3.5248	0.0012	1.2054	9826.3	0.4568	0.85316
76	euryceros	W	5	548	7	2518	76	8465
AMNH648	Tragelaphus	BRO	4.8596	0.0021	0.6451	14625.	0.4876	0.79851
26	euryceros	W	42	548	36	32518	513	1324
	Tragelaphus	B-G	1.9599	0.0018	0.2230	6104.3	0.4227	0.62207
FMNH1417	imberbis	INT	655	555	76	43952	16757	1501
	Tragelaphus	B-G	1.5716	0.0026	0.2083	3586.9	0.4117	1.15938
FMNH1418	imberbis	INT	34	605	365	89323	59596	8114
FMNH2693	Tragelaphus	B-G	1.3108	0.0018	0.3416	7127.7	0.5652	1.07321
9	imberbis	INT	795	44	01	32995	05796	8597
FMNH2713	Tragelaphus	B-G	1.8968	0.0045	0.2664	7583.6	0.4544	0.69333
2	imberbis	INT	895	94	975	15289	94398	9727
	Tragelaphus	B-G	1.7233	0.0045	0.2666	6139.7	0.6616	0.95046
FMNH1420	imberbis	INT	095	085	58	1137	34046	7421
	Tragelaphus	B-G	1.6017	0.0023	0.2083	7072.6	0.4508	1.17816
FMNH1423	imberbis	INT	645	33	15	37796	29854	1069
	Tragelaphus	B-G	2.9909	0.0017	0.1499	9161.9	0.5823	0.95467
FMNH1427	imberbis	INT	87	085	495	84062	2256	7416
FMNH2716	Tragelaphus	B-G	1.9328	0.0052	0.2664	5225.7	0.5092	0.63282
4	imberbis	INT	16	45	575	96011	53979	8157
FMNH2791	Tragelaphus	B-G	1.4893	0.0046	0.2666	8327.9	0.5520	0.75338
8	imberbis	INT	025	71	76	43005	60228	8965
	Tragelaphus	B-G	1.6308	0.0041	0.3417	9427.9	0.4711	0.70172
FMNH1416	imberbis	INT	02	335	355	44431	31255	8552
	Tragelaphus	B-G	1.4385	0.0032	0.2668	9864.2	0.5898	0.91427
FMNH1412	imberbis	INT	965	375	94	6584	40478	9805
FMNH2794	Tragelaphus	B-G	1.9737	0.0054	0.2669	7702.1	0.5357	0.87340
8	imberbis	INT	23	415	05	13033	30954	9116
FMNH2765	Tragelaphus	B-G	2.0524	0.0056	0.2671	3814.1	0.4606	0.93660
8	imberbis	INT	825	895	49	98208	33746	2756

	Tragelaphus	B-G	1.3700	0.0042	0.2670	5589.4	0.5252	1.05823
FMNH1425	imberbis	INT	775	41	02	44536	27132	2644
	Tragelaphus	B-G	2.3332	0.0039	0.2089	10309.	0.4937	0.82404
FMNH1431	imberbis	INT	525	265	145	69833	51695	2974
FMNH1900	Tragelaphus	B-G	2.5295	0.0035	0.1518	8048.1	0.6152	1.14450
2	imberbis	INT	72	5685	095	3503	931	0106
FMNH2693	Tragelaphus	B-G	3.6666	0.0020	0.1501	6559.8	0.4236	1.33172
3	imberbis	INT	935	57	61	42741	02341	7358
FMNH2693	Tragelaphus	B-G	3.2950	0.0007	0.1499	10572.	0.3712	0.67185
5	imberbis	INT	585	095	255	15926	18114	4237
AMNH497	Tragelaphus	B-G	1.5940	0.0053	0.1513	4920.5	0.4319	0.80699
82	imberbis	INT	385	08	16	03133	23045	3514
AMNH197	Tragelaphus	B-G	2.0623	0.0035	0.2098	6835.7	0.4129	0.72774
48	imberbis	INT	735	535	035	86327	0433	9659
AMNH194	Tragelaphus	B-G	2.1051	0.0049	0.2674	8561.2	0.3556	0.47793
52	imberbis	INT	31	32	72	659	67022	0721
AMNH248	Tragelaphus	B-G	2.0632	0.0030	0.2083	4089.8	0.3112	1.20216
76	imberbis	INT	33	705	365	18011	35058	2144
AMNH298	Tragelaphus	B-G	2.2858	0.0039	0.2664	9114.2	0.4445	0.63526
75	imberbis	INT	31	235	52	5412	51081	6192
FMNH1866	Tragelaphus		1.8740	0.0060	1.3744	1458.9	0.3594	0.67305
77	spekii	VG	175	845	8	25904	13996	4437
FMNH2607	Tragelaphus		2.0821	0.0039	0.8200	1467.4	0.4493	1.06755
7	spekii	VG	85	545	685	31414	73186	4232
FMNH2607	Tragelaphus		1.4873	0.0064	1.4663	2374.6	0.4563	0.96225
8	spekii	VG	45	075	21	72746	0626	9406
FMNH2607	Tragelaphus		2.5787	0.0044	1.2450	4227.8	0.3063	0.92519
4	spekii	VG	055	61	365	68149	29461	5407
FMNH2068	Tragelaphus		1.0705	0.0060	1.0850	2750.6	0.3264	0.82980
1	spekii	VG	5	7125	835	08435	90572	9507
FMNH2068	Tragelaphus		2.1927	0.0057	1.1504	1634.0	0.4142	0.85519
6	spekii	VG	055	145	895	63668	71718	8645
FMNH2607	Tragelaphus		1.6483	0.0058	0.9668	4534.1	0.3289	0.84903
9	spekii	VG	09	695	915	98456	39439	9044
FMNH1866	Tragelaphus		2.0883	0.0067	1.3504	1875.6	0.3814	0.89352
72	spekii	VG	555	53	805	3843	26419	9452
RMCA83-	Tragelaphus		1.4115	0.0053	0.9503	2406.3	0.4896	0.80840
006M846	spekii	VG	39	15	025	7545	26336	6765
RMCA83-	Tragelaphus		1.3388	0.0063	1.2138	2234.6	0.5502	1.08192
006M538	spekii	VG	315	3	61	78133	81923	7703
RMCA83-	Tragelaphus		1.4011	0.0016	0.9209	3734.2	0.4643	1.22358
006M978	spekii	VG	005	235	55	14894	35062	3372
RMCA83-	Tragelaphus	_	2.8469	-	0.9851	1985.1	0.3487	0.75958
006M154	spekii	VG	5	0.0041	4	5487	9	7
RMCA83-	Tragelaphus	-	2.0215		1.0326	3528.2	0.4265	-
006M876	spekii	VG	4	0.0038	5	6548	8	0.86548
	1							

RMCA83-	Tragelaphus		1.8952		1.2326	4125.2	0.3654	0.76546
006M654	spekii	VG	6	0.0059	48	884	87	8
NMNH468	Tragelaphus		1.6587	0.0052	0.8515	1587.1	0.3987	0.96546
75	spekii	VG	84	2	4	684	87	8
NMNH548	Tragelaphus		2.3651		0.7165	1984.3	0.4012	0.92124
76	spekii	VG	8	0.0037	48	518	26	5
NMNH264	Tragelaphus		1.5978		1.0874	3287.5	0.3698	
86	spekii	VG	4	0.0042	9	48	74	0.85468
NMNH164	Tragelaphus		1.6255		0.9125	3164.5	0.4159	
85	spekii	VG	8	0.0033	4	88	87	1 07154
NMNH648	Tragelaphus		Ũ	0100000	0 7981	41204	0 4298	1.0,101
56	snekii	VG	1 9874	0 004	5	878	57	1 09784
NMNH164	Tragelaphus		2.0695	0.001	0.8321	3987 5	0 4875	0.98765
89	snekii	VG	8	0.0037	54	487	15	4
NMNH569	Tragelanhus		1 8547	0.0027	0 8688	2685 1	0 3548	
75	snekii	VG	8	0.0042	7	6498	7	0 85487
NMNH597	Tragelanhus		2 6264	0.0012	,	1859 5	0 3987	0.85154
53	snekii	VG	8	0.0029	1 1 5 4 7	468	87	6
FMNH1343	Tragelanhus	.0	3 2011	0.0029	0.4193	10955	0 4 1 8 4	0 64690
67	strensiceros	GEN	955	0.002	2	18074	33418	3696
07	Tragelanhus	<b>O</b> LIV	1 2492	0.0066	0 3415	8971 1	0 5235	1 20032
FMNH1414	strensiceros	GEN	69	32	19	73574	77837	8269
FMNH2719	Tragelanhus	ULI	2 6645	0.0037	0 2675	59524	0 4 2 0 7	0 75309
8	strensiceros	GEN	2.0043	42	0.2075	17071	01973	5769
5 FMNH3443	Tragelanhus	ULI	2 2909	0.0037	0 2666	10924	0 4924	1 03730
2	strensiceros	GEN	2.2909	0.0057	855	86206	51993	595
EMNH3//3	Tragelanhus	ULIV	2 6003	0.0050	0 2082	11592	0 4654	0.06055
3	strensiceros	GEN	655	855	0.2082 945	87352	28536	804
5 FMNH3443	Tragelaphus	ULIN	2 6088	0.0030	0 2088	0565 3	0 / 1 / 0	0 70358
/	strepsiceros	GEN	2.0988	305	25	76522	1807/	6205
4 EMNILI2442	Tragalaphus	ULIN	1 6064	0.0040	0 2 4 2 2	10660	0.6182	0295
5	stropsiooros	GEN	065	555	285	10000.	66522	0.00100
J NMNH580	Tragelaphus	ULIN	1 6500	0.0038	0 4337	6052.5	0.5040	0237
200	stropsiooros	GEN	1.0 <i>5</i> 0 <i>9</i>	10	0.4557	2186	0.3040	1207
200 NIMNIH580	Tragalaphus	UEN	1 5006	19	02	2180 7460.6	03078	0.86552
204	stransiaaras	CEN	1.3900	0.0049	0.4100	71162	12472	4507
204 A MNILI469	Tragalaphus	GEN	2 0 4 4 1	443	/3	/1103	134/3	439/
75	stransiaaras	CEN	2.9441	26	0.3420	9123.0 67740	5927	5071
/ J	Tragalanhua	GEN	44	30 0.0051	1	0//42	0.2821	0.00007
	ragelaphus	CEN	1.8270	0.0031	0.2079	/ 800.1	0.3621	0.99007
	strepsiceros	GEN	83	283	433	90922 (5(5 5	23804	9444
AMNH4/8	Tragelaphus	CEN	1.2440	0.0054	0.2070	0303.3 62079	0.3010	0.81/33
	strepsiceros	GEN	275	105	005	03078	40//9	3028
AIVIINEI49/	ragelaphus	CEN	2.1123	0.0048	0.2000	34/1.3	0.3099	0.04038
	strepsiceros	GEN	000	12	02	52295 10745	420/3	81U3
AIMINH498	i ragelaphus	OEM	2.3103	0.0043	0.2664	10/45.	0.494/	1.30341
/0	strepsiceros	GEN	4	200	915	8282	29861	2105

AMNH478	Tragelaphus		3.1203	0.0040	0.1508	6005.7	0.3688	0.83348
96	strepsiceros	GEN	28	645	395	21573	01915	1426
AMNH497	Tragelaphus		1.7352	0.0046	0.1499	9027.3	0.4745	0.83255
85	strepsiceros	GEN	8	67	695	65846	62255	8192
FMNH1297	Tragelaphus		3.0050	0.0045	0.2083	11281.	0.4108	0.96223
32	strepsiceros	GEN	065	33	125	43092	3212	6676
FMNH1648	Tragelaphus		2.0370	0.0049	0.1500	9718.7	0.4779	1.10987
7	strepsiceros	GEN	84	2	655	64167	93358	4629
FMNH6485	Tragelaphus		3.1655	0.0037	0.1500	11083.	0.4917	0.69218
6	strepsiceros	GEN	43	435	14	5564	94063	5345
FMNH6948	Tragelaphus		2.4314	0.0054	0.2666	7276.5	0.4734	0.87374
5	strepsiceros	GEN	055	265	885	62026	15628	0413
FMNH4687	Tragelaphus		3.1664	0.0048	0.1500	8459.2	0.4845	1.03075
6	strepsiceros	GEN	385	268	08	55567	14805	6851
FMNH5948	Tragelaphus		2.3875	0.0029	0.2789	7526.3	0.5126	0.86548
4	strepsiceros	GEN	1215	487	5154	1548	498	4545
FMNH4685	Tragelaphus		2.6687	0.0052	0.3464	9875.2	0.3987	0.69878
6	strepsiceros	GEN	8154	4878	8512	16547	4628	94545
RMCA83-	Tragelaphus		1.2656	0.0039	0.1988	10859.	0.4165	0.93157
006M497	strepsiceros	GEN	8132	5651	7412	65685	48621	81545
RMCA83-	Tragelaphus		3.0548	0.0031	0.2154	6985.1	0.3879	0.73216
006M648	strepsiceros	GEN	42365	56465	87515	65426	84612	5448

		Membe		EPLSA			HASF	HASF
Specimen	Taxon	r	ASFC	R	SMC	TFV	C9	C81
KNM-	Alcelaphini	Allia	2.4609	0.0019	0.2692	4930.2	0.5959	0.6962
ER17889	sp.	Bay	01	61	02	11117	76053	23764
KNM-	Tragelaphu	Allia	2.4299	0.0032	0.6794	12460.	0.7463	1.2190
ER17890	s sp.	Bay	795	715	645	65812	71761	6948
KNM-	Tragelaphu	Allia	3.1566	0.0036	0.8082	8132.4	0.5516	1.4814
ER42604	s sp.	Bay	165	83	775	85875	00035	5984
KNM-	Tragelaphu	Allia	3.4778	0.0026	0.6931	12887.	0.7103	0.8068
ER42655	s sp.	Bay	045	015	6183	30307	83818	25485
KNM-	Tragelaphu	Allia	3.9480	0.0020	0.7499	10128.	0.5780	1.1231
ER42663	s sp.	Bay	215	59	035	28305	67292	52467
KNM-	Tragelaphu	Allia	3.2459	0.0030	0.9086	12353.	0.5230	0.8120
ER42673	s sp.	Bay	59	07	825	39044	98438	73315
KNM-	Tragelaphu	Allia	4.1230	0.0037	0.8693	8304.0	0.6186	0.8828
ER42685	s sp.	Bay	465	085	35	26325	31097	82928
KNM-	Tragelaphu	Allia	2.6580	0.0023	0.7969	12625.	0.6267	0.6577
ER42686	s sp.	Bay	805	265	2327	34027	25316	11386
KNM-	Tragelaphu	Allia	4.7388	0.0020	0.6505	7968.0	0.6269	1.1976
ER42689	s sp.	Bay	63	71	555	76129	12731	78889
KNM-	Tragelaphu	Allia	3.1516	0.0017	0.8085	10256.	0.7306	0.7919
ER42694	s sp.	Bay	46	685	4	51326	13434	11924
KNM-	Tragelaphu	Allia	2.3995	0.0016	0.7667	9739.2	0.6053	1.0848
ER42823	s sp.	Bay	925	58	375	17806	09416	47547
KNM-	Tragelaphu	Allia	2.7211	0.0030	0.8671	13694.	0.6229	0.7004
ER42831	s sp.	Bay	19	24	925	89911	09262	07349
KNM-	Tragelaphu	Allia	3.7189	0.0028	0.9341	11755.	0.7072	1.2529
ER42953	s sp.	Bay	31	035	1511	99358	75427	34552
KNM-	Tragelaphu	Allia	2.6978	0.0017	0.6091	9026.7	0.8101	0.8565
ER42974	s sp.	Bay	245	515	99	94839	29851	26755
KNM-								
ER79465	Tragelaphu	Allia	4.3156	0.0019	0.7671	11520.	0.4998	1.1437
4	s sp.	Bay	48	955	655	70831	96807	1602
	Aepyceros	Denen	2.1324	0.0042	0.4816	5317.6	0.4813	
AL114-7	sp.	Dora	68	8	48	51465	54	0.8354
	Aepyceros	Denen	1.6984		0.5268	4921.5	0.3984	0.7651
AL 385-1	sp.	Dora	545	0.004	74	15	54	5
	Aepyceros	Denen	1.7984		0.4148	6108.6	0.4356	0.6926
AL794-1	sp.	Dora	65	0.0034	78	546	987	48
	Aepyceros	Denen	1.6986		0.5664	5425.4		0.7354
AL158-16	sp.	Dora	15	0.0037	87	31797	0.4182	37477
	Aepyceros	Kada	1.2878		0.2478	6264.3	0.3759	0.9845
AL450-1	sp.	Hadar	45	0.0041	13	548	31	1
	Aepyceros	Kada	1.4846		0.2684	5627.3	0.4122	0.8544
AL 463-1	sp.	Hadar	54	0.0037	5	15	64	1

Appendix II: Specimen numbers and raw data for fossil specimens

	Aepyceros	Kada	2.0656		0.3168	7345.9	0.4065	0.8642
AL870-1	sp.	Hadar	48	0.0049	45	8454	48	06
	Aepyceros	Kada	1.8654		0.2845	5972.3	0.3865	0.7548
AL939-2	sp.	Hadar	65	0.0037	156	5145	152	1
	Aepyceros	Kada	1.4813	0.0046	0.2665	5482.8	0.3312	0.6354
AL479-6	SD.	Hadar	945	11	435	7357	55662	37477
	1	Sidi						
	Aepyceros	Hakom	3.5261		0.2971	8185.3	0.5298	0.9451
AL50-35	sp.	a	468	0.0021	5	515	1	5
	-1	Sidi						
	Aepvceros	Hakom	3.9584		0.2976	9815.6	0.4965	0.8231
AL 222-2	sp.	a	51	0.0033	1	548	45	2
	-F.	Sidi						
	Aepvceros	Hakom	2.9884		0.3015	8752.4	0.5598	0.9968
AL1242-1	sp.	a	515	0.0024	48	8765	745	21
	Alcelaphini	Denen	2.1662	0.0049	0.4220	4564.1	0.4215	0.8601
AL135-17	sp	Dora	91	21	665	01844	86349	51003
	Alcelaphini	Denen	2.4890	0.0059	0.3121	5766.1	0.4736	0.9632
AL310-2	sp	Dora	78	33	89	16738	61604	11544
	Alcelaphini	Denen	1 6803	0 0047	0 4095	5600 7	0 6718	0 8985
AL55-35	sn	Dora	95	89	705	04758	14019	30909
AL116-	Alcelaphini	Denen	2 1546	0,7	0 4654	8731 1	0 5131	0.9513
146	sn	Dora	5	0.0031	86	254	35	24
1.0	Alcelaphini	Denen	2 8784	010001	0 4451	6264 5	0 5103	1 0326
AL181-4	sn	Dora	<u>-</u>	0 0048	3251	45	515	45
	Alcelaphini	Denen	2 1648	0.00.0	0 4298	8206.6	0 4987	0 9031
AL196-5	sn	Dora	4	0.0033	784	46	54	5
	Alcelaphini	Denen	2 3165	0.0022	0 3863	6214 6	0 4265	0 8754
AL162-2	sn	Dora	4	0.0042	3515	46	468	31
	Alcelaphini	Denen	2 5987	0.0012	0 3845	7628.6	0 4687	51
AL 398-18	sn	Dora	84	0.0021	44	465	1	0 9451
112370 10	Alcelaphini	Denen	1 7045	0.0059	0 5736	6989 1	0 6236	1 0498
AL310-3	sn	Dora	605	49	945	98931	36446	56342
1125105	Alcelanhini	Denen	2 3079	0.0052	0 4820	4873.8	0 5025	0.8347
AI 406-25	sn	Dora	355	21	165	1992	3542	83203
11L 100 25	Alcelanhini	Kada	1 6578	0 0077	0 3382	3148.5	0 3503	0 7724
AT 451-1a	sn	Hadar	86	0.0077	505	5864	76082	0.7724
712101 Iu	Alcelanhini	Kada	1 3136	0.0062	03131	5425.4	0 4 5 6 8	0.8721
AI /51-1b	sn	Hadar	1.5150	0.0002	6/15	31707	/07/1	8/375
112451-10	sp. Alcelanhini	Kada	1 18/6	))	0 2315	3513.6	0 308/	0.7616
AT 485 1	sn	Kaua Hadar	51	0.0075	0.2313 A	545	65	5/
AL403-1	sp. Alcelanhini	Kada	1 08/15	0.0075	0 3 3 1 5	28463	0.4015	0.8154
AT 406 26	Alcelapiiiii	Kaua Uodor	1.9045	0 0096	0.5515	2040.5	0.4013	0.0134
AL400-20	sp.	Sid:	1	0.0000	4	515	/	0
	Alcolonhini	Hakom	1 0761	0.0040	0 3100	5170 8	0 5256	0 7767
AI 225 6	Alcelapiiiii	11aKUIII	1.9201 625	215	0.3190 50	J1/7.0 70001	0.3330	56260
AL223-0	sp.	a	033	515	39	∠0094	00318	30209

		Sidi						
	Alcelaphini	Hakom	2.3095	0.0035	0.3984	6849.3	0.4987	0.8794
AL126-58	sp.	а	654	08	541	5145	64	5
		Sidi						
	Alcelaphini	Hakom	2.1687		0.4131	6347.6	0.5129	0.9026
AL325-12	sp.	а	684	0.0039	32	548	4	5468
		Sidi						
	Alcelaphini	Hakom	3.3081	0.0044	0.4420	7612.4	0.5777	0.8891
AL597-1	sp.	а	12	21	31	41309	72237	0245
	Antilopini	Denen	4.1597	0.0032	0.6254	10597.	0.5684	1.1648
AL773-3	sp.	Dora	8	84656	4879	36588	1354	79
	Antilopini	Kada	4.3165	0.0035	0.6876	9834.5	0.5988	1.3684
AL452-2	sp.	Hadar	468	49878	1654	48755	745	87
	Antilopini	Kada	3.8646	0.0041	0.5978	10587.	0.6865	1.0649
AL496-2	sp.	Hadar	54	65487	4654	31655	461	879
	Antilopini	Kada	3.1649	0.0029	0.6187	8268.6	0.6987	0.8616
AL452-15	sp.	Hadar	87	76547	6454	48785	8545	487
	Antilopini	Kada	2.9846	0.0026	0.7132	9597.2	0.7364	0.9813
AL444-27	sp.	Hadar	513	78465	6468	34687	878	4
	Antilopini	Kada	3.5876	0.0021	0.5684	9234.8	0.7598	1.3264
AL807-1	sp.	Hadar	4654	87465	6513	7845	7545	845
		Sidi						
	Antilopini	Hakom	5.1465	0.0037	0.6755	13685.	0.6059	1.0468
AL251-5	sp.	а	63	64651	126	66587	48	78
		Sidi						
	Antilopini	Hakom	4.4876	0.0029	0.5876	10268.	0.7164	1.2648
AL255-6	sp.	а	5465	25513	1315	74688	84654	7864
		Kada	0.9266	0.0069	1.0680	3564.1	0.4736	0.8632
AL535-2	Bovini sp.	Hadar	77	8	705	01844	61604	11544
		Kada	1.5092	0.0062	0.8346	4821.6	0.4187	0.7241
AL587-5	Bovini sp.	Hadar	765	78	135	67768	99156	77844
		Kada	0.8356	0.0066	1.0666	2989.1	0.4279	0.5778
AL716-3	Bovini sp.	Hadar	08	31	65	98931	61115	62749
		Kada	1.0368	0.0070	0.9425	1414.2	0.5247	0.8375
AL717-1	Bovini sp.	Hadar	687	66	405	3049	18888	93961
		Kada	0.5844	0.0080	1.2149	3980.9	0.3532	0.4502
AL1251-4	Bovini sp.	Hadar	045	995	32	41995	01023	86345
		Denen	3.0846	0.0061	0.5135	5684.6	0.3987	0.8984
AL161-4	Gazella sp.	Dora	545	84546	487	24651	1313	5645
		Denen	2.4987	0.0042	0.4978	4982.3	0.4164	0.7165
AL613-1	Gazella sp.	Dora	545	8465	46545	4451	86	69
		Kada	2.1978	0.0047	0.4898	5168.1	0.4325	0.8265
AL713-2	Gazella sp.	Hadar	465	63512	74654	14513	6846	4687
		Kada	2.3648	0.0055	0.5798	3385.0	0.4698	0.8946
AL713-3	Gazella sp.	Hadar	751	98746	76547	84515	75465	1313
AL132-7	Gazella sp.	Sidi	2.5987	0.0058	0.5987	6157.9	0.4982	0.6898

		Hakom	546	74545	5465	34654	648	7846
		а						
		Sidi						
		Hakom	1.9784	0.0039	0.6268	5984.2	0.4987	0.7684
AL666-24	Gazella sp.	а	513	98465	45645	95144	6545	54
	Hippotragi	Denen	1.5846	0.0063	0.3165	2197.7	0.4501	0.8135
AL1043-1	ni sp.	Dora	545	48784	4687	85485	78465	4654
	Hippotragi	Denen	2.0845	0.0059	0.4598	3984.6	0.3192	0.7946
AL427-2	ni sp.	Dora	4	78454	7545	84651	7854	513
	Hippotragi	Denen	1.9784	0.0055	0.4368	4129.1	0.5198	0.8874
AL133-37	ni sp.	Dora	513	31355	45642	98785	7846	13213
	Hippotragi	Kada	1.3468	0.0048	0.3846	2894.3	0.3878	0.8976
AL810-2	ni sp.	Hadar	7	9465	5465	1548	946	45351
	Hippotragi	Kada	1.2845	0.0049	0.4162	2358.8	0.4218	0.9451
AL610-1	ni sp.	Hadar	645	65132	6548	46545	76847	31
		Denen	1.2648	0.0051	0.7651	3826.3	0.4198	0.8551
AL157-4	Kobus sp.	Dora	78	3487	321	2154	654	3
		Denen	0.9846	0.0029	1.1876	5394.3	0.3987	0.7965
AL155-2	Kobus sp.	Dora	512	4645	546	45	654	31
		Denen	1.5987	0.0048	0.8654	4462.3	0.4029	0.8315
AL1334-1	Kobus sp.	Dora	451	9765	351	5456	8984	46
		Denen	2.1546	0.0036	1.2654	4862.3	0.4998	0.9265
AL155-7	Kobus sp.	Dora	578	9845	87	1545	78	48
		Denen	1.6984	0.0024	1.3468	3051.3	0.5165	0.9845
AL1227-1	Kobus sp.	Dora	513	87545	78	554	4867	6
		Denen	0.8465	0.0036	1.4987	3945.2	0.3532	0.7561
AL1230-1	Kobus sp.	Dora	487	98454	84	1524	1548	654
		Denen	1.3465	0.0031	0.9846	5192.3	0.3987	0.8613
AL1276-1	Kobus sp.	Dora	4687	68456	545	548	684	2354
		Denen	2.3984	0.0041	0.9984	4592.1	0.4517	0.9126
116-191	Kobus sp.	Dora	56	68465	654	3245	98	5487
		Denen	1.9846	0.0037	1.3968	2976.7	0.4538	0.8637
AL210-1	Kobus sp.	Dora	545	96465	7654	61354	7684	4654
		Kada	0.9876	0.0029	1.1354	1103.8	0.4324	0.7613
AL1352-2	Kobus sp.	Hadar	51321	76453	54	31545	965	3
	1	Sidi						
		Hakom	2.1654	0.0023	0.9984	3975.1	0.3265	0.9762
AL1311-1	Kobus sp.	а	687	6878	6545	3215	4687	664
	-	Sidi						
		Hakom	2.8165	0.0018	0.9464	4394.6	0.3264	0.6987
AL1218-1	Kobus sp.	а	4687	6487	51	487	687	654
	Neotragus	Kada	2.6987	0.0019	0.6246	10268.	0.6264	0.9884
AL454-9	sp.	Hadar	45641	8453	87	15465	867	35132
	1	Sidi						
	Neotragus	Hakom	5.2354	0.0018	0.6568	9763.8	0.7126	1.0336
AL200-5	sp.	а	687	65431	76546	45468	46786	548
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		Sidi						
	Neotragus	Hakom	3.9684	0.0017	0.7164	12648.	0.6935	0.8953
AL211-2	sp.	а	54	63214	87846	26487	1454	5132
		Sidi						
	Neotragus	Hakom	4.2654	0.0031	0.6865	8753.6	0.5698	0.9531
AL249-7	sp.	а	687	54687	43213	54658	75456	3548
	-	Kada	0.9987	0.0081	1.2648	986.35	0.5984	1.0264
AL515-1	Oryx sp.	Hadar	46514	65469	765	4587	651	867
	• •	Kada	1.3688	0.0059	1.0631	1168.6	0.6798	0.9754
AL447-3	Oryx sp.	Hadar	754	8454	254	548	46545	543
	• •	Kada	0.7984	0.0076	1.4987	1987.3	0.4879	0.9264
AL447-2	Oryx sp.	Hadar	51321	98452	656	2154	131	6867
	Pelorovis	Denen	1.3987	0.0053	0.6943	2168.3	0.4987	0.8135
AL167-2	sp.	Dora	654	6487	132	54548	654	4576
	Pelorovis	Denen	1.0654	0.0067	0.8654	2489.3	0.5135	0.8965
AL116-39	sp.	Dora	87	98465	87	5454	457	1321
	Pelorovis	Kada	1.3654	0.0061	0.7165	1972.3	0.3968	0.7132
AL709-3	sp.	Hadar	51	64876	4687	5454	4654	64568
	Pelorovis	Kada	1.2654	0.0059	0.7898	3381.6	0.4587	0.7945
AL713-1	sp.	Hadar	87	87651	7654	54687	65456	45315
	Reduncini	Denen	0.9846			2984.2	0.4987	0.8945
AL 116-6	sp.	Dora	51	0.0073	1.2654	654	8654	13
	Reduncini	Denen			1.0354	1026.3	0.3654	0.9764
AL134-15	sp.	Dora	1.1648	0.0056	86	51454	87	1351
	Reduncini	Denen	1.5987		0.8654	2873.1	0.4126	1.0354
AL156-1	sp.	Dora	84	0.0059	31	6584	84768	8
	Reduncini	Denen	0.8351		0.9765	3821.8	0.4632	0.7652
AL154-57	sp.	Dora	54	0.0058	4	64	648	513
	Reduncini	Denen	0.6874		1.3568	3384.6	0.3497	0.6998
AL118-13	sp.	Dora	51	0.0071	476	54	8465	46513
	Reduncini	Denen	1.0326			2493.3	0.2978	0.9894
AL118-20	sp.	Dora	9	0.0069	1.8645	545	45	51
	Reduncini	Denen	0.9715		1.4987	3935.6	0.4258	0.8645
AL524-6	sp.	Dora	46	0.0063	45	54444	79	13
	Reduncini	Denen	0.7598		1.6984	921.87	0.4978	0.7984
AL116-77	sp.	Dora	42	0.0041	5	36	131	654
	Reduncini	Denen	1.3874		0.8641	1138.6	0.5369	0.8321
AL158-76	sp.	Dora	6	0.0052	35	543	8751	6547
	Reduncini	Denen	1.3987		0.9735	2462.8	0.6198	0.8364
AL341-10	sp.	Dora	4	0.0069	1	7651	784	84
	Reduncini	Denen	1.3654		0.7651	2069.6	0.5978	0.7985
AL279-8	sp.	Dora	8	0.0046	3	846	46	43
	Reduncini	Denen	0.9984		1.1654	1576.6	0.4987	0.7126
AL153-5	sp.	Dora	65	0.0061	68	87461	846	48
	Reduncini	Denen	0.8144		1.2687	3358.6	0.3654	0.6976
AL181-6	sp.	Dora	87	0.0048	87	5548	687	64
	1							

	Reduncini	Denen	0.8687		1.3684	2692.6	0.3884	0.7065
AL169-7	sp.	Dora	4315	0.0064	654	54	651	846
	Tragelaphu	Denen	1.2649	0.0026	0.7676	10652.	0.5038	0.7534
AL116-94	s sp.	Dora	74	815	07	85091	94066	14207
	Tragelaphu	Denen	1.9451	0.0051	0.8097	5566.2	0.4315	0.6838
AL154-59	s sp.	Dora	175	12	13	5568	48011	86041
	Tragelaphu	Denen	2.3794	0.0061	0.7499	7720.3	0.5161	0.5829
AL158-17	s sp.	Dora	405	095	215	32497	32501	50261
	Tragelaphu	Denen	1.4811	0.0042	0.9698	6551.1	0.4749	0.5570
AL158-5	s sp.	Dora	18	965	355	27628	34198	02915
	Tragelaphu	Denen	2.1764	0.0048	1.0382	6527.1	0.3073	0.6754
AL158-6	s sp.	Dora	825	745	335	6338	56036	37447
	Tragelaphu	Denen	1.6268	0.0055	0.8849	5469.2	0.6266	0.7530
AL161-30	s sp.	Dora	47	965	82	14604	61264	25096
	Tragelaphu	Denen	2.6448	0.0075	1.1126	7907.7	0.5822	0.7484
AL167-28	s sp.	Dora	47	995	72	75471	56758	80789
	Tragelaphu	Denen	1.9828	0.0039	0.8338	4722.0	0.4897	0.8577
AL185-12	s sp.	Dora	73	065	79	3982	24215	78914
	Tragelaphu	Denen	2.5501	0.0054	0.6997	7059.5	0.4673	0.7057
AL567-1	s sp.	Dora	945	8	695	92552	9309	53969
	Tragelaphu	Denen	1.4850	0.0044	0.8767	6001.2	0.3749	0.7348
AL86-1	s sp.	Dora Sidi	805	805	035	12512	64349	69539
	Tragelaphu	Hakom	3.2201	0.0033	0.7409	8062.7	0.4973	0.8802
AL146-2	s sp.	a Sidi	705	215	33	3506	26206	36322
	Tragelaphu	Hakom	3.0762	0.0029	0.5638	7126.1	0.5840	0.9961
AL222-12	s sp.	a Sidi	13	095	425	01481	66825	07172
	Tragelaphu	Hakom	2.8587	0.0031	0.7079	11614.	0.4265	0.8383
AL25-17	s sp.	a Sidi	06	485	155	78684	86679	58138
	Tragelaphu	Hakom	3.1118	0.0027	0.6448	7529.9	0.5440	0.7070
AL539-12	s sp.	a Sidi	525	57	725	83075	11206	0521
	Tragelaphu	Hakom		0.0036	0.6195	10487.	0.6188	0.8553
AL885-1	s sp.	а	2.6043	85	46	15134	77217	05241
	Ugandex	Denen	2.3468	0.0041	0.4987	5134.9	0.4976	0.9163
AL133-12	sp.	Dora	67864	64879	86465	84651	5451	54
	Ugandex	Denen	2.1654	0.0055	0.5135	4031.6	0.5543	0.8264
AL169-3	sp.	Dora	87	98755	4876	4513	265	6548
	Ugandex	Denen	1.8646	0.0046	0.5597	5752.7	0.5132	0.8035
AL185-9	sp.	Dora	513	13548	65465	1513	648	4567
	Ugandex	Denen	2.0687	0.0037	0.6186	4983.3	0.4876	0.8940
AL185-30	sp.	Dora	64	65451	5456	5158	12531	54564
AL158-	Ugandex	Denen	1.6874	0.0049	0.4364	4267.6	0.4626	0.9034

100	sp.	Dora	513	75451	6876	87646	4687	6876
	Ugandex	Denen	2.6845	0.0049	0.5284	3357.9	0.5361	0.8164
AL362-9	sp.	Dora	13	33154	5645	6626	48	876
	Ugandex	Denen	1.9846	0.0055	0.4598	5384.9	0.5846	0.7596
AL907-2	sp.	Dora	513	98765	45465	76513	1526	54
	Ugandex	Denen	1.7946	0.0056	0.5026	4137.0	0.4987	0.9164
AL279-13	sp.	Dora	5132	84654	4688	2154	6545	68764
	Ugandex	Denen	2.7975	0.0042	0.4168	3982.3	0.6310	0.9120
AL1305-3	sp.	Dora Sidi	121	68764	79846	13546	5154	46846
	Ugandex	Hakom	2.9845	0.0031	0.4265	5826.8	0.5749	0.9305
AL265-1	sp.	a Sidi	6435	64654	46879	4513	751	8464
	Ugandex	Hakom	3.0875	0.0034	0.4897	5034.6	0.6132	0.7920
AL327-24	sp.	a Sidi	45	16469	86545	54651	6448	3664
	Ugandex	Hakom	3.2982	0.0037	0.4364	6138.6	0.5031	0.8579
AL224-15	sp.	a Sidi	5454	64513	65413	84513	544	4654
	Ugandex	Hakom	2.9897	0.0043	0.5136	4873.3	0.5842	0.8264
AL332-5	sp.	a Sidi	86456	16548	4687	21655	6254	878
	Ugandex	Hakom	2.4987	0.0029	0.4698	5394.2	0.5684	0.8546
AL237-12	sp.	a Sidi	68465	78646	76463	64874	5166	6846
	Ugandex	Hakom	2.7751	0.0034	0.4316	5762.9	0.5987	0.8126
AL886-1	sp.	а	65	87865	48798	6513	6565	4876
KNM-	Aepyceros	Kanapo	2.3278	0.0030	0.4502	8649.5	0.6444	1.0552
KP106	sp.	i	519	38	065	57176	22804	87541
KNM-	Alcelaphini	Kanapo	1.7861	0.0055	0.7505	3166.8	0.3847	0.6151
KP31733	sp.	i	75	24	735	61599	43197	45643
KNMKP7	Alcelaphini	Kanapo	1.3084	0.0034	0.8090	5740.8	0.3022	0.6433
3C	sp.	i	54	55	34	04173	37359	15361
KNMKP4	Alcelaphini	Kanapo	1.4195	0.0040	1.2665	6615.5	0.5032	0.6204
684	sp.	i	96	08	29	23163	10614	0796
KNMKP1	Alcelaphini	Kanapo	1.2916	0.0032	0.5664	4496.5	0.4381	0.7027
648	sp.	i	365	485	755	44482	22689	83721
KNMK01	Alcelaphini	Kanapo	1.7732	0.0037	0.7667	5969.2	0.4457	0.8182
645	sp.	1	445	045	09	88811	28588	01827
KNM-	Hippotragu	Kanapo	2.8316	0.0065	0.2675	3106.9	0.3149	0.7935
KP29274	s sp.	1	815	02	445	21376	65682	8124
KNM-	Hıppotragu	Kanapo	1.2866	0.0042	0.3890	2165.2	0.3146	0.7169
KP32526	s sp.	1	645	465	635	03595	53099	6124
KNM-	Madoqua	Kanapo	5.3047	0.0045	1.0350	5956.2	0.4998	0.8/83
KP103	sp.	1	95	4/5	8	63898	65145	/1543
KNM-	Madoqua	Kanapo	4.6155	0.0044	0.8510	16220.	0.5034	0.7068

KP30206	sp.	i	985	125	23	87274	88497	37529
KNM-	Madoqua	Kanapo	2.8354	0.0032	0.9170	17759.	0.6234	0.9102
KP30207	sp.	i	315	94	895	14335	76176	87001
KNM-	Madoqua	Kanapo	2.6203	0.0027	0.7672	17580.	0.4645	0.6783
KP30416	sp.	i	535	7	72	3078	67433	07352
KNM-	Madoqua	Kanapo	4.0728	0.0035	159.32	6775.2	0.8660	1.7627
KP30427	sp.	i	415	11	48995	54756	9766	01094
KNM-	Madoqua	Kanapo	3.9541	0.0037	0.8370	17634.	0.3977	0.7306
KP30537	sp.	i	955	0105	41	00437	70092	76057
KNM-	Madoqua	Kanapo	2.3060	0.0029	1.1551	14040.	0.5076	0.7849
KP36832	sp.	i	34	55	315	57707	72705	99177
KNM-	Madoqua	Kanapo	1.8347	0.0024	0.8900	14903.	0.3099	0.5414
KP36835	sp.	i	345	025	285	81333	55302	55649
KNM-	Madoqua	Kanapo	4.7322	0.0034	0.9501	19250.	0.4481	0.6092
KP36840	sp.	i	09	915	895	72913	27289	30377
KNM-	Raphicerus	Kanapo	3.6907	0.0043	0.2112	4931.5	0.6612	0.7582
KP30273	sp.	i	08	755	86	60042	35727	43616
KNM-	Raphicerus	Kanapo	2.6917	0.0021	0.3645	5799.5	0.7477	1.2119
KP30443	sp.	i	205	49	01	82372	52354	92446
KNM-	Raphicerus	Kanapo	4.4691	0.0016	0.5239	72203.	0.4803	1.1978
KP93	sp.	i	82	29	4	21969	04244	19262
KNM-	Simatheriu	Kanapo	1.9862	0.0063	0.5014	1485.6	0.5266	1.1371
KP29265	m sp.	i	31	09	75	55024	15502	84905
KNM-	Simatheriu	Kanapo	1.4262	0.0077	0.7753	2137.1	0.6757	0.8727
KP96	m sp.	i	38	99	525	72914	36075	62666
KNM-	Simatheriu	Kanapo	0.7382	0.0064	0.4501	953.60	0.3707	0.9517
KP8746	m sp.	i	69	205	68	69815	02738	15851
KNM-	Tragelaphu	Kanapo	3.5290	0.0030	0.1506	13242.	0.5148	1.4090
ER30395	s sp.	i	08	675	38	35152	41696	75416
KNM-	Tragelaphu	Kanapo	4.4025	0.0020	117.57	13268.	0.9102	1.1044
KP109	s sp.	i	735	38	71395	23962	06601	15547
KNM-	Tragelaphu	Kanapo	3.8073	0.0032	0.1500	13705.	0.4048	0.7666
KP29273	s sp.	i	665	21	715	49366	78526	36045
KNM-	Tragelaphu	Kanapo	4.9993	0.0012	0.1503	16610.	0.2533	0.4532
KP30421	s sp.	i	875	82	82	678	27419	91166
KNM-	Tragelaphu	Kanapo	12.218	0.0014	0.1500	17649.	0.1517	0.3067
KP32545	s sp.	i	1905	855	04	16246	29583	31334
KNM-	Tragelaphu	Kanapo	3.6626	0.0018	0.1502	13897.	0.7019	1.0177
KP32570	s sp.	i	24	865	03	55428	99726	28737
KNM-	Tragelaphu	Kanapo	4.9199	0.0012	0.1499	8655.8	0.4012	0.6744
KP32573	s sp.	i	885	23	06	03027	0644	46812
KNM-	Tragelaphu	Kanapo	4.0011	0.0036	0.1502	15139.	0.4657	0.9580
KP32574	s sp.	i	275	135	21	10903	40501	06844
KNM-	Tragelaphu	Kanapo	2.1932	0.0043	0.1499	9706.8	0.2651	0.4294
KP32829	s sp.	i	755	46	835	04205	66705	9546
KNM-	Tragelaphu	Kanapo	2.1045	0.0008	0.1500	10711.	0.3379	0.7770

KP32881	s sp.	i	91	1	63	22821	55359	8943
KNM-	Tragelaphu	Kanapo	3.3390	0.0018	1.3387	11329.	0.3266	0.5842
KP67	s sp.	i	45	095	745	82757	99747	53805
KNM-	Tragelaphu	Kanapo	5.2907	0.0016	0.1503	14582.	0.5323	0.7222
KP76	s sp.	i	56	21	42	67458	8821	30472
LAET25-	Alcelaphini	Laeotil	1.6468	0.0054	1.1499	6927.4	0.3716	0.7489
2246	sp.	Beds	75	65	835	02754	66098	08174
LAET75-	Alcelaphini	Laeotil	2.5946	0.0063	0.7499	7011.0	0.3363	0.6407
793	sp.	Beds	72	89	705	54853	98442	31496
LAET76-	Alcelaphini	Laeotil	2.7913	0.0055	0.9820	9492.3	0.4020	0.9408
7E-23	sp.	Beds	945	415	3855	02415	29077	61103
LAET76-	Alcelaphini	Laeotil	2.4390	0.0048		5889.1	0.5733	0.9320
7E-25	sp.	Beds	135	195	1.2499	56774	97374	31295
LAET75-	Alcelaphini	Laeotil	2.4446	0.0039	0.8350	6908.9	0.4103	1.0217
18-592	sp.	Beds	265	09	3335	51455	34399	0111
Laet75-	Alcelaphini	Laeotil	1.1672	0.0048	1.2625	4816.5	0.6013	0.8868
248	sp.	Beds	375	015	41	7711	26088	34575
LAET75-	Alcelaphini	Laeotil	1.5496	0.0059	1.1811	8044.7	0.3154	0.5276
2900	sp.	Beds	95	475	24	33116	41584	07726
LAET75-	Alcelaphini	Laeotil	1.9478	0.0020	0.8149	3423.1	0.5291	1.6222
4624	sp.	Beds	855	735	898	95817	91673	73819
	Alcelaphini	Laeotil	1.1672	0.0048	1.0665	6036.1	0.6013	0.8868
Laet75-56	sp.	Beds	375	015	41	86299	26088	34575
LAET76-	Alcelaphini	Laeotil	2.8632	0.0038	1.1250	6194.5	0.3205	0.5912
18-354	sp.	Beds	595	83	646	24061	14019	83469
LAET337	Antidorcas	Laeotil	2.6368	0.0017	0.5301	8598.8	0.5263	0.9331
3-00	sp.	Beds	785	59	98	97841	59079	42765
LAET75-	Antidorcas	Laeotil	4.8409	0.0013	0.3381	10484.	0.4030	1.0023
18-596	SD.	Beds	215	355	77	95727	14162	73112
LAET75-	Antidorcas	Laeotil	5.2489	0.0008	0.9817	6976.5	0.5885	1.0150
910	sp.	Beds	25	215	665	50413	73583	51344
, <b>-</b> .	-F.	Laeotil	3.2782	0.0044	0.5247	3560.1	0.4600	1.2764
EP816-03	Bovini sp.	Beds	485	19	35	18978	13949	37001
LAET75-	- • · · · · · · · · · · · · · · · · · ·	Laeotil	3.9603	0.0017	0.7101	4816.5	0.6851	1.3216
117	Bovini sp	Beds	285	595	17	7711	20608	9176
LAET75-	20 mil spi	Laeotil	2 1475	0 0040	0 6667	2235.4	0 3231	0 6973
1890	Bovini sp	Beds	715	16	885	72122	13433	14264
Laet75-	Bovini sp.	Laeotil	2 2026	0 0007	0 8304	4080 3	0 8043	1 3197
2.77	Bovini sp	Beds	64	67	01	59523	75311	22659
LAET75-	Bovini sp.	Laeotil	3 7335	0.0058	0 5803	6801.9	0 4199	0.8256
3052	Bovini sp	Beds	465	475	07	18438	43834	48195
LAET78-	Dovini sp.	Laeotil	1 2125	0 0040	0 6611	3431.6	0 7776	1 3692
4760	Bovini sn	Beds	56	44	025	00407	67412	01784
LAET78-	Dovini sp.	Laeotil	2 6893	0 0041	1 0045	5177.0	0 4246	0 7533
4761	Bovini sn	Beds	725	845	3	82571	99179	94466
I AFT78	Bovini sp.	Lapotil	1 7500	0.0016	0 7077	4631 0	0 /612	1 20/1
LAL1/0-	Dovini sp.	Lacoth	1./307	0.0010	0.1941	TUJ1.9	0.4012	1.4041

4763		Beds	43	56	17	42162	58014	53749
LAET78-		Laeotil	2.1367	0.0040	0.8401	4572.6	0.5051	1.2807
4826	Bovini sp.	Beds	71	14	67	26719	06023	64192
LAET78-	1	Laeotil	2.3222	0.0016	0.4988	5001.6	0.3281	0.6696
5176	Bovini sp.	Beds	155	065	25	78567	82815	33363
	1	Laeotil	3.3925	0.0037	0.5226	4281.4	0.3061	0.5766
EP497-00	Gazella sp.	Beds	505	45	15	12861	36192	77341
LAET75-	1	Laeotil	2.4512	0.0016	0.6342	2437.2	0.3546	0.6881
2741	Gazella sp.	Beds	38	67	495	60456	90069	10861
LAET78-	-	Laeotil	1.7175	0.0049	0.4867	4294.4	0.3820	0.8359
5198	Gazella sp.	Beds	59	45	425	27981	08146	44301
LAET75-	-	Laeotil	1.4566	0.0021	0.5505	5856.8	0.5366	1.6755
3520	Gazella sp.	Beds	99	715	52	76502	75699	03866
EP1268-	Hippotragu	Laeotil	1.0906	0.0042	0.3514	4800.9	0.6690	0.8478
00	s sp.	Beds	02	12	84	28882	8798	9888
LAET75-	Hippotragu	Laeotil	1.3433	0.0057	0.2703	8941.3	0.5894	0.7231
2949	s sp. (?)	Beds	605	055	685	80548	92669	02041
LAET75-	Madoqua	Laeotil	3.9077	0.0010	0.8504	12160.	0.4216	0.9867
5287	sp.	Beds	655	13	66	99281	27386	93913
LAET74-	Madoqua	Laeotil	2.7674	0.0031	0.9126	14011.	0.9084	1.9290
1982	sp. (?)	Beds	3	52	46	05485	3698	41254
LAET75-	Madoqua	Laeotil	4.8452	0.0016	0.6514	15858.	0.6153	1.2026
479	sp. (?)	Beds	99	3	545	15926	92967	52864
LAET75-	Parmulariu	Laeotil	1.4707	0.0047	1.1507	2663.3	0.3544	0.6271
1417	s sp.	Beds	25	605	61	46649	69646	82659
LAET75-	Parmulariu	Laeotil	2.1141	0.0065	0.9821	5911.4	0.4326	0.7726
2829	s sp.	Beds	82	68	495	99064	45734	17773
LAET75-	Parmulariu	Laeotil	0.7785	0.0051	1.6750	2474.0	0.6420	1.0697
2855	s sp.	Beds	805	665	559	10561	18139	79161
LAET75-	Parmulariu	Laeotil	0.9544	0.0044	1.4996	4466.1	0.1991	0.4691
2937	s sp.	Beds	3	895	1	92982	72321	62384
LAET75-	Parmulariu	Laeotil	1.2638	0.0038		2365.2	0.3977	0.8077
3376	s sp.	Beds	285	54	0.8506	22808	7613	786
LAET75-	Parmulariu	Laeotil	0.8201	0.0043	1.0800	1187.6	0.4784	1.0139
3458	s sp.	Beds	11	825	735	7524	90095	9669
LAET75-	Parmulariu	Laeotil	1.5567	0.0051	1.3600	5820.3	0.3345	0.9416
356	s sp.	Beds	62	06	565	81607	27639	70152
LAET75-	Parmulariu	Laeotil	2.0223	0.0070	0.9706	5377.3	0.4990	0.7606
357	s sp.	Beds	325	8	03	18689	19774	38786
LAET75-	Parmulariu	Laeotil	1.3235	0.0065	0.8950	3096.1	0.3017	0.8197
832	s sp.	Beds	31	265	005	70645	11341	79929
LAET75-	Parmulariu	Laeotil	1.1258	0.0077		4159.4	0.3234	0.5314
1669	s sp.	Beds	125	64	1.1499	60592	30523	03999
LAET75-	Simatheriu	Laeotil	3.0665	0.0030	0.4503	8932.1	0.4683	0.7690
616	m sp.	Beds	905	15	215	2449	85619	1904
LAET78-	Simatheriu	Laeotil	2.4788	0.0061	0.5208	6187.0	0.3996	0.6994

2437	m sp. (?)	Beds	04	08	347	89877	01059	23408
LAET78-	Simatheriu	Laeotil	2.2965	0.0022	0.6499	7784.0	0.4643	1.2019
4623	m sp. (?)	Beds	425	12	635	78839	80863	85378
LAET75-	Tragelaphu	Laeotil	4.0167	0.0019	0.7502	11169.	0.6042	0.8212
1300	s sp.	Beds	25	265	425	77475	63033	55208
LAET76-	Tragelaphu	Laeotil	3.1679	0.0031	0.6415	13351.	0.5226	1.0714
18-622	s sp.	Beds	235	21	235	57434	73149	59457
LAET752	Tragelaphu	Laeotil	3.6062	0.0028	0.8301	15714.	0.6808	0.9759
915	s sp.	Beds	99	22	745	69775	68684	01865
LAET75-	Tragelaphu	Laeotil	5.1162	0.0018	0.6904	14015.	0.7708	0.8198
722	s sp.	Beds	72	265	89	20803	44419	55393

# Curriculum vitae

## Jessica Renee Scott

Doctoral Program in Environmental Dynamics University of Arkansas 113 Ozark Hall Fayetteville, AR 72701

## **Education**

University of Arkansas, Ph.D. in Environmental Dynamics, expected December 2011 <u>Committee Chair:</u> Peter S. Ungar <u>Committee Members:</u> J. Michael Plavcan, Walter S. Manger and Rene Bobe <u>Dissertation:</u> Dental Microwear Texture Analysis of Pliocene Fossil Bovids from Four Early Hominin Sites in Eastern Africa: Implications for Paleoenvironmental Dynamics and Human Evolution

University of Arkansas, M.A. in Anthropology, 2007 <u>Committee Chair:</u> Peter S. Ungar <u>Committee Members:</u> Jerome C. Rose, Jesse Casana and Mark Teaford <u>Thesis:</u> Dental Microwear Texture Analysis of the Archaeolemurids and Megaladapids, Two Families of Subfossil Lemurs from Madagascar

University of Arkansas at Little Rock, B.A. in Anthropology, 2005

#### **Research Interests**

Environmental and ecological context of human evolution in East and South Africa Relationship between climatic and evolutionary change Community ecology of fossil mammals Dietary reconstruction using dental microwear and stable isotope analyses

## **Field Experience**

Amarna, Egypt	The Amarna Project	March-Apr	il 2008
Freighter Gap, Wyomi	ng The Great Div	vide Basin Project	June-July 2007
Beza Mahafaly, Madag	gascar The Ring-Tai	led Lemur Project	July-August 2006
LaSuerte, Costa Rica	Advanced Pr	imate Ecology J	uly-August 2005
Myakka City, Florida	Lemur Conse	ervation Foundation	May-June 2005
Toltec Mounds, Arkan	sas Arkansas Arc	chaeological Survey	July-October 2004

#### **Collections Experience**

Smithsonian National Museum of Natural History, Washington D.C. National Museum of Ethiopia, Addis Ababa Kenya National Museums, Nairobi Royal Museum of Central Africa, Tervuren, Belgium The Field Museum, Chicago Arkansas Archaeological Survey American Museum of Natural History, New York The Amarna Trust, Egypt

## **Other Research Experience**

Koobi Fora Plio-Pleistocene Bovid Project University of Arkansas Stable Isotope Lab, Department of Biology University of Arkansas Soil Lab, Department of Geosciences Graduate Research Assistant, J. Michael Plavcan Graduate Research Assistant, Peter S. Ungar

### **Grants and Awards**

NSF Doctoral Dissertation Improvement Grant (0925822), 2009-2011, \$15000 Walton Doctoral Academy Fellowship, August 2007- May 2011 University of Arkansas Conference Grant for AAPA meeting, April 2010, \$1100 University of Arkansas Conference Grant for AAPA meeting, April 2009, \$1100 University of Arkansas Conference Grant for AAPA meeting, April 2008, \$1100 University Study Abroad Grant for work at Amarna, Egypt, March- April 2008, \$500 University of Arkansas Conference Grant for AAPA meeting, April 2007, \$550 University of Arkansas Conference Grant for AAPA meeting, April 2007, \$550 Katherine J. Hardie Award for Outstanding Graduate in Anthropology, May 2005 UALR Student Fieldwork Award in Anthropology, May 2005, \$1000

#### **Professional Service**

Kenya National Museum, Invited Lecturer, May 2010 Department of Biological Sciences, Brown Bag Lunch Speaker, 2010 Day of Darwin Celebration, University of Arkansas, 2009 Pearson Publishing Text Development Seminar, March 2009 Fayetteville Middle School Science Fair Judge, 2008 University Days for high school students, 2005, 2006, 2007, 2008, 2009 Department of Anthropology, Colloquium Speaker, University of Arkansas, 2007 Primate Enrichment Coordinator, Little Rock Zoo, 2003, 2004, 2005

#### **Membership in Professional Organizations**

American Association of Physical Anthropologists

Society of Vertebrate Paleontology East African Association for Palaeoanthropology and Paleontology Dental Anthropology Association Sigma Xi Phi Kappa Phi Lambda Alpha

# **Publications**

Scott, J.R., in press.Dental microwear texture analysis of extant African Bovidae.Mammalia.

Stynder, D., **Scott, J.R.**, Schubert, B., Ungar, P.S., in review. Dental microwear texture analysis of fossil felids and hyaenids from Langebaanweg, South Africa.

Pontzer, H., Scott, J.R., Lordkipanidze, D., Ungar, P.S., in press. Dental microwear and diet in the Dmanisi hominins. Journal of Human Evolution.

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Merceron, G.M., **Scott, J.R.**, Scott, R.S., Geraads, D., Spassov, N., Ungar, P.S., 2009. Seed predation for an early Colobine as a link between frugivory and folivory?Evidence from dental microwear texture analysis of *Mesopithecus* (Late Miocene of Eurasia).Journal of Human Evolution.doi:10.1016/j.jhevol.2009.06.009

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Krueger, K., **Scott, J.R.**, Kay, R., Ungar, P., 2008. Comparisons of dental microwear texture attributes between facets in three primate taxa. American Journal of Physical Anthropology 137, 485-490.

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# **Abstracts**

**Scott, J.R.**, 2012. Dental microwear texture analysis of the fossil Bovidae from the Hadar hominin site: implications for paleoenvironment. American Association of Physical Anthropologists.

Ungar, P.S., **Scott, J.R.**, McNulty, K.P., Harcourt-Smith, W., Dunsworth, H.M., 2012.Environments of early Miocene Rusinga Island and Songhor: evidence from the dental microwear of tragulids. American Association of Physical Anthropologists.

**Scott, J.R.**, 2011. Dental microwear texture analysis of extant African Bovidae. Society of Vertebrate Paleontologists.

**Scott, J.R.**, Ungar, P.S., Schubert, B.W., Stynder, D.D., 2011. Dental microwear texture analysis of fossil hyaneids from Langebaanweg, South Africa. American Association of Physical Anthropologists.

Zolnierz, M.S., Delezene, L., Kimbel, W.H., **Scott, J.R.**, Ungar, P.S., 2011. Premolar microwear in *Australopithecus afarensis* and *A. africanus*. American Association of Physical Anthropologists.

Schmitt, E., Schubert, B.W., **Scott, J.R.**, Ungar, P.S., 2010. Analysis of the bonecrushing behavior of *Canis dirus* using dental microwear texture analysis. Society of Vertebrate Paleontology 2010.

**Scott, J.R.**, Grine, F.E., Teaford, M.F., Ungar, P.S., 2010. Premolar microwear texture analysis of *Australopithecus africanus*. American Journal of Physical Anthropology, S50.

**Scott, J.R.**, Krueger, K.L., Kemp, B., Rose, J.C., 2009. Dental microwear texture analysis of workers from Amarna, an eighteenth dynasty site in Egypt.American Journal of Physical Anthropology S48, 223.

Krueger, K.L., **Scott, J.R.**, Kay, R.F., Ungar, P.S., 2008. Comparisons of dental microwear texture attributes between facets in three primate taxa. American Journal of Physical Anthropology S46, 135.

**Scott, J.R.**, Ungar, P.S., Jungers, W.L., Godfrey, L.R., Scott, R.S., Simons, E.L., Teaford, M.F., Walker, A., 2007. Dental microwear texture analysis of the archaeolemurids and megaladapids, two families of subfossil lemurs from Madagascar. American Journal of Physical Anthropology S44, 212.

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