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Using springbok (Antidorcas) dietary proxies to reconstruct inferred palaeovegetational changes over 2 million years in Southern Africa

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ABSTRACT

The reconstruction of past vegetation and climatic conditions of the Cradle of Humankind, Gauteng Province, South Africa, has been approached using various proxies (such as micromammals, speleothems, faunal and floral presence and stable carbon isotopes). Elisabeth Vrba's seminal studies (1974; 1975) on the fossil record of this region indicated dramatic faunal turnover based on species extinction and speciation data. This turnover was thought to have been driven by increasing aridity and spreading grasslands. These reconstructions however, are continuously being refined and adapted in light of advancing techniques (such as dental microwear textural analysis) and terrestrial proxies, such as speleothems.

However, more recent studies show varying proportions from wooded towards more grassland-dominated habitats, with the most common reconstruction being the heterogeneous 'mosaic' habitat. Here we re-evaluate the findings of a transition from woodland to grassland conditions in the fossil record from Member 4 Sterkfontein to Member 5 Sterkfontein and the deposits of Swartkrans. To approach the palaeovegetation changes through time via a different angle, we focus on the diet of the springbok (genus *Antidorcas*), represented throughout this temporal period from geological members dating from 2.8–0.8 Ma. We use detailed dietary methods (dental linear measurements, mesowear, microwear, and stable carbon isotope analysis) to explore past changes in diets of springbok that can be used to indicate the prevailing vegetation conditions. Our results presented here broadly agree with previous palaeoenvironmental reconstructions, in indicating increased grassland post ca 1.7 Ma, with some suggestion of more heterogeneous habitats for Swartkrans Member 2 (ca 1.65–1.07 Ma). We find that there is support for the implementation of a multi-disciplinary approach to produce more accurate and robust reconstructions of past diets and by extension, of palaeovegetation conditions, if the selected herbivore species is a mixed-feeder, like the springbok.

1. Introduction

Climatic aridification and spread of grasslands are a key feature of African Plio-Pleistocene environmental changes (e.g. Sepulchre et al., 2006; Ségalen et al., 2007; Hopley et al., 2007; deMenocal, 2011; Kaya et al., 2018), which are inferred to have caused dramatic habitat changes that impacted on our ancestors, as well as other fauna. There is broad agreement, from a range of different palaeoenvironmental proxies, (from climatic proxies such as deMenocal's (1995, 2004) marine sediment sequences to terrestrial habitat proxies such as Avery's (2001) micrommamals analyses, that Africa experienced gradually increasing aridity (Bobe and Eck, 2001; Bobe and Behrensmeyer, 2004; Avery, 2001; deMenocal, 1995, 2004 amongst others) and a spread of grasslands through time after 5 Ma, with a resulting radiation in grassland-adapted taxa (Vrba, 1973, 1985). Yet recent studies cast doubt on this prevailing trend for east Africa (Blumenthal et al., 2017; Kaya et al., 2018), heralding the need to revisit this long-standing hypothesis.

An increasing number of hominin species are now known to have been present in the southern African region throughout this temporal period: *Australopithecus sediba* (Berger et al., 2010), *Australopithecus africanus* (Dart, 1925; Clarke, 2008) *Paranthropus robustus* (Broom, 1938), *Homo habilis* (Hughes and Tobias, 1977) and *Homo erectus/ergaster* (Leakey, 1960; Kuman and Clarke, 2000; Antón et al., 2014).

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During this time of climate and habitat change, we see the disappearance of *Australopithecus* and the appearance of *Homo* and *Paranthropus*. What role, if any, did climate-driven habitat shifts have in the appearance and extinction of these hominins? Here, we are attempting to use dietary variations in other members of the faunal community for insights into wider patterns of vegetation and habitat change.

Within ecological proxies, the diversity of ungulate mammals and notably bovids has been frequently studied (e.g. Vrba, 1974; Spencer, 1997). Amongst bovids, the species of the genera *Antidorcas* is continuously present throughout this temporal period of interest in South Africa.

There are several Antidorcas species, of which only one remains today, Antidorcas marsupialis, Modern springbok are typically aridadapted, mixed feeders, with a relatively wide habitat tolerance primarily in the more arid areas of south and southwestern Africa; South Africa, Namibia, Botswana and Angola (Lehmann, 2015; Castelló, 2016). Extant A. marsupialis will graze on young grasses when available, swapping to browse on low shrubs and succulents, when young grasses are not available. Springbok can survive without drinking water, as long as their food contains > 10% water content (Estes, 1991). Historic records make mention of the mass migrations of these animals in search of better quality grazing after the rains (Skinner and Louw, 1996). Social organization alters according to season, sex and age range of the individual springbok (Skinner and Chimimba, 2005). When considering dietary signals from fossil Antidorcas, social and migratory behaviour should be considered to account for the range of signals obtained (e.g. Nagy and Knight, 1994; Bednekoff and Ritter, 1997; Stapelberg et al., 2008).

Fossil springbok will henceforth be referred to by genus (i.e. "*Antidorcas*"). In the fossil record, there were several *Antidorcas* species; based on tooth enamel carbon stable isotopes, *A. recki, A. australis, A. bondi and A. marsupialis* appear to have been mixed feeders to varying degrees, with the exception of *A. bondi*, which was suggested to have been a specialist grazer (Brink and Lee-Thorp, 1992).

The earliest species, Antidorcas recki (Schwarz, 1932) was presumed to tend towards browsing (Luyt, 2001). It is assumed that A. recki is the species from which Antidorcas australis (Hendey and Hendey, 1968) and/or Antidorcas marsupialis (Zimmerman, 1780) and A. bondi (Cooke and Wells, 1951) descends. Antidorcas bondi, had a very specific small body-size and specialist dentition (with characteristically small, but very high-crowned, hyposdont teeth) and a unique dietary ecology (based on stable carbon isotope data) that enabled it to exist as part of a grazing succession in southern Africa prior to its extinction at the end of the Pleistocene/early Holocene (Brink, 2016; Brink et al., 2013; Brink and Lee-Thorp, 1992). Brink and Lee-Thorp (1992) proposed A. bondi to be the smallest member of a grazing succession feeding on new grass shoots whose growth was stimulated by the regular 'mowing' of grasses by extant and extinct species, such as the giant buffalo (Pelorovis antiquus), giant hartebeest (Megalotragus priscus), large equid (Equus capensis) and the medium-sized hartebeest (Damaliscus niro) in a similar way to that proposed for the Serengeti in East Africa by Gwynne and Bell (1968).

The presence and palaeoecology of *A. australis* as a potentially separate species, or a subspecies is debated, but remains unresolved (e.g. Gentry, 1978, 1992, 2011; Vrba, 1973). *Antidorcas australis*, was originally described from the Middle Stone Age (MSA) site of Swartklip (Cape Province, South Africa) by Hendey and Hendey (1968) as a transitional form of *Antidorcas* between the ancestral *A. recki* and the descendant *A. marsupialis*, consequently tentatively assigned as a subspecies of *Antidorcas marsupialis*. *A. australis* was subsequently elevated by Vrba (1973) to a separate species, supported by Hendey (1974) as a valid taxon, when found alongside *A. recki*. However, Vrba (1976) later doubted this elevation, suggesting *A. australis* was perhaps only a southern Cape endemic form. Gentry (2010) suspects *A. australis* and *A. marsupialis* to be synonymous, but not all authors agree (e.g. Klein, 1980; Faith, 2014).

Initially all *Antidorcas* specimens are considered together as *'Antidorcas'* with no prior assumptions based on taxonomy and a second time, distinguishing specimens according to their assigned taxonomic category (see Table 1 and Fig. 1).

The number of species of *Antidorcas* in the past is important in understanding the overall function of the palaeoecosystem. If herbivore species can only coexist by species 'packing' into narrow niche spaces to reduce competition (e.g. Richie and Olff, 1999; Codron et al., 2008) then we should see that *A. australis* possesses indications of a distinct dietary niche, and or/different body sizes compared to other *Antidorcas* congeners. Can dietary signals from dental specimens of proposed *Antidorcas australis*, combined with contemporaneous *Antidorcas marsupialis*, establish the likelihood from dental and dietary evidence, of *Antidorcas australis* as a separate species?

From the dietary indicators of *Antidorcas*, we establish a broad palaoevegetation reconstruction. One way to reconstruct palaeovegetation cover would be to examine direct evidence of the plants themselves (fossil plants, pollen and phytoliths) preserved at sites like Sterkfontein and Swartkrans, Cradle of Humankind, Gauteng Province, South Africa. However, since plant fossils and pollen are not commonly preserved in the Cradle of Humankind sites (but see Bamford, 1999), we use the dietary evidence from a mixed feeding herbivore that is commonly preserved at these sites, and one which is relatively abundant. Herbivores are considered faithful reflectors of prevailing habitat conditions, over decades of detailed palaeoenvironmental research that used various aspects of herbivore palaeoecology (Vrba, 1974, 1975; Plummer and Bishop, 1994; Spencer, 1997; Reed, 1997; Bobe and Eck, 2001; amongst many others).

A key transition in the southern African fossil record is between Sterkfontein Member 4 (where Australopithecus africanus is abundant) and Member 5 (where we have no Australopithecus, but instead Homo and Paranthropus). The typical palaeoenvironmental story is one of wetter woodland and closed habitats up to and including Sterkfontein Member 4 (c. 2.8-2.0 Ma), with the area becoming more open with drier, grassland-dominated habitats by Sterkfontein Member 5 times which overlap with Swartkrans Members (2-0.8 Ma) (Bamford, 1999; Kuman and Clarke, 2000; Luyt, 2001). Mosaic habitats (Brophy, 2011; Reynolds et al., 2015) and local variations have been reported between the sites lying within the Cradle of Humankind, presumably due to the varying proximity to the Blaaubank River (Avery, 2001; Reynolds and Kibii, 2011). The cave stratigraphy and dating of these South African deposits have received much attention (e.g. Hopley and Maslin, 2010; de Ruiter et al., 2008) and a summary of the dates of the deposits are provided in Table 2. There are very few genera of herbivores that occur in both Member 4 and Member 5; which is chiefly characterized by a dramatic turnover in species (e.g. Vrba, 1974, 1975; de Ruiter, 2003a, 2003b; Reynolds and Kibii, 2011). We focus on the only single and successful lineage that is present in Member 4 (as Antidorcas recki) and in later Member 5 (as A. marsupialis).

The majority of previous studies typically employ only one or two methods (e.g. microwear and/or stable isotope analyis) to tackle aspects of palaeoenvironmental questions. Each method reveals important aspects relevant to the overall palaeoenvironmental picture, but these methods also have inherent limitations. For example, microwear indicates primarily the last few weeks as new micrometric marks occur after each meal eaten (Teaford and Oyen, 1989), mesowear requires observer experience and therefore large sample sizes to mitigate for this (Kaiser and Fortelius, 2003), and isotopes are informative only of a short window in the early years of life during enamel formation (Reade et al., 2015). Comparisons between species are also complicated by

Table 1

Features typical of each Antidorcas species found in Southern Africa, used to aid in species identification of fossil dentition. Characteristic features used to establish taxonomic assignment of fossil dental specimens (Vrba, 1973; Cooke and Wells, 1951; Gentry, 2010, 2011). Typical M² dimensions established according to the type specimens, from corresponding reference as stated in the table.

Species	Antidorcas bondi	Antidorcas recki	Antidorcas marsupialis	Antidorcas cf. australis
Reference Figure Characteristic dental features	 Cooke and Wells, 1951 See Fig. 1, Images F-H Upper: Strongly developed styles (are often basally (i.e. towards the body of the tooth) pinched, giving a more prominent impression). Upper: Complicated folding of enamel walls of central cavities, with a greater tendency towards dumbeell- shaped central cavities and increased curvature Extreme hypsodonty Lower: a depression runs from occlusal surface to root the molar walls appear to "bow out", giving the whole lingual molar roles awary look (more similar to impala than the other <i>Antidorcus</i> species). Lower: Sometimes present: lingual "rib" (metastylid) in the other <i>Antidorcus</i> species). Lower: Central cavities are centrally constricted, creating a dumbbell appearance. Laver BM3 is larger and more complex than in <i>A. marsupialis</i>. Typically Retains PM2 PM3 is larger and more complex than in <i>A. marsupialis</i>. Lower: Occasionally goat folds haal pillars are present. Tipchens of folds basal pillars are present. Tipchens of folds basal pillars are present. Tipchens of folds basal pillars are present. 	 Schwarz, 1932 See Fig. 1, Images A-B Unbent central cavities Buccal lophs are V-shaped (more so than the more rounded ones of A marupialis) Has styles on upper molars, which are more often V-shaped Upper: Less strongly pronounced concave posterior part of lateral wall (than in A. marupialis/A. bond). Upper: Stronger styles than A. marsupialis, with a flatter and even concave wall between mesostyle and metastyle. P.2 is usually retained (sometimes absent). Lower: Central cavities straighten in relatively early wear stages. Large M₃ metastyle. Typically Retains PM₂ PM₃ is larger and more complex than in A. marupialis. 	 Zimmernan, 1780 See Fig. 1, Image I Lower: Lingual walls are straight (less outwardly bowed than in the impala). Lower: Central cavities straighten in relatively early wear stages Large M₃ metastyle, with noticeably enlarge third (hypoconulid) lobes Lacks PM₃ Reduced PM₃ 	 Hendey and Hendey, 1968 See Fig. 1, Images C-E Molars are bucco-lingually narrower in mersiodistal length (Vrba, 1970) Mower: Central cavities straighten in relatively early wear stages Large M₃ metastyle. Large PM₃ Reduced PM₃ Narrower than A. marstpialis and appear more gracile (Vrba, 1970)
Mean M ² dimensions	spectes. 16mm × 9mm ^a	$13.3\mathrm{mm} imes 6.3\mathrm{mm}^\mathrm{b}$	17.4 mm (Female) 18 mm (Male) $\times 10.9 \text{ mm}^{\circ}$	$12.5 \mathrm{mm} imes 7.95 \mathrm{mm}^{\mathrm{d}}$
^a Cooke and Wells, 19: ^b mesiodistal length \times	:1; dimensions written as length \times breadth (in mm). bucco-lingual width. Based on mean measurements taken of <i>A. reck</i>	ci from Sterkfontein and Swartkrans by L. Sewell (20)	(6).	

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5 Ę. g ⁴ Based on mean measurements (by L. Sewell and Vrba, 1973) of specimens assigned to A. *australis*, mesiodistal length × buccolingual width.



Fig. 1. Examples of Antidorcas maxillary (upper) dentition. All photographs taken by L. Sewell unless otherwise stated*. A-B: Antidorcas recki (A = KA2610; *B = KNM-ER 6–18), C-E: Antidorcas australis as identified by Vrba (1973): (C = SK 3055; D and E = SK 2115), F–H: Antidorcas bondi (F = SF 592; G = KA 2472; H = SK 2366), I: modern Antidorcas marsupiualis. [Key: KA = Kromdraai, KNM-ER = Kenya National Museum-East Rudolph [*photograph by Lucile Crété], SK = Swartkrans, SF = Sterkfontein]. Scale bars equal 10 mm.

tooth development varying inter-specifically (Reade et al., 2015). There is, therefore, support for a multi-disciplinary approach to produce more holistic, accurate and robust reconstructions of past diets (Schubert et al., 2006) and by extension, of palaeovegetation. Consequently, we take a multi-method approach here to understand when, and how, each method can be used. Our palaeoenvironmental picture obtained from measurements of dental molars, mesowear and microwear analyses, along with new stable isotope data (collected by SCR and PJH in 2002) are compared to the palaeoenvironmental signals gained from stable isotope analysis by Lee-Thorp et al. (2007).

In this paper, we report on our findings from each method; these methods are typical of much of the current research in palaeoanthropology. The results presented are a subset of results for the wider project underway, used here to reflect on the accuracy and limitations of each method. We use these datasets to address two main questions: 1) can palaeodietary information be used to reconstruct habitat conditions through time; and 2) what additional detail can we gain about the dietary differences between the different *Antidorcas* species in the fossil record?

2. Materials and methods

2.1. Materials

A sample of 287 fossil *Antidorcas* teeth from Sterkfontein and Swartkrans, Cradle of Humankind, Gauteng Province, South Africa were examined for this study. All species of *Antidorcas* present in the deposits are considered. Specimens are taxonomically identified following the criteria set out in Table 1.

Modern springbok specimens were also examined (n = 33 individuals). Only permanent upper and lower second molars (M^2 and M_2) were selected for study in the first instance. Where unavailable, permanent upper and lower first or third molars are used for use-wear analyses instead. Substitutions of the permanent molar teeth have been shown to produce comparable dietary signals (Scott, 2012; Ramdarshan et al., 2017 for dental microwear textural analysis, and Kaiser and Solounias, 2003 for dental mesowear analysis, respectively). Upper and lower second molar were separated for metric (linear dental measurements) analyses.

All statistical analyses were run in software package SPSS (v.23). All results presented here are used to showcase particular aspects of each methodology, highlighting where each method can be implemented to obtain the most accurate results. All data are rank transformed prior to analysis to mitigate for small and varying sample sizes.

2.1.1. Collections used

Sterkfontein Member 4 and 5 (Stw53 Infill, East and West), Swartkrans Member 031 (Lower Bank and Hanging Remnant), Member 2 and Member 3. Fossil collections are housed at the Evolutionary Studies Institute (ESI), University of the Witwatersrand, Johannesburg and the Ditsong Museum of Natural History (DM), Pretoria, South Africa. Additional material from modern Mammalogy collections (wild caught) was studied to establish the anticipated range of variation. Modern collections are housed at the Natural History Museum (NHM), London and the National Museums Scotland (NMS), Edinburgh, United Kingdom and the Ditsong Museum of Natural History, Pretoria, South

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Table 2

Dates ranges and methods used to obtain these dates for each member used. Maximum date ranges are indicated below each Member row. Member deposits ordered chronologically relative to one another (oldest to youngest) based on the dates provided in the table.

Member	Date	Method	Reference
Sterkfontein Member 4	2.8–2.4 Ma	Bovid biochronology	Vrba, 1976, 1980
	~2.5 Ma	Primate biochronology	Delson, 1988
	2.8–2.6 Ma	Mammalian biochronology	McKee, 1993
	~2.1 Ma	ESR (electron spin resonance)	
		Palaeomagnetism	Schwarcz et al., 1994
	2.15–2.14 Ma	Uranium-lead	
	2.65–2.01 Ma	ESR, and palaeomagnetism	Partridge, 2005
			Pickering and Kramers, 2010
	2.8–~2.0 Ma		Herries and Shaw, 2011
Date range: 2.8-2.0 Ma.			
Sterkfontein Stw infill	2.6–2.0 Ma	Biochronology	Kuman and Clarke, 2000
	1.8–1.4 Ma	Biostratigraphy	Herries et al., 2009
	1.8–1.5 Ma	ESR and palaeomagnetism	Herries and Shaw, 2011
Date range: 2.6–1.4 Ma.			
Sterkfontein M5 East	2.0–1.7 Ma	Biochronology and archaeology	Kuman and Clarke, 2000
		Dating seriation	
	1.4–1.1 Ma	ESR, isotopes and	Herries et al., 2009
	1.4–1.2 Ma	palaeomagnetism	Herries and Shaw, 2011
Date range: 2.0–1.1 Ma.			
Sterkfontein M5 West	1.7–1.4 Ma	Biochronology and archaeology Dating seriation	Kuman and Clarke, 2000
	1.3–0.8 Ma	ESR and palaeomagnetism	Herries et al., 2009
	1.3–1.1 Ma		Herries and Shaw, 2011
Date range: 1.7–0.8 Ma.			
Swartkrans Member 1 LB	1.7 Ma	Biostratigraphy	Vrba, 1985; Churcher and Watson, 1993; de Ruiter, 2003a, 2003b; Brain, 1995, Vrba, 2000
	1.8 Ma	Biostratigraphy	
Swartkrans Member 1 HR	1.6 Ma	Biostratigraphy	Vrba, 1982, 1985; Delson, 1984; Brain, 1995; Berger et al., 2002; de
			Ruiter, 2003a, 2003b
Swartkrans Member 1	2.0–1.4 Ma	ESR	Curnoe et al., 2001
(combined)	2.0 (± 0.02) Ma	U-Pb bovid enamel dating	Albarède et al., 2006
Date range: 2.0–1.4 Ma.			
Swartkrans Member 2	1.36 (± 0.29) Ma (1.65–1.07 Ma)	U-Pb enamel dating	Balter et al., 2008
	1.44 ± 0.05	U-Pb enamel dating	Albarède et al., 2006
	1.7–1.1 Ma	Biochronology	Brain, 1995; Vrba, 1995a, 1995b: Herries et al., 2009
		Dating seriation	
Date range: 1.7–1.07 Ma.		C C	
Swartkrans Member 3	0.83(± 0.21)Ma (1.04–0.61 Ma) 0.988 ± 0.003	U-Pb enamel dating	Balter et al., 2008
	1.5–0.7 Ma	U-Pb enamel dating	Albarède et al., 2006
		Biochronology	Brain, 1993; Vrba, 1995a, 1995b; Herries et al., 2009
Date range: 1.5–0.61 Ma.			

Africa.

Additional modern collections were used for dental microwear textural analysis (DMTA). In addition to NHM and NMS, the following institutions' collections were used: MNHM Paris, France; Zoologisches Museum Berlin, Germany; Staatliches Museum für Naturkunde of Stuttgart, Germany; Senckenberg Naturkundemuseum of Frankfurt, Germany; NHM Basel, Switzerland.

2.2. Methods

Due to the uncertainty surrounding the taxonomic status of *A. australis*, (according to features in Table 1), *A. australis* and *A. marsupialis* fossil specimens were analysed together under '*Antidorcas* sp.'

Other species of specific dietary preference, such as the grazing blesbok (*Damaliscus pygargus/dorcas*) are occasionally used to establish the fossil dietary parameters and understand the dietary signals gained from *Antidorcas*.

2.2.1. Linear measurements of teeth

Tooth dimensions obtained from measurements of mesiodistal length (MDL) and bucco-lingual width (BLW) are considered for this study. Measurements were taken in millimeters with carbon fiber composites digital calipers (resolution: 0.1 mm/0.001").

2.2.2. Mesowear

Mesowear provides an indication of an individual's dietary signal through their lifetime. Mesowear scores are informative on the abrasiveness of the diet and can be used as a reflection of the grazing/browsing composition of diet. Grazers' molars typically exhibit low occlusal relief and blunter cusps due to the abrasive nature of a predominantly grazing diet. Browsers on the other hand, normally display higher occlusal relief, typically with sharper cusps (Fortelius and Solounias, 2000). Following the protocol established by Fortelius and Solounias (2000), numerical values are given to each tooth based on occlusal relief (High or Low) and cusp shape (Sharp, rounded or blunt) as set out in Fig. 2. The numerical value (score) is given as follows: Low Blunt:0, Low Rounded:1, Low sharp:2, High rounded:3, High sharp:4. Upper molars are used for mesowear analysis as lower molars have a tendency towards blunting and are consequently likely to be exhibiting a predominance of grazing signals (Kaiser and Fortelius, 2003).

2.2.3. Dental microwear textural analysis (DMTA)

Dental microwear textures provide insight into the diet over the last few weeks/days of an individual animal's lifetime (Teaford and Oyen,

Fat	$\wedge \wedge$	Steep valley between cusps	HIGH Occlusal Relief
	\sim	Shallow valley between cusps	LOW Occlusal Relief
	\land	Sharp cusps	SHARP Cusp Shape
TO		Rounded cusps	ROUNDED Cusp Shape
		No distinct facets	BLUNT Cusp Shape

Fig. 2. Mesowear scoring figure: Traditional mesowear scoring system. Established by Fortelius and Solounias (2000), figure modified from Merceron et al. (2007).



Fig. 3. DMTA image: Dental microwear texture analysis is run on the disto-buccal facets of the buccal cuspids along the lower arch. (Modified from Merceron et al., 2016.)

1989; Schulz et al., 2013; Merceron et al., 2016; Ramdarshan et al., 2016). Upper and lower molars were used for microwear analysis to allow direct comparisons across species and deposits. Because values from all positions (M1 to M3) of permanent molars can be used together in statistical analysis as their dental microwear textures have been shown to be statistically similar (see Ramdarshan et al., 2017). Data on lower molars can be grouped with upper molars if specific facets are considered (see below; see Ramdarshan et al., 2017). This is beneficial when dealing with isolated fossil dentition that may not be as clearly identifiable to a particular tooth.

Dental microwear texture analysis (DMTA) was conducted according to Merceron et al. (2016). Molds of the dental enamel facets of interest, on the disto-buccal facets of the protoconid (or on the hypoconid if the former facet is broken), and on mesio-lingual facets of the protocone (Fig. 3), were created using polyvinvylsiloxane elastomer. Ramdarshan et al. (2017) have shown that there are significant differences in dental microwear textures between the disto-buccal facets of the protoconid on lower teeth and the lingual facet of the paracone on upper; but not with the protocone from upper molars (see discussion in Ramdarshan et al., 2017). This mold is cut to provide a relatively small and flat surface, for viewing under the surface profilometer (Leica DCM8 confocal profilometer). The prepared molds are placed under a Leica DCM8 confocal profilometer using white light confocal technology with a Leica $100 \times$ objective (Numerical aperture = 0.90; working distance = 0.9 mm). The lateral resolution is an (x, y) interval of 0.129 µm, with a vertical numerical step of 1 nm. Scans are taken in the centre of the facet, where possible, to ensure repeatability and to limit subjectivity. Scans are treated using LeicaMap to eliminate artefacts, such as abnormal peaks following procedures detailed in the supplementary material in Merceron et al. (2016). Data analyses for DMTA are run through Sfrax and Toothfrax software, as set out by Scott et al. (2006).

Complexity (Area-scale fractal complexity, *Asfc*) measures the roughness of the dental facet surface. Scale of maximum complexity (*Smc*) shows the scale range *Asfc* is calculated from (taking the steepest part of the relative area against the scale of the curve). Anisotropy considers the direction of wear and relative lengths of vectors (depth profiles), essentially providing a view of direction of wear and a glimpse into heterogeneity (Scott et al., 2006). Multiple scratches with no pitting would produce anisotropic surfaces with a high *epLsar* value, whereas a surface with scratches in opposing directions coupled with pits of varying depths and sizes would produce a complex (high Asfc)

and isotropic (low *epLsar*) surface and be more indicative of browsing diets (Scott, 2012; Scott et al., 2006). Homogeneity of wear is provided through *HAsfc* values (Heterogeneity of Area-scale fractal analysis). Higher *HAsfc* values indicate greater variation in diet. Homogeneity of diet tends to show in homogeneity of surface texture (i.e. lower HAsfc values) (Scott et al., 2006). *HAsfc* are calculated through a 9- and an 81-cell mesh over each scanned surface. *Tfv* (Textural fill volume) values represent the surface texture based on fill volume at different scales (i.e. the relief of the surface) (Scott et al., 2006). Considerable research has focused on the nature of microwear signatures in whether dust and grit cloud the dietary signals obtained. Ungar et al. (2016) and Merceron et al. (2016) show that overall, microwear data is informative of dietary preferences and thus, in this case also informative about palaeovege tation.

Data are rank transformed and a one-way ANOVA with post hoc Tukey's HSD and Fisher's LSD tests are performed, once to establish species dietary differences and a second time, to understand dietary differences (for all *Antidorcas* species combined) between members.

2.2.4. Stable carbon isotope analysis

Stable carbon isotope analysis was made possible via temporary export and destructive sampling permits granted by SAHRA (South African Heritage Resource Agency). Stable carbon isotope values provide a dietary signal from the early years of an individual animal's lifetime. The carbon values are reflective of the C3/C4 composition of the diet during the time of dental-enamel formation. Carbon isotopes indicate the photosynthetic pathways used by the vegetation and subsequently consumed by the animal sampled. C3 plants (indicative of a predominantly browsing diet) are more depleted in carbon (δ^{13} C) than C₄ plants (indicative of a predominantly grazing diet) (Vogel, 1978; Luyt, 2001). Isotope analysis complements use-wear (mesowear and microwear) analysis. Isotopes contrast with microwear (meals of last days to weeks) and mesowear (averaged lifetime) results, by providing the early years (during dental enamel formation) signal. To obtain the isotopic information, approximately 5-10 mg of enamel powder was extracted from each tooth using a diamond-tipped drill bit, as a bulk sample. Samples were then cleaned using standard pre-treatment methods for the removal of organic and carbonate contaminants (Lee-Thorp et al., 1997; Van der Merwe et al., 2003). Carbon and oxygen isotopic measurements were obtained using a VG SIRA 12 mass spectrometer at the University of Liverpool (see Hopley et al., 2006 for further details). Long-term laboratory reproducibility (o) is better than $\pm 0.1\%$ for both isotope ratios.

3. Results

3.1. Linear measurements of teeth

Modern springbok are sexually dimorphic, with males having larger body size and larger, more robust horn sizes (e.g., Skinner and Chimimba, 2005; Brakora, 2014). To rule out sexual dimorphism, we tested modern *Antidorcas* data (of known sex) for significant differences in MDL and BLW measurements. No significant differences were found (Mann-Whitney *U* test p > 0.05 for occlusal length; p > 0.05 for bucco-lingual width). These results show that male and female springbok teeth do not differ significantly in size. Consequently, we consider that sexual dimorphism in the fossil species is unlikely to confound the potential size differences where they are observed in the fossil record.

Both *A. recki* and *A. bondi* can clearly be distinguished from *A. marsupialis* based on their consistently smaller size (Table 3). In contrast, there is considerable overlap between specimens previously assigned to *Antidorcas australis* (Vrba, 1973) compared to those assigned

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Table 3

Mean M^2 measurements for each Antidorcas species. Measurements are separated according to deposit (site and Member). MDL = mesiodistal length; BLW = bucco-lingual width.

Species	Provenance	N	Mean MDL	Mean BLW
Antidorcas marsupialis/ australis	Swartkrans Member 1	3	14.1	9.6
	Swartkrans Member 2	1	14.7	9.4
	Swartkrans Member 3	22	14.8	9.4
	Cave of Hearths	5	15.0	11.3
Antidorcas marsupialis	Modern	44	16.3	11.7
Antidorcas recki	Sterkfontein Member 4	1	14.7	6.0
	Cooper's Cave (D)	1	14.7	10.4
Antidorcas bondi	Swartkrans Member 2	11	13.9	8.6
	Cave of Hearths	3	15.1	8.5
Antidorcas sp. ^a	Swartkrans Member 1	1	13.4	10.4
	Swarktrans Member 2	18	14.4	9.8
	Swartkrans Member 3	2	13.1	11.4

^a In this table, Antidorcas sp. refers to specimens identifiable only to genus level.

to Antidorcas marsupialis. For example, in Swartkrans Member 1, the bucco-lingual width of A. australis (n = 10) ranges from 4.5–12.5 mm and A. marsupialis (n = 3) ranges from 5.5–10.2 mm.

Statistical analysis of mesiodistal length (MDL) and bucco-lingual width (BLW) (see Fig. 4) showed that *Antidorcas bondi* and *Antidorcas* sp. are significantly different for Swartkrans Member 2 (MDL p = 0.014; BLW p = 0.08), perhaps suggesting the presence of another *Antidorcas* species, other than *Antidorcas bondi*. BLW significantly increased (Kruskal-Wallis test) from Swartkrans Member 1 to Swartkrans Member 2 for specimens assigned to the following taxa: *Antidorcas* sp. (p = 0.012), *A. bondi* (p = 0.005) and *A. marsupialis* (p = 0.004). These results suggest the presence of two species of *Antidorcas* in Swartkrans Member 1 and Member 2, both of which exhibit dental size changes through time. Whilst some specimens are identifiable only to genus (*Antidorcas* sp.), we can assert that the bucco-lingual width increases for the *Antidorcas* genus from Swartkrans Member 1 to Swartkrans Member 2.

Antidorcas bondi second molar mesiodistal length significantly increases between Swartkrans Member 1 (1.7–1.1 Ma) and Swartkrans Member 2 (0.7–0.6 Ma). The mean mesiodistal length of the sample of *A. bondi* second molars significantly increased (Independent samples Kruskal-Wallis Test (p = 0.018), with post hoc independent samples *t*-test p = 0.038) from Member 1 times (mean length 12.9 mm) to Member 2 times (mean length 14.5 mm).

3.2. Mesowear

Mesowear scores from this study are given in Table 4. There is a good level of agreement with our data and the existing dietary inferences for each species.

Specifically, the extinct *A. bondi* differs from the modern *A. marsupialis* in having higher frequency of molars with low occlusal relief, together with rounded to blunt cusps. This is suggestive of more grazing dietary habits than the modern forms, as previously stated by Brink and Lee-Thorp (1992). Although low in sample size, *A. recki* exhibits a primarily browsing mesowear signature of high relief, with sharp cusps, thus supporting previous isotopic results by Luyt (2001) and other

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Fig. 4. Scatter plot of *Antidorcas* linear measurements. Morphological measurements of permanent A) lower second molars for all members (bucco-lingual width against mesiodistal length) according to species. (*Antidorcas bondi* n = 9, *Antidorcas recki* n = 5, *Antidorcas* sp. (*A. australis/marsupialis*) n = 28, modern *Antidorcas marsupialis* n = 36). Modern and fossil *A. marsupalis* and *A. australis* show significant overlap in dimensions. B) upper second molars for all members (bucco-lingual width against mesiodistal length) according to species.

Table 4

Antidorcas mesowear values. Relief (%high/low) and cusp shape (%sharp, rounded and blunt), 'N' = number of specimens Table A: Mesowear by species, B: Mesowear for all the *Antidorcas* genus through time, (ordered oldest to youngest according to member), C: Mesowear through time, separated for each *Antidorcas* species. *Antidorcas* sp. represents *Antidorcas* identifiable only to genus. "*Antidorcas marsupialis*" represents specimens taxonomically assigned to *A. australis* or *A. marsupialis* (with the exception of the modern specimens).

Anidarous sp., Antidorus shorid 27 63 97 44 56 0 Antidorus shorid 43 73 25 10 97 40 0 0 Antidorus shorid 42 74 26 48 40 12 Antidorus manapialis 42 74 26 48 60 12 Antidorus manapialis 42 73 26 48 50 8 B. Provenance N % High % Low % Sharp % Nowded % Bitmedia Stefontein 31 48 52 48 2 0 0 Stefontein Member 10 60 40 40 60 0 0 Stefontein Member 3 100 0 100 0 0 0 0 Stefontein Member 1 3 100 0 100 0 0 0 0 0 Stefontein Member 2 22 64 56 41 52 7 0 Sterdontein Member 3 Species	A: Species	N	% High		% Low	% Sharp		%Rounded	%Blunt
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	Modern	Antidorcas marsup	ialis	24	62.5	37.5	33	58	8

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Fig. 5. Biplots (Asfc on X axis vs epLsar on Y axis) of modern ruminants and extinct species of Antidorcas from South African Cradle of Humankind sites. The two lower plots show overlap, and therefore mixed-feeding signatures, when compared to modern obligate browsing and obligate grazing species.

studies of postcranial ecomorphology of this species from Olduvai Gorge, Tanzania (Plummer and Bishop, 1994).

3.3. Microwear

The dental microwear textures of modern *A. marsupialis* display microwear values expected for mixed feeders with medium values in complexity and high values in anisotropy.

This data shows a wide *Asfc* and *epLsar* distribution for fossil *Antidorcas* also, indicative of mixed-feeding dietary habits, similar to modern springbok (Fig. 5).

As shown in Tables 5 and 6, significant differences between species emerge.

Tables 5 and 7 show that all extinct species of Antidorcas differ from

the modern grazing taxa-Alcelaphus buselaphus and Syncerus caffer, in having higher Asfc. All fossil Antidorcas differ from the leaf-browsing Giraffa camelopardalis, in having higher Asfc. All fossil Antidorcas, excluding A. recki have higher *epLsar* than the two modern species of browsing ruminants. Finally, all fossil Antidorcas species have higher *Tfv* than modern species (except modern springbok; Tables 5 and 7).

As indicated (Table 9), there are no significant differences for *Antidorcas* microwear texture parameters between Members (through time), suggestive of varying degrees of mixed habitats being available throughout this temporal period.

Considering Table 8, the highest *Asfc* and lowest *epLsar* values for Sterkfontein Member 4 could support the notion of greater woodland in the area, with *Antidorcas* including slightly more browse in the diet, than at Swartkrans (Member 2 and 3).

Table 5

Summary dental microwear texture parameter statistics. Mean (M), standard deviation (SD) and standard error of the mean (SEM) for all species. Antidorcas sp. refers to those identifiable only to genus level.

Species		Ν	Asfc			epLsar			HAsfc	9		HAsfc	81		TFV2		
			М	SD	SEM	М	SD	SEM	М	SD	SEM	М	SD	SEM	М	SD	SEM
Fossil	Antidorcas sp.	7	4.61	2.79	1.06	0.00585	0.00221	0.00083	0.56	0.14	0.05	0.86	0.26	0.10	72,548.6	12,914.2	4881.1
	A.bondi	12	6.16	4.05	1.17	0.00479	0.00229	0.00066	0.52	0.34	0.10	0.85	0.44	0.13	63,763.9	8943.1	2581.6
	A.marsupialis/australis	10	3.98	3.79	1.20	0.00692	0.00395	0.00125	0.47	0.33	0.10	0.75	0.36	0.12	63,742.3	12,917.9	4085.0
	A.recki	5	7.51	4.25	1.90	0.00368	0.00250	0.00112	0.52	0.27	0.12	1.06	0.49	0.22	56,845.1	16,825.5	7524.6
Modern	Alcelaphus buselaphus	31	1.59	0.70	0.13	0.00536	0.00183	0.00033	0.34	0.21	0.04	0.59	0.22	0.04	50,449.0	8450.2	1517.7
	Cephalophus sylvicultor	27	3.50	2.63	0.51	0.00304	0.00173	0.00033	0.34	0.13	0.02	0.61	0.21	0.04	4.0349.2	8450.2	1517.7
	Giraffa camelopardalis	16	1.97	0.91	0.23	0.00217	0.00176	0.00044	0.40	0.33	0.08	0.68	0.49	0.12	27,251.7	19,050.1	4762.5
	Syncerus caffer	23	1.53	0.84	0.18	0.00456	0.00254	0.00053	0.32	0.19	0.04	0.57	0.28	0.06	46,026.2	8556.2	1784.1
	Damaliscus pygargus	8	2.38	0.58	0.21	0.00453	0.00308	0.00109	0.56	0.31	0.11	0.90	0.36	0.13	60,687.7	7822.0	2765.5
	Antidorcas marsupialis	16	3.32	2.40	0.60	0.00733	0.00360	0.00090	0.53	0.32	0.08	1.02	0.64	0.16	63,222.4	18,965.2	4741.3

Table 6

DMTA analysis of variance. One-way ANOVAs on dental microwear textural parameters to explore variations between modern and extinct species of ruminants. df: degrees of freedom, SS: sum of squares, MS: mean of squares.

lf	SS	MS	F	р
)	118,199.2	13,133.2	9.91	0.00000
45	192,110.8	1324.9		
)	89,219.0	9913.2	6.50	0.00000
45	221,091.0	1524.8		
)	47,319.2	5257.7	2.90	0.00350
45	262,990.8	1813.7		
)	61,267.8	6807.5	3.96	0.00015
45	249,042.2	1717.5		
)	164,841.0	18,315.7	18.26	0.00000
45	145,469.0	1003.2		
	f 45 45 45 45 45	f SS 118,199.2 192,110.8 89,219.0 45 221,091.0 45 2262,990.8 61,267.8 45 249,042.2 164,841.0 45 145,469.0	f SS MS 118,199.2 13,133.2 192,110.8 1324.9 45 192,110.8 45 221,091.0 45 221,091.0 45 262,990.8 45 249,042.2 45 249,042.2 45 164,841.0 18,315.7 45 145,469.0	f SS MS F 118,199.2 13,133.2 9.91 45 192,110.8 1324.9 9.91 45 221,091.0 9913.2 6.50 45 221,091.0 1524.8 6.50 45 262,990.8 1813.7 2.90 45 61,267.8 6807.5 3.96 45 249,042.2 1717.5 18.26 45 164,841.0 18,315.7 18.26

3.4. Stable carbon isotopes

 C_3 plants are more depleted in ¹³C relative to C_4 plants (Vogel, 1978; O'Leary, 1981). Therefore, tooth enamel with higher carbon isotope values is indicative of the consumption of more C_4 plants. We assign each fossil enamel carbon isotope value to the dietary classifications of Lee-Thorp et al. (2007); see Tables 10, 11 & 12.

In line with results from the other dietary proxies, Table 11 shows *Antidorcas bondi* was generally eating a very different diet to other *Antidorcas* species and is more in line with *Damaliscus* (grazing) diets (other than during Swartkrans Member 2 times c.1.65–1.07 Ma). The higher mean carbon isotope values of *A. bondi* are indicative of C₄ plants being consumed, suggestive of a grazing and mixed feeding diet (Cerling and Harris, 1999). Of the 19 fossils classified as *Antidorcas* sp. (Table 8), two specimens (SK 14123 and SK11073) have carbon isotopic compositions consistent with *A. marsupialis*, and can be tentatively assigned to this species; the remaining specimens have higher carbon isotope values and are likely to belong to *A. bondi*.

The blesbok, *Damaliscus pygargus (dorcas)* is typically considered to be a grazing species and indeed, *Damaliscus* is expected to largely be a grazing genus (Estes, 1991). At Sterkfontein and Swartkrans, isotope data shows *Damaliscus* sp. to be true to expected dietary preferences and a small sample is included here to establish most-likely grazing parameters.

Stable carbon isotope results (Tables 11 & 12) support the dietary categories and vegetation signals obtained from use-wear (microwear and mesowear) analyses. However, *Antidorcas bondi* shows slight indications of more mixed feeding in Swartkrans Member 2, away from the specialized grazing signals apparent from the other methodologies. This perhaps lends support to the presence of more heterogeneous habitats for Swartkrans Member 2.

4. Discussion

This paper has used data from various dietary proxies of an abundant antelope to attempt a detailed reconstruction of the palaeovegetation conditions through time at the important hominin-bearing sites of the Cradle of Humankind. Each proxy, predictably, reflects a different aspect of the diets of the springboks. Combined, we have tried to use these datasets to address two main questions: 1) can palaeodietary

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information be used to reconstruct habitat conditions through time; and 2) what additional detail can we gain about the dietary differences between the different Antidorcas species in the fossil record? In response to the first question, the dietary proxies from the Antidorcas examined in this study do yield palaeovegetation reconstructions that broadly agree with habitat reconstructions based on other palaeoenvironmental proxies. Our results suggest increased grassland post c.1.7 Ma, with some suggestion of more heterogeneous habitats for Swartkrans Member 2 (c.1.65-1.07 Ma). A reduced grassland presence in Swartkrans Member 2 is potentially indicated with the assumed-to-be grazing Antidorcas species, A. bondi, yielding a slightly mixed feeding dietary signal from stable carbon isotope values. Whilst it is only slightly indicative of more mixed feeding practices. Antidorcas bondi specimens from Swartkrans Member 2 do exhibit dietary behavioral difference when compared to A. bondi from other members, in which A. bondi is shown to be a more faithful grazer (from stable carbon isotope values). Lee-Thorp et al. (2007) suggested the lower δ^{13} C values were indicative of either a remnant component of C₃ vegetation or less specialized dietary taxa than their modern counterparts. However, dental microwear textures studied here suggest A. bondi had a more mixedfeeding diet than previously assumed.

The second question regarding dietary niches in the various fossil species of *Antidorcas* is more complex. Our dietary reconstructions indicate a wide dietary range represented within the *Antidorcas* lineage, as we would expect from a mixed-feeder that survived changing habitat and vegetation conditions.

In contrast to Brink and Lee-Thorp (1992) all the fossil species of Antidorcas display high values in dental microwear texture complexity (Asfc) which exclude grazing habitats. DMTA supports that all species of Antidorcas were mixed feeding, including herbaceous monocots and browse (shrubs and foliages with lignified tissues, possibly with seeds and fruit). A. recki displays a lower epLsar than the other Antidorcas species (see Table 5), and significantly lower than the fossil A. marsupialis (Table 7). The dental microwear texture for fossil A. bondi here shows it to have practiced greater mixed feeding than shown by the fossil A. marsupialis sample (Fig. 5). Dental microwear textures on A. recki are consistent with mesowear and isotopic analysis in all supporting a higher browsing component in the diet compared to the other Antidorcas species. Browsing is more likely to result in higher heterogeneity due to the nature and range of the browse vegetation consumed. Interestingly, both fossil A. marsupialis/australis and A. recki have higher heterogeneity of diet, indicative of more mixed feeding diets, higher than that of modern Antidorcas marsupialis, a known mixed-feeder. Indeed, all DMTA supports Antidorcas being a predominantly mixed-feeding genus.

In interpreting these diets and what they mean in the wider context of habitat changes and the faunal community, we must also consider the role of biotic factors within the faunal community. Codron et al. (2008) highlight the role that niche specialization can play on changing diets of herbivores within a community, with both environmental and biotic (competition) factors contributing to the forcing of dietary differentiation into specific niches (C_3 or C_4). *Antidorcas* may have been in competition with other herbivore species such as the impala (*Aepyceros melampus*).

Significant differences in tooth metrics analysed in this study indicate that the ancestral *A. recki* possessed smaller overall dental dimensions than modern *A. marsupialis*, supporting the results from previous studies (e.g. Vrba, 1970, 1973) and strongly suggesting that *A. recki* possessed a smaller body size (Gentry, 1978) than *A. marsupialis*. Body size can be informative of palaeoenvironmental conditions, and increases in body size can possibly be explained as an advantage in more seasonal environments where there is a longer period of fasting, a phenomenon known as fasting endurance (Millar and Hickling, 1990;

 Table 7

 Pairwise differences for microwear texture parameters between modern ruminants and extinct species of Antidorcas from South African fossil sites. Significance at p < 0.05 for a given variable is indicated by variable abbreviations below the diagonal for Fisher's LSD tests and above the diagonal for both Tukey's HSD and Fisher's LSD tests (see text for details). Antidorcas sp. represents specimens identifiable only to genus. Fossil 'Antidorcas marsupialis' represents Antidorcas marsupialis' represents Antidorcas marsupialis' represents Antidorcas marsupialis'
 au

LSD \ HSD tests		Antidorcas bondi	Antidorcas sp	Antidorcas marsupialis	Antidorcas recki	Alcelaphus buselaphus	Cephalophus sylvicultor	Giraffa camelopardalis	Syncerus caffer	Damaliscus pygargus	Antidorcas marsupialis
	Antidorcas bondi					asfc		asfc	asfc		
Fossil	Antidorcas sp					asfc		asfc	asfc		
	Antidorcas marsupialis					asfc			asfc		
	Antidorcas recki					asfc			asfc		
	Alcelaphus buselaphus	asfc	asfc	asfc	asfc		asfc				asfc
	Cephalophus sylvicultor					asfc			asfc		
Modem	Giraffa camelopardalis	asfc	asfc	asfc	asfc		asfc				
	Syncerus caffer	astc	asfc	asfc	asfc		asfc				asfc
	Damaliscus pygargus	astc				astc			astc		
	Antidorcas marsupialis	asic				asrc			asrc		
	Antidorcas bondi							epLsar			
Fossil	Antidorcas sp							epLsar			
	Antidorcas marsupialis						epLsar	epLsar			
	Antidorcas recki			epLsar							
	Alcelaphus buselaphus						epLsar	epLsar			
	Cephalophus sylvicultor	epLsar	epLsar	epLsar		epLsar					epLsar
Modem	Giraffa camelopardalis	epLsar	epLsar	epLsar		epLsar			epLsar		epLsar
	Syncerus caffer						epLsar	epLsar			
	Damaliscus pygargus							epLsar			
	Antidorcas marsupialis				epLsar		epLsar	epLsar	epLsar	epLsar	
	Antidorcas bondi										
Fossil	Antidorcas sp								HAsfc 9		
	Antidorcas marsupialis										
	Antidorcas recki		1								
	Alcelaphus buselaphus		HAsfc 9								
	Cephalophus sylvicultor		HAsfc 9								
Modem	Girarra camelopardalis	0.3-01-	HASTC 9								
	Syncerus carrer	HASIC 9	HASIC 9			0 -J- VII		0 -7 - 0	0 -7- 0		
	Damaiscus pygargus					HASIC 9		HASIC 9			
	Antidorcas marsupialis					HASIC 9		HASIC 9	HASIC 9		
	Antidorcas bondi										
Fossil	Antidorcas sp										
	Antidorcas marsupialis								19.01		
	Antidorcas recki		10						HAStc 81		
	Alcelaphus buselaphus		HASIC 81		HASTC 81						HASTC 81
	Giraffa camelonardalis		HASE 81		HASIC 01 HASIC 81						
Modem	Syncerus caffer	- HAsfc 81	HAsfc 81		HAsfc 81						HAsfc 81
	Damaliscus pygargus					HAsfc 81	HAsfc 81	HAsfc 81	HAsfc 81		
	Antidorcas marsupialis					HAsfc 81	HAsfc 81	HAsfc 81	HAsfc 81		
	Antidorcas bondi	-				TfA2	Tf\2	Tf/2	Tfv2		
1000	Antidorcas sp					Tf\2	Tfv2	TfA2	Tfv2		
	Antidorcas marsupialis					Tfv2	Tfv2	TfA2	Tfv2		
	Antidorcas recki		Tfv2					Tfv2			
	Alcelaphus buselaphus	Tfv2	Tfv2	Tfv2			Tfv2	Tfv2			TfA2
	Cephalophus sylvicultor	Tfv2	Tfv2	Tfv2	Tfv2	Tfv2				Tfv2	Tf/2
Modem	Giraffa camelopardalis	Tfv2	Tf/2	Tf/2	TfV2	Tf/2	C it	¢ F	TfA2	Tfv2	Tf/2
	Syncerus catter	142	142	142		0.54	11/2	1772	¢	241	11/7
	Damaliscus pygargus					11V2	1142	11/2	11/2		
	Antidorcas marsupiaris					INZ	7/11	IN2	7/11		

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Table 8

Dental microwear textural parameters (mean and standard deviation). All Antidorcas species are pooled together, with fossil Antidorcas separated by provenance (site and Member 'M'), where sufficient sample sizes allowed, to indicate the prevailing trend through time. 'Modern' represents the modern Antidorcas marsupialis sample.

Provenance	Ν	Asfc			epLsar			HAsfc	9		HAsfc	81		TFV2		
		М	SD	SEM	М	SD	SEM	М	SD	SEM	М	SD	SEM	М	SD	SEM
Sterkfontein M4	5	7.51	4.25	1.90	0.00368	0.00250	0.00112	0.52	0.27	0.22	1.06	0.49	0.22	56,845.1	16,825.5	7524.6
Swartkrans M2	12	1.50	3.84	1.11	0.00589	0.00238	0.00069	0.50	0.21	0.06	0.80	0.26	0.07	69,556.8	11,937.3	3446.0
Swartkrans M3	7	2.80	1.19	0.45	0.00697	0.00336	0.00127	0.41	0.17	0.06	0.70	0.23	0.09	59,850.9	12,984.3	4907.6
Cave of Hearths	10	6.17	4.28	1.35	0.00482	0.00338	0.00107	0.59	0.43	0.13	0.93	0.53	0.17	65,679.1	9409.6	2975.6
Modern	16	3.32	2.40	0.60	0.00733	0.00360	0.00090	0.53	0.32	0.08	1.02	0.64	0.16	63,222.4	18,965.2	4741.3

Table 9

One-way ANOVAs on dental microwear textural parameters to explore variations between Members (Sterkfontein Member 4, Swartkrans Member 2, Swartkrans Member 3, Cave of Hearths and Modern springbok). df: degrees of freedom, SS: sum of squares, MS: mean of squares.

	df	SS	MS	F	р
Asfc					
Sample	4	15,620.9	3905.2	1.42	0.242
Error	45	123,612.4	2746.9		
epLsar					
Sample	4	27,589.5	6897.4	2.02	0.107
Error	45	153,388.5	3408.6		
HAsfc 9					
Sample	4	1952.6	488.2	0.15	0.964
Error	45	151,665.9	3370.4		
HAsfc 81					
Sample	4	7396.2	1849.1	0.75	0.563
Error	45	110,802.2	2462.3		
Tfv2					
Sample	4	13,843.6	3460.9	1.64	0.181
error	45	94,932.1	2109.6		

Table 10

Dietary categories according to Stable carbon isotope values. Dietary distinctions according to δ^{13} C values and proportion of C3/C4 vegetation dominance in diet. Values from Lee-Thorp et al. (2007).

Dietary category	C3/C4 dominance	$\delta^{13}C$
Grazer	C ₄	> -3.0%
Mixed feeder	C ₃ & C ₄	< -3.0% > -9.0‰
Browser	C ₃	< -9.0%

Reynolds, 2007).

One of the outstanding problems in the fossil record is whether the extinct, Antidorcas australis is a valid species or not. The original species description emphasizes the difference in horncore shape, with A. australis having somewhat straighter horns than the lyrate form typical of A. marsupialis (Cooke and Wells, 1951). At present, we do not have enough data to definitively identify clear dietary distinctions between fossils assigned to A. australis and those assigned to A. marsupialis. The sample of fossil Antidorcas marsupialis/australis examined in this study does exhibit greater variation in measurements than modern Antidorcas marsupialis, which could be argued to lend support to the idea of the presence of multiple species (Brophy et al., 2014). Further landmarkbased measurements and additional techniques, such as geometric morphometric analysis (to analyse shape changes) would be required to settle the question of this species distinction. It may be that the A. australis is differentiated only on its horncores and that dietary differences alone would not be sufficient to indicate distinctions between A. australis and A. marsupialis.

Another important aspect of modern springbok ecology is the migratory behaviour. Historic records show that springbok would trek in herds of thousands of individuals to seek out fresh grazing after the rains (Skinner and Louw, 1996). At present, there is little way to determine whether other extinct species of *Antidorcas* exhibited this behaviour or not. If so, *A. australis*, may have been a migrant to the Cradle region at various times. Certainly, migration would have helped reduce competition in grazing species.

As may be expected (e.g. Davis and Pineda-Munoz, 2016), there is some disparity between dietary signals from DMTA compared to those from stable carbon isotopes. DMTA indicates all *Antidorcas* fossil species were mixed feeders, whereas carbon isotopes indicate higher C4 signals (i.e. grazing) for these antelopes, particularly the case for *A. bondi*. This combination would support a mixed feeding dietary preference for *Antidorcas*, altering their dietary intake according to seasonal availability of resources. This suggests that the Cradle of Humankind region would have had seasonally available resources with significant amounts of both browse and graze. If severe seasonal droughts decimated the herbaceous layer (where grasses are abundant) then the antelopes may have switched to browse as tree/shrub roots go deeper in the soil to obtain water, compared to the herbaceous plants.

5. Summary

In this paper, we examined fossil *Antidorcas* dentition using a range of dietary proxies, from these results we attempted to infer the prevailing vegetation conditions between 2.8 and 0.8 Ma for the region where these *Antidorcas* lived and died, the Cradle of Humankind, South Africa. The majority of past research has used a single methodology on multiple species from assemblages, to address palaeoenvironmental questions. Here, we implemented a multiproxy analysis on a single mixed-feeding, herbivorous and abundant genus (*Antidorcas*) to provide a detailed palaeovegetation picture for the Sterkfontein and Swartkrans area through time. Aside from some subtle variations, preliminary conclusions from our study are found to be in broad agreement with those found through isotopic analysis by Lee-Thorp et al. (2007), of increased grassland presence in South Africa from ca 0.1.7 Ma.

Notably, we have been able to suggest from this multi-method approach, that methods can be reliably substituted for one another where time or cost constraints apply. Where methods such as mesowear, are non-destructive, quick and cheap to implement, these techniques can be used to accurately pinpoint where to concentrate the more time-consuming, expensive and destructive techniques, such as stable carbon isotope analysis. However, a more holistic, robust and detailed palaeovegetation reconstruction can be achieved by using complementary methodologies. The complementary use of methods such as mesowear analysis, stable isotope analysis and dental microwear texture analyses for instance, allows us to question whether *Antidorcas bondi* was typically a grazer or not. This approach could be implemented at other sites across the world for palaeovegetation reconstructions, where the

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Table 11

Stable carbon and oxygen isotope data for Antidorcas fossils from Swartkrans. Antidorcas sp. indicates specimens only identifiable to genus level (specimens processed at the University of Liverpool).

Number	Species	δ^{13} C	δ ¹⁸ Ο	Member
SK5882	Antidorcas sp.	1.6	-1.5	Swartkrans Member 2
SK5882	Antidorcas sp.	1.8	-0.8	Swartkrans Member 2
SK5990	Antidorcas sp.	- 4.8	1.6	Swartkrans Member 2
SK10555	Antidorcas sp.	-1.1	0.1	Swartkrans Member 2
SK10555	Antidorcas sp.	-1.6	0.2	Swartkrans Member 2
SK4064	Antidorcas sp.	-0.6	0.6	Swartkrans Member 2
SK6118	Antidorcas sp.	1.1	0.6	Swartkrans Member 2
SK2264	Antidorcas sp.	-3.8	-0.9	Swartkrans Member 2
SK4083	Antidorcas sp.	-0.4	1.8	Swartkrans Member 2
SK4083	Antidorcas sp.	-0.4	1.5	Swartkrans Member 2
SK4080	Antidorcas sp.	-1.7	0.1	Swartkrans Member 2
SK2292	Antidorcas sp.	-5.9	-1.3	Swartkrans Member 2
SK4633	Antidorcas sp.	-0.6	-1.1	Swartkrans Member 2
SK11899	Antidorcas sp.	2.1	-0.8	Swartkrans Member 2
SK6106	Antidorcas sp.	-0.2	0.3	Swartkrans Member 2
SK11073	Antidorcas sp.	-12.7	-1.3	Swartkrans Member 2
SK 3014	Antidorcas sp.	-1.1	-0.8	Swartkrans Member 2
SK 14123	Antidorcas sp.	-12.7	0.8	Swartkrans Member 2
SK14070	Antidorcas marsupialis	-12.9	-1.9	Swartkrans Member 2
SK4054	Antidorcas marsupialis	-11.6	0.2	Swartkrans Member 2
SK 5958	Antidorcas marsupialis	-9.1	-0.8	Swartkrans Member 2
SK 4081	Antidorcas marsupialis	-10.3	0.4	Swartkrans Member 2
SK 2953	Antidorcas marsupialis	-15.0	-2.1	Swartkrans Member 2
SK 3055	Antidorcas marsupialis/australis	-11.0	2.1	Swartkrans Member 2
SK 2366	Antidorcas bondi	-5.2	-2.2	Swartkrans Member 2
SK 40592	Antidorcas sp.	-2.2	0.4	Swartkrans Member 5
SKX 41059	Antidorcas australis/marsupialis	-7.3	-1.1	Swartkrans Member 5

Table 12

Stable carbon isotope descriptive statistics for *Antidorcas and Damaliscus* species present in each member. References: where more than one value is given in original reference, a mean value has been calculated and used here. Luyt, 2001¹; Van der Merwe et al., 2003²; Lee-Thorp et al., 2000³; Lee-Thorp et al., 1994⁴; New isotope data (this study)⁵; Vogel, 1978⁶; Sponheimer et al., 2003⁷. Dietary categories following category distinctions as set out in Table 6. *N.B. Modern *Antidorcas marsupialis* would be expected to be a mixed feeder. To aid comparison between the modern and fossil data, modern enamel values were increased by 1‰ to account for the post-industrial decrease in the δ^{13} C value of atmospheric CO₂ (Francey et al., 1999).

Species	Ν	$Mean \; \delta^{13}C$	Standard deviation	$Minimum \; \delta^{13}C$	Maximum $\delta^{13}C$	Dietary category
Sterkfontein Member 4						
Antidorcas recki	10 1,2	-11.4	1.063	-14.0	-5.0	Browser
Antidorcas bondi	2 ^{1,2}	-1.3	0.020	-1.0	-1.0	Grazer
Damaliscus sp.	9 ¹	-1.8	1.256	-7	3.0	Grazer
Sterkfontein Member 5						
Antidorcas recki	2 1	-11.7	0.945	-13.0	-11.0	Browser
Damaliscus sp.	5 ¹	-1.3	1.055	-5.0	1.0	Grazer
Antidorcas sp.	3 1	-5.4	3.133	-10.0	1.0	Mixed
Sterkfontein Member 5 East/Oldowan						
Damaliscus sp.	5 ¹	-2.7	0.965	-5.0	1.0	Grazer
Antidorcas sp.	1 ¹	-9.2	x	-9.2	-9.2	Browser-Mixed
Sterkfontein Member 5 West/Acheulean						
Antidorcas recki	2 1	-11.7	0.945	-13.0	-11.0	Browser
Damaliscus sp.	7 1	-0.9	0.540	-3.0	1.0	Grazer
Swartkrans Member 2						
Antidorcas recki	1 4	-12.9	x	-12.9	-12.9	Browser
Antidorcas bondi	9 ³	-3.3	0.334	-5.0	-2.0	Mixed-Grazer
Antidorcas sp.	27 ^{4,5}	-5.1	1.076	-15.0	2.0	Mixed
Modern						
Antidorcas marsupialis	2 ^{6,7}	-13.1*	4.00	-17.0*	-9.0*	Browser

preserved faunal community is restricted in some way but where at least one continuous (mixed feeding) herbivore lineage is represented.

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