Faunal Assemblage Seriation of Southern African Pliocene and Pleistocene Fossil Deposits

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ABSTRACT Fossil assemblages from the Pliocene and Pleistocene of southern Africa were seriated in order to give a better idea of their relative chronology. Time-sensitive mammals were selected for calculation of the Faunal Resemblance Index among 17 site units. On the basis of a logistical seriation and subsequent site analysis, the following sequence of sites was deemed most probable: Makapansgat Member 3, Makapansgat Member 4, Taung Dart deposits, Sterkfontein Member 4 and Taung Hrdlička deposits, Sterkfontein Member 5 (in part) and Kromdraai B, Kromdraai A and Swartkrans Member 1, Swartkrans Member 2, Swartkrans Member 3, Plovers Lake, Cornelia, Elandsfontein Main Site, Cave of Hearths Acheulian levels, Florisbad and Equus Cave and Klasies River Mouth. © 1995 Wiley-Liss, Inc.

Associations of the early hominid fossils with a wealth of Pliocene and Pleistocene fauna from southern Africa provide vital clues to the past environments and evolutionary processes that shaped human evolution. Due to the nature of the cave deposits in which these ancient assemblages are typically found, where absolute geochronological dating techniques are inhibited, the absolute and even relative temporal positions of these sites have proved difficult to ascertain. Nevertheless, it is of pivotal importance in studies of mammalian evolution, and particularly in paleoanthropological research, to establish a chronology with which we can assess the evolutionary roles of the varied fauna found in southern African cave deposits.

Understanding the timing of deposition in southern African hominid fossil sites is of continuing importance, as illustrated by the influence dates have on interpretations of hominid phylogeny and theories of the causes of evolution. For example, the proposed late date for the type specimen of *Australopithecus africanus* from Taung, at circa 1 mya (Partridge, 1973, 1982, 1985; Vogel and Partridge, 1983; Vogel, 1985) led to the suggestion that the Taung child may have been close to the hypodigm of *A. robustus* (Tobias, 1973, 1975, 1978; Wolpoff, 1974) or that it represented a derived form of *A. afri*canus (Tobias, 1991a,b). Recent faunal dating of Taung (Delson, 1988; Cooke, 1990; McKee, 1993a), placing the site in time closer to Sterkfontein and Makapansgat, opens the possibility that the Taung hominid is a more typical or even primitive representative of *A. africanus*.

Turner (1983, 1990) and Vrba (1985a, 1988, 1992) have argued that changes in global climate during the Pliocene *caused* mammalian evolutionary changes in Africa. Recently Hill et al. (1992) have suggested that the origin of *Homo* corresponded to these putative climatic changes on the basis of the age of a specimen they feel represents the genus. However, in southern Africa it is

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not clear whether the apparent faunal turnovers seen in the fossil record predate or postdate any climatic change, or whether or not some of the faunal turnovers are indeed real (McKee, 1991, in press). Thus an understanding of cause and effect in mammalian evolution requires at least an understanding of the succession of sites if not the actual age of deposition.

Opportunities to determine the ages of sites by geological dating have been limited by the nature of cave deposits. Sedimentological correlations of sites (Butzer, 1974), nick-point analysis of cave openings (Partridge, 1973), sedimentological rates (Partridge, 1985), and isotopic techniques (Vogel and Partridge, 1983; Vogel, 1985) have provided unsatisfactory results (De Swardt, 1974; Butzer et al., 1978; Tobias et al., 1993; McKee, 1993a). Palaeomagnetic chronologies have given mixed results, with some success at Makapansgat (Brock et al., 1977; McFadden et al., 1979; McFadden, 1980) and Kromdraai (Jones et al., 1986), but few clues at other sites due to the dynamics of cave sediments. Electron spin resonance (ESR) has provided age estimates for various sites, including Border Cave (Grün et al., 1990a), Klasies River Mouth (Grün et al., 1990b), and Sterkfontein (Schwarcz et al., 1993), but these are dependent on assumptions regarding rates of uptake of uranium (Grün and Stringer, 1991). Nevertheless, there is considerable potential in the application of ESR dating with necessary caveats and corroborating evidence from other techniques.

Chronologies based on faunal correlations have thus been the primary means of dating southern African fossil sites. Many attempts have been made (reviewed by McKee, 1993a), but the most relevant work has its cornerstone in the faunal spans delineated by Ewer and Cooke (1964) and Cooke (1967). The oldest of these temporal associations was the Sterkfontein Faunal Span, comprising the *Australopithecus*-bearing deposits of Sterkfontein, Makapansgat, and Taung, as well as the Older Gravels of the Vaal River Basin. The subsequent Swartkrans Faunal Span encompassed deposits from the Sterkfontein extension site (most of which is now referred to as Sterkfontein Member 5), Swartkrans, and Kromdraai. The Vaal-Cornelia Faunal Span included the Vaal Younger gravels, Elandsfontein (formerly known as Hopefield or Saldanha), and Cornelia, followed by the Florisbad-Vlakkraal Faunal Span encompassing the important *Homo*-bearing sites of Florisbad and Broken Hill.

As progress is made in both excavation and taxonomic identification of southern African fossil fauna, sufficient samples are emerging for the refinement of the early methods that simply recognized faunal spans. Toward this end, biochronologies within taxonomic families have been established for bovids (Vrba, 1975, 1982, 1985b), cercopithecids (Delson, 1984, 1988), carnivores (Hendey, 1974; Turner, 1990), and suids (Harris and White, 1979). A combined picture of all the known mammalian fauna may then be useful in the seriation of individual site units within and between the faunal spans.

There are numerous factors that may lead to similarities or differences of faunal representation in site assemblages that must be considered when attempting to establish biochronologies across taxonomic boundaries. Temporal determinants, those most useful for faunal dating, interplay with other sampling biases so that only a fraction of the total faunal community living at any one time would be represented in a cave deposit. A summary of the key factors that may lead to similarities or differences among faunal assemblages in southern African cave deposits can be found in Table 1.

Although faunal dating may have its limitations due to the complex network of factors that determine site assemblages, it remains as the most reliable tool for determining the sequence of fossil sites in southern Africa. Our objective here is to establish a chronological framework for understanding mammalian evolution in the Pliocene and Pleistocene of southern Africa through the seriation of site units. In order to achieve this with a total mammalian perspective, issues arising from the complicating factors discussed above must be resolved with an appropriate method.

TABLE 1. Factors	causing similarities	s or differences in
faunal represente	ition at southern Af	rican cave sites

Environmental factors
Faunal community
Floral community
Demography
Climate
Altitude
Terrain
Geographic proximity of sites
Availability of water
Cave size and depth
Taphonomic factors
Taphonomic agents
Bone preservation
Mixing of deposits
Interpretive factors
Selective excavation and preparation
Taxonomic biases
Temporal factors
Origins
Radiations
Extinctions

METHODS AND ASSUMPTIONS

Lists of identified mammalian species and genera were compiled from published literature for 23 Neogene site units (Fig. 1 and Table 2). This amounted to 246 species of 159 genera and 44 families.

It was our primary goal to seriate the sites based on similarities of identified species between site unit assemblages. The initial step is to create a matrix exhibiting the number of identified species shared between each pair of site units (Table 3A); the diagonal from top left to bottom right shows the number of identified species for each site unit. Specific identifications with "cf." or "aff." were included for the analysis, as in Turner and Wood (1993).

The ideally seriated matrix should exhibit an ordering of figures within each row, increasing from left to right toward the diagonal, and decreasing away from the diagonal. In other words, sites closer in time should share a successively greater number of species than more temporally distant sites. The nontemporal factors listed in Table 1, however, severely complicate the ideal, and additional steps are necessary to place the focus of the seriation on the temporal components of the data.

Intrinsic standardization of the data is necessary in order to alleviate the problems engendered by differential sampling; a well

sampled or faunally diverse site is more likely to yield species identified from elsewhere than is a site with a paucity of fauna. Simpson's (1960) Faunal Resemblance Index (FRI) provides a partial solution to the sampling problem and is employed here. For each pair of sites, the FRI is calculated by taking the number of species shared (C) divided by the total number of species found in the smaller sample (N_1) , or simply FRI = C/ $N_1 \times 100$. These values are displayed in Table 3B. The advantage of the method is that a small sample can exhibit an FRI value comparable to that of a much larger sample. thus standardizing the figures in the matrix. It should be noted, however, that a site with a large sample still has a greater probability of sharing species with other sites and the use of an FRI does not address this issue.

Complications from nontemporal factors of site similarities can cause spurious correlations. For example, a long-lived but ecologically limited species, *Hippopotamus amphibius*, appears at the oldest and youngest sites, but not at sites that span the time in between. Thus the analysis must be limited to the species that are most sensitive to time and not restricted to highly specialized environments.

Evaluation of temporally sensitive species can be a somewhat subjective task. Any criteria used to label a species as time-sensitive must be seen as an underlying assumption of the analysis. For this analysis we established the following objective criteria for identifying such species: (1) The species must occur at more than one fossil site; species known from only one site that show evolutionary relationships to species from other sites can be used as corroborating evidence. (2) The species must not be present throughout the entirety of the sequence (given the assumption of an oldest site). (3) Species that are highly sensitive to local environmental conditions may cause spurious correlations among sites of similar conditions from vastly different times, and thus should be excluded. Known habitat preferences of extant species can be used to infer similar tendencies of the same or related species in the past, but the potential for bias in exclud-

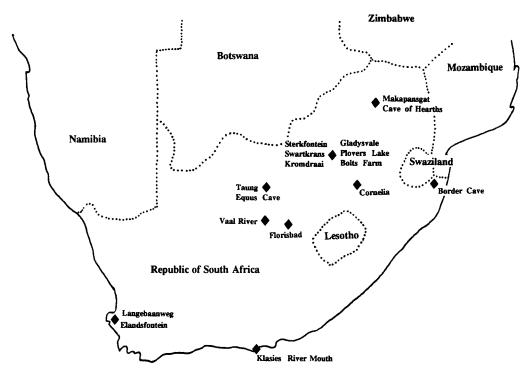


Fig. 1. Map of southern African fossil sites.

ing species on this basis must be acknowledged. Thus species that occur at sites which, on the basis of independent evidence, appear to have been environmentally distinct (e.g., Taung and Makapansgat), were not excluded. It should be remembered that most of the southern African fossil sites are cave deposits that have sampled mammals from a mosaic of local environments.

Virtually all of the 66 time-sensitive species identified using the above criteria showed some degree of irregularity in their appearances at various sites, as they are not free from the factors listed in Table 1. It thus must be assumed that sampling of such species from site units is random with respect to time. The time-sensitive species are listed in Table 4 along with their distribution among the sites used in the analysis. The matrices in Table 5 are based on time-sensitive species alone.

Once time-sensitive species were considered, a number of sites had to be excluded from the analysis due to small sample sizes. The small samples of time-sensitive species produced unrealistically large FRI values. These sites included Langebaanweg, the Taung Dart Deposits, the Vaal Older River Gravels, and Border Cave. Additional sites were excluded because it was known a priori that fauna from deposits of vastly different time periods were mixed into single assemblages; these sites were Bolt's Farm and Gladysvale (Berger, 1993). It must be acknowledged at the outset, however, that some mixing of fossils may have occurred at other sites with multiple deposits.

Logistical seriation of the sites was based on the matrix of FRI values for time-sensitive species. A computer program was written which tested different site sequences in order to minimize differences among adjacent sites showing "backward steps": figures that decrease in value toward the diagonal are "backward steps" because in an ideal matrix the FRI values should always increase toward the diagonal. This was accomplished by summing the absolute value of differences between adjacent sites in a row showing "backward steps." The site unit or-

238

TABLE 2. Site units and codes used in Tables $3-5^1$

Site unit	Code
Langebaanweg 'E' Quarry	LANG
Makapansgat Member 3	MAK3
Makapansgat Member 4	MAK4
Taung Dart deposits	TAUD
Taung Hrdlička deposits	TAUH
Sterkfontein Member 4	STS4
Sterkfontein Member 5	STS5
Older Vaal River Gravels	VAAL
Bolt's Farm	BOLT
Kromdraai A	KROA
Kromdraai B	KROB
Gladysvale	GLVH
Swartkrans Member 1	SWT1
Swartkrans Member 2	SWT2
Swartkrans Member 3	SWT3
Ployers Lake	PLOV
Cornelia	CORN
Elandsfontein Main Site	EFTM
Cave of Hearths (Acheulian levels)	COHA
Border Cave (MSA levels)	BCVM
Florisbad springs	FLOS
Equus Cave (MSA levels)	EQUU
Klasies River Mouth (MSA I and II levels)	KLAS
Extant species in southern Africa	EXTA

¹Species identified from each site were compiled by one of us (JKM) from the following references: Berger (1993), Brain (1981), Brain and Watson (1992), Brickhill (1976), Brink (1987), Butzer et al. (1974), Churcher (1970), Collings (1972), Collings et al. (1976), Cooke (1978, 1988, 1990, 1991, 1993a,b), Cooke and Wells (1947), de Graaf (1960, 1988), Freedman (1957), Greenwood (1955), Helgren (1979), Hendey (1981), Hooijer (1958), Klein (1976, 1977), Klein et al. (1991), Klein and Cruz-Uribe (1991), McKee (1991, 1993a,c), Randall (1975), Pocock (1987), Skinner and Smithers (1990), Szalay and Delson (1979), Turner (1986), Vrba (1981, 1987a,b), Watson (1993). An unpublished species list was provided by J.F. Thackeray for Plovers Lake.

der producing the minimum sum was deemed the best seriation. As the number of possible seriated orders is a factorial of the number of site units, only 10 sites could be seriated at any one time and the complete sequence was compiled from overlapping seriations. For the overall ordering of the sequences, an a priori assumption was made that sites with fewer extant species were earlier in time.

Use of the method outlined above minimizes the effects of nontemporal factors but is far from foolproof. Initial tests of the process revealed that a number of principles must be imposed on the logistcal seriation and interpretation of the matrix. (1) When available, stratigraphic relations among site units must be used. For example, Swartkrans Member 3 must follow Swartkrans Member 2, although this detracts from an ideal sequence of FRI values. (2) Where multiple sequences are possible, each must be treated as tenable. The exclusion of certain sites often changes the sequence among other sites, and these situations must be scrutinized carefully. However, penecontemporaneous sites may be expected to seriate in alternating positions. (3) Each site must be considered in light of corroborating evidence from evolutionary sequences of lineages, human artifacts (stone tools, etc.), and absolute dating, whenever such information is available.

RESULTS

The seriation of the FRI values presented in Table 5B led to the following optimum series of 10 sites per run:

- A. MAK4 MAK3 TAUH STS4 STS5 KROB KROA SWT1 SWT3 SWT2
- B. MAK3 MAK4 STS4 TAUH KROB KROA SWT1 SWT2 SWT3 PLOV

These two sequences maintained consistency with the further removal of individuals sites. The position of the Taung Hrdlička assemblage alternated with Sterkfontein Member 4, depending on the inclusion of Sterkfontein Member 5. As discussed below, this may be because of near contemporaneity of the two alternating sites. Although Makapansgat Member 4 was sometimes positioned before Member 3, and Swartkrans Member 3 sometimes seriated before Member 2, forcing the stratigraphic orders in sequence did not affect the relative positions of the other site units.

In the presumed middle of the site sequence, the seriation results were quite consistent and confirmed the relations suggested by sequences A and B above. For example:

- C. TAUH STS5 KROA SWT1 SWT3 SWT2 PLOV CORN EFTM COHA
- D. TAUH KROB KROA SWT1 SWT2 CORN EFTM COHA EQUU KLAS

Optimal seriations of the more recent sites produced a more variable picture, depending on the sites used for analysis. The main variants are illustrated by the following sequences:

EXTA	101 105 105 105 105 105 105 105 105 105	EXTA	$\begin{smallmatrix} & & & & & & \\ & & & & & & & \\ & & & & $
BCVM	0000-00000000000000-40400000	BCVM	100 00 100 00 100 00 100 00 100 00 100 00
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tdex (F)	822248832966999999468888888888888888888888888888	EQUU	986566688888888888888888888888888888888
lance in FLOS	001040100200000040588849	FLOS	0 0 0 0 1 0 0 0 0 0 0 0 0 0 0 0 0 0 0 0
faunal resemblance index (FRI) ¹ EFTM COHA FLOS FOIIII KI.	201130200000000000000000000000000000000	COHA	$\begin{smallmatrix} & & & & \\ & & & & \\ & & & & \\ & & & & $
faunal E.FTM	84419409000041131212121212409000000	EFTM	° ° ° ° ° ° ° ° ° ° ° ° ° ° ° ° ° ° °
site units: (A) absolute frequencies of shared species and (B) is know know aiven swer, swer, swer, play CORN		CORN	$^{10}_{33}$
pecies (៲ ៲៵៷៷៰៰៷៷៹៷ <mark>ៜ</mark> ៹៝៷៰៰៲៸ៜ៷៹៷៹៷៹៷	PLOV	7 2 2 2 3 3 3 2 8 1 0 4 3 3 3 5 5 5 5 5 3 3 2 5 2 5 5 1 1 3 8 8 5 5 5 5 5 5 5 5 5 5 5 5 5 5 5 5 5
hared s	21 0 0 1 0 2 1 0 0 0 0 0 0 0 0 0 0 0 0 0	SWT3	$\begin{smallmatrix} & 1 \\ & 0 \\ & $
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requend	2 3 5 5 5 5 5 5 5 5 5 5 5 5 5 5 5 5 5 5	SWT1	6 8 8 8 8 8 8 8 8 8 8 8 8 8 8 8 8 8 8 8
solute f	2,8 119512 2552 2552 9 11 12 2 9 15 2 2,8 119512 2552 2552 9 11 12 2 9 15 2 2,8 119512 2 15 2 15 2 15 2 15 2 15 2 15 2	B GLVH	5 2 2 3 2 4 5 5 2 2 2 5 5 5 5 5 5 5 5 5 5 5 5 5
(A) ab KROA	19 3 10 11 2 2 5 5 5 5 5 2 2 7 1 2 5 1 2 5 1 2 5 1 1 2 5 5 5 5 5 5 5 5	KROA	$\begin{smallmatrix} & 6 \\$
e units. KROB	300345066846686712121212121211211211	KROB	833 33 33 33 33 33 33 33 33 33 33 33 33
ican sit STS5	17-2265233866523386652331111111111111111111111111111111111	STS5	$\begin{smallmatrix} & & & & & \\ & & & & & \\ & & & & & \\ & & & & & \\ & & & & & \\ & & & & & & & \\ & & & & & & \\ & & & & & & & \\ & & & & & & & \\ & & & & & & & \\ & & & & & & & \\ & & & & & & & \\ & & & & & & & \\ & & & & & & & \\ & & & & & & & \\ & & & & & & & \\ & & & & & & & \\ & & & & & & & \\ & & & & & & & \\ & & & & & & & \\ & & & & & & & \\ & & & & & & & \\ & & & & & & & \\ & & & & & & & \\ & & & & & & & \\ & & & & & & $
southern African	2 2 2 2 2 2 2 2 2 2 2 2 2 2 2 2 2 2 2	BOLT	°°°°°°°°°°°°°°°°°°°°°°°°°°°°°°°°°°°°°
t south	12 - x x - x x 2 0 0 2 2 2 2 2 2 2 2 2 2 2 2 2 2 2	TAUH	39 ~ 9 9 9 1 2 9 3 3 3 3 3 2 5 0 0 2 2 4 2 3 3 3 4 9 9 9 1 2 9 5 3 3 3 3 3 2 5 0 0 2 2 4 2 3 3 3 3 3 3 5 0 0 0 2 2 4 2 3 3 3 3 3 3 3 3 3 3 3 3 3 3 3 3
vaal	84000000444488488000000	VAAL	2 5 5 5 5 5 5 5 5 5 5 5 5 5 5 5 5 5 5 5
tared sp		S	000 000 000 000 000 000 000 000
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3.	≈;;;;;;;;;;;;;;;;;;;;;;;;;;;;;;;;;;;;	MAK4	1008 1009 110 12 12 12 12 12 12 12 12 12 12 12 12 12
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UNA I	440-c00-0000-000-0000-0000-0000-0000-00	LANG	0 0 0 1 0 4 0 2 0 0 2 0 0 2 0 0 0 0 0 0 0 0 0 0
	LANG MAK4 MAK4 MAK4 MAK4 TSB TSB TSB TSB KR0B BOLT STS5 KR0B SWT1 SWT2 SWT2 SWT2 SWT2 SWT2 SWT2 SWT2 SWT2		LANG MAKE MAKE MAKE TAUD TAUH TAUH TAUH TAUH TAUH TAUH TAUH TAUH

¹Site codes are listed in Table 2.

SOUTHERN AFRICAN FAUNAL ASSEMBLAGE SERIATION

TABLE 4.	Time-sensitive	mammalian	species w	ith site uni	it occurrences	marked by an ast	$erisk^1$
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	К 3	К 4	S 4	U H	S 5	O B	O A	Т 1	Т 2	Т 3	o v	R N	T M	H A	s^{o}	U U	A S	T A
Artiodactyla																		
Alcelaphus buselaphus														*		*	*	*
Antidorcas australis											*		*					
Antidorcas bondi			*			*	*				*			*	*	*		
Antidorcas marsupialis								*	*	*					*	*		*
Antidorcas recki			*		*	*	*	*	*	*		*	*					
Connochaetes gnou												*	*		*	*		*
Damaliscus dorcas															*	*		3
Damaliscus niro												*	*		*			
Hippotragus cookei	*		*															
Hippotragus leucophaeus													*				*	,
Makapania broomi	*		*															
Megalotragus priscus											*	*	*	*	*	*		
Oreotragus major	*	*	*	*	*													
Pelea capreolus							*	*		*	*					*	*	
Pelorovis antiquus												*	*	*	*		*	
Raphicerus campestris								*	*	*						*		;
Redunca arundinum							*	*				*	*	*			*	
Redunca darti	*		*															
Redunca fulvorufula																*	*	
Simatherium kohllarseni	*	*																
Syncerus caffer																*	*	
Taurotragus oryx							*		*	*	*	*	*	*	*	*	*	
Tragelaphus strepsiceros							*	*		*	*		*	*		*	*	
Kolpochoerus paiceae												*	*			·	-	
Metridiochoerus andrewsi									*	*	*		*					
Notochoerus scotti	*	*		*														
															*	*	*	
Phacochoerus aethiopicus						*	*	*	*	*				Ť	*	*	*	
Phacochoerus modestus			*			Ŧ	*	4	*	Ŧ		+						
Potamochoeroides shawi	÷		*															
Carnivora																		
Lycaon pictus													*		*	*		
Vulpes pulcher							*	*										
Dinofelis barlowi	*		*					*										
Homotherium crenatidens	*		*		*		*											
Megantereon cultridens			*		*	*	*	*		*								
Panthera pardus			*	*		*	*	*	*	*	*					*	*	
Chasmaporthetes silberbergi			*		*			*	-									
Chasmaporthetes nitidula			*		*				*	*								
Hyaena brunnea			*				<u>ب</u>	*	*				*					
	*		*				*	Ŧ	~				Ŧ			*	+	
Pachycrocuta bellax					*		*		ىك	*								
Aonyx capensis									-4-				*		*	÷.	* *	
Mellivora capensis													*			1	*	
Atilax paludinosus								*							•	÷	Ť	-
Herpestes ichneumon								Ŧ					*	*			*	
Suricata suricatta									*	*			~			*		
Primates																		
Australopithecus africanus	*	*	*															
Australopithecus robustus						*		*	*	*								
Homo sapiens													*	*	*	*	*	
Cercopithecoides williamsi	*	*	*	*		*		*										
Papio izodi			*	*														
Papio angusticeps						*	*											
Papio robinsoni						*	*	*	*	*	*							
Papio cynocephalus														*		*	*	
Parapapio broomi	*	*	*															
Parapapio whitei	*	*	*															
	*	*	*			*	*	*										
Parapapio jonesi							-	-	*	*								
Theropithecus oswaldi								*	÷	Ŧ			*					
roboscidea																		
Elephas recki			*		*													
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Perissodactyla Equus burchelli					*					*				*	بو	*		
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Equus capensis	ىتە				*		*	*	*	÷	*	*	*	*	*	*		
Hipparion libycum	*						*	*	*	*		*						
Iyracoidea																		
Procavia transvaalensis		*		*			*	*	*	*	*							
Rodentia																		
Mystromys antiquus	*		*	*		*		*										
Proodontomys cookei	*	*	*	*		*	*	*										
Hystrix makapanensis	*						*											
Elephantulus antiquus	*	*	*	*	*	*	*											

¹Site codes are listed in Table 2.

	TABLE 5.		hared tin	ve-sensitiv	e species	at southe	List of shared time-sensitive species at southern African site units: (A) absolute frequencies of shared species and (B) faunal resemblance index (FRI) ¹	ı site uniı	s: (A) abs	olute freq	uencies o	f shared s	pecies and	l (B) faun	al resemb	lance inde	$x (FRI)^1$	
	MAK3	MAK4	STS4	TAUH	STS5	KROB	KROA	1.TWS	A SWT2	SWT3	PLOV	CORN	EFTM	COHA	FLOS	EQUU	KLAS	EXTA
11110	5		5	9	-	U	0	C	- 	-	-	-	<	9		<		
CALM I	17	3	1		# (. -			4 •	, ,	- 0	•		•	•	> <	
MAK4	10	11	x	¢	N	4	4	4	-	-	1	•	5	•	0	•	0	0
STS4	17	œ	26	7	6	ი	11	12	4	5 2	7	-	7	-		က	5	0
TAUH	9	9	7	6	5	5 D	4	5	0	2	2	0	0	0	0	1	1	1
STS5	4	5	6	67	Π	e	7	ų	က	5	1	7	2	2	2	2	0	1
KROB	5	4	6	л С	က	13	10	10	5	9	က	7	٦	1	Н	2	1	Г
KROA	00	4	11	4	7	10	24	16	6	13	8	9	9	9	4	80	9	7
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	MAK3	MAK4	STS4	TAUH	STS5	KROB	KROA	SWT1	SWT2	SWT3	PLOV	CORN	EFTM	COHA	FLOS	EQUU	KLAS	EXTA
MAK3	100	16	81	67	36	38	38	33	9	5	0	6	0	0	0	0	0	0
MAK4	- 16	100	73	67	18	36	36	36	6	6	6	0	0	0	0	0	0	0
STS4	81	73	100	78	82	69	46	46	24	24	18	6	10	80	2	12	Π	8
TAUH	67	67	78	100	22	56	44	56	22	22	22	0	0	0	0	11	11	11
STS5	36	18	82	22	100	27	64	45	27	45	6	18	18	18	18	18	0	6
KROB	38	36	69	56	27	100	77	77	38	46	27	18	80	80	æ	15	æ	8
KROA	38	36	46	44	64	77	100	67	53	62	73	55	29	46	27	33	33	29
SWT1	33	36	46	56	45	77	67	100	82	81	55	45	38	31	20	36	39	38
SWT2	9	6	24	22	27	38	53	82	100	100	55	45	47	15	27	47	24	41
SWT3	ъ	6	24	22	45	46	62	81	100	100	73	45	43	31	33	52	33	48
PLOV	0	6	18	22	6	27	73	55	55	73	100	27	55	45	36	64	36	36
CORN	6	0	6	0	18	18	55	45	45	45	27	100	82	45	55	36	27	27
EFTM	0	0	10	0	18	8	29	38	47	43	55	82	100	62	60	57	56	57
COHA	0	0	80	0	18	80	46	31	15	31	45	45	62	100	62	85	69	69
FLOS	0	0	7	0	18	8	27	20	27	33	36	55	60	62	100	87	40	67
EQUU	0	0	12	11	18	15	33	36	47	52	64	36	57	85	87	100	83	92
KLAS	0	0	Π	11	0	œ	33	39	24	33	36	27	56	69	40	83	100	94
EXTA	0	0	œ	11	6	æ	29	38	41	48	36	27	57	69	67	92	94	100
¹ Site code	Site codes are listed in Table 2	in Table 2.																

- E. STS5 KROA SWT1 SWT2 PLOV CORN EFTM FLOS EQUU KLAS
- F. SWT2 SWT1 KROA PLOV CORN EFTM FLOS COHA EQUU KLAS
- G. SWT1 SWT2 SWT3 PLOV KROA CORN EFTM EQUU COHA FLOS

Because of the high FRI value for Plovers Lake and Kromdraai A, two of the sequences place the latter site after Swartkrans. But note in sequence F that the first three sites are in reverse order from that expected; they seriated correctly among themselves in order to allow for the similarities between Kromdraai A and Plovers Lake.

None of the Middle Stone Age (MSA) sites (Florisbad, Equus Cave, and Klasies) takes consistent positions in the seriation. A close look at the species from the late Pleistocene sites shows why: there are few distinctive faunal trends following the time of Elandsfontein deposition, thus site correlations are somewhat more spurious. Nevertheless, there is consistency in the sequence of Plovers Lake, Cornelia, Elandsfontein, and Cave of Hearths, and one of the MSA sites always comes out as the most recent site. This is illustrated by the following sequence covering the entire time span:

H. MAK3 TAUH STS4 KROB SWT1 PLOV CORN EFTM COHA FLOS

Site analysis

Makapansgat

Makapansgat Member 3 has long been thought to be the oldest of the southern African fossil sites, and this is borne out by the faunal seriation. Over two-thirds of its species are extinct, and it shares no time-sensitive species with the late Pleistocene Homobearing sites. The assemblage includes particularly ancient species from East Africa such as the chalicotheriid, Ancylotherium hennigi, a primitive buffalo, Simatherium kohllarseni, and two primitive suids, Potamochoeroides shawi (Bender, 1992) and Notochoerus scotti (Cooke, 1993).

Member 4 at Makapansgat has its greatest faunal associations with Member 3, hence their alternating positions in the seriations. Both deposits contain *Australopithe*- cus africanus as well as the ancient artiodactyls, Simatherium cf. kohllarseni and Notochoerus scotti, and Eisenhart (1974) had noted that the cercopithecids are virtually identical. Makapansgat Member 4 does pick up more extant fauna, but this is largely due to the microfauna from "rodent corner" which may have only a tenuous association with the large mammals in other parts of the cave. Although a stratigraphic separation between the two "members" exists in some parts of the cave, the difference may not have been of a geologically significant time period and Members 3 and 4 may be better viewed as different parts of the same cave exhibiting slightly different taphonomic processes (Maguire et al., 1985).

Taung

Most of the Taung fauna came from the Hrdlička deposits, which have their greatest affiliations with Sterkfontein Member 4. The faunal seriation of the time-sensitive species supports the recently espoused view that the Hrdlička deposits are roughtly contemporaneous with Sterkfontein Member 4 (McKee, 1993a). As the association is as strong with Swartkrans Member 1 as it is with Makapansgat Member 3, a slightly later age may be tenable.

The Taung hominid skull, however, probably came from a cave system associated with the Dart deposits, and has been posited to be considerably older than the age represented by the Hrdlička faunal assemblage (McKee, 1993b). There are few time-sensitive species yet known from the Dart deposits, so a faunal age is difficult to ascertain by the seriation method used here. All of the seven species likely to be from the Dart deposits are extinct, and only two time-sensitive species (A. africanus and Parapapio broomi) occur at other sites, so an age predating Makapansgat is conceivable. On the other hand, the occurrence of four unique, extinct species, as witnessed in the Dart deposits, is not unexpected at a time following deposition at Makapansgat (McKee, 1994). The Dart assemblage, including the type specimen of Australopithecus africanus. most likely falls between the times of Makapansgat (Members 3 and 4) and Sterkfontein Member 4 (McKee, 1993a).

Sterkfontein

Sterkfontein Member 4 has been extremely well sampled through 27 years of continuous excavation. As it has yielded considerably more of the late Pliocene fauna known from other Transvaal sites than does Makapansgat, it most likely postdates Makapansgat. However, Sterkfontein Member 4 shares many ancient species with Makapansgat, and thus shows the last appearance of seven time-sensitive species (Table 4), including A. africanus.

Member 5 at Sterkfontein marks the first appearance in southern Africa of the genus *Homo* as well as of stone tools. Its strongest faunal affinities are with Sterkfontein Member 4, followed by Kromdraai A. Being that it has lost some of the older species, notably *A. africanus* and *Notochoerus scotti*, it is best placed following the Taung Hrdlička deposits. It does not have as strong an affinity to Swartkrans as does Kromdraai B, so the seriation places Sterkfontein Member 5 before Kromdraai B.

Sterkfontein Member 5 thus falls between two sites with which it has very little in common. There are a number of possible explanations for this. It may have had a distinctly different environment from that of Taung, as it does today, and thus would have more in common with Sterkfontein Member 4. However, it seems to be nearly contemporaneous with Kromdraai B, yet shares fewer than half of all its species, and only 27% of its time-sensitive species with this nearby site in the same valley. This may be attributable to taphonomic considerations in which early Homo is a major contributor to the Sterkfontein Member 5 deposit (Vrba, 1975, 1976, 1980; Brain, 1981, 1985; McKee, 1991), but not to Kromdraai which has been viewed as a part death trap and part feeding lair of opportunistic predators (Vrba, 1981; Brain, 1981). Indeed, Kromdraai B has remarkably few bovids, and Sterkfontein Member 5 has a paucity of cercopithecids.

A final possibility for the interpretation of Sterkfontein Member 5, as suggested by Vrba (1985), is that temporally distinct deposits are mixed within the "member." Archaeological evidence from recent excavations (Kuman, 1994) has increased the likelihood of this interpretation. The timesensitive species, however, fit conformably within the seriated position of Member 5 (Table 4).

Kromdraai

Kromdraai A and B are most similar to each other, but Kromdraai B, with Australopithecus (Paranthropus) robustus, is most likely older. Both units are the same FRI of time-sensitive species with Makapansgat Members 3 and 4, and Kromdraai B has proportionally stronger ties to Sterkfontein Member 4. Kromdraai A, on the other hand, is the first site to show a large increase in the number of extant time-sensitive species as well as those species characteristic of the Cornelia Faunal Span. This is largely due to the increased proportion of bovid species known from the site.

A case could be made for Kromdraai A following Swartkrans Member 1, as suggested by the seriation. This is largely due to the similarity between Kromdraai A and Plovers Lake, but there are other considerations. Dinofelis piveteaui of Kromdraai A is seen to be an evolutionary descendent of D. barlowi, known from Swartkrans Member 1 and earlier sites (Cooke, 1991). Swartkrans Member 1 also has stronger faunal ties to Sterkfontein Member 4 and the Taung Hrdlička deposits. On the other hand, Kromdraai A has a strong resemblance to Kromdraai B and Sterkfontein Member 5, and has not yielded as many extant species as Swartkrans Members 2 and 3, so we suggest that on the current evidence it comes just before or very near in time to Swartkrans Member 1.

Swartkrans

The Swartkrans Formation harbors both *A. robustus* and *Homo* together for the first time. Each of the three members share the most species, including time-sensitive species, among themselves, and it has been suggested that they do not span a long period of time (Brain, 1988, 1993; Delson, 1988). However, Member 1 is clearly distinct from the others. It is the last site unit in the seriation to have strong faunal ties to Makapansgat, Taung, and Sterkfontein, and thus

sees the last appearance of eight time-sensitive species (Table 4). Recently Grine et al. (1993) have suggested a further link between Swartkrans Member 1 and Sterkfontein Member 5 with their identification of similarities in the hominid material, which they tentatively assign to *Homo habilis* sensu lato; Clarke (1985), however, refers the SK 847 cranium from Member 1 to *Homo* erectus. It should be noted as well that Swartkrans Member 1 also picks up five first appearances in the fossil record (or more if Kromdraai A follows it in time), thus clearly separating it from the early Australopithecus sites.

Swartkrans Members 2 and 3 then lose most of the fauna from sites of the Sterkfontein Faunal Span. By the time of Member 3, nearly half of the time-sensitive species, and 62% of all the species, are those that are extant today. Member 3 also provides the last glimpse of A. robustus.

There is a most remarkable similarity between Members 2 and 3 at Swartkrans, with an FRI for time-sensitive species of 100 and an overall FRI of 96. This may be because of a very close time frame covered by the deposition of the two members, but it is also possible that there has been some inadvertent mixing of specimens from the two members due to the decalcification of Member 3.

Plovers Lake

Plovers Lake, a cave site in the Blaaubank River Valley with Sterkfontein, Swartkrans, and Kromdraai, is the most recent addition to the known faunal assemblages of southern Africa. The fauna from Plovers Lake resembles most closely Kromdraai A and Swartkrans Member 3, and fits most consistently at or near the time of the latter. Plovers Lake presents the last appearance of Papio robinsoni (or Papio hamadryas robinsoni) and Procavia transvaalensis.

Cornelia

Cornelia is the last faunal site before the appearance of archaic *Homo sapiens* in the fossil record of southern Africa. It is especially significant for the last appearance of *Hipparion libycum*, the last time-sensitive species from Makapansgat Member 3. Cornelia has its greatest affinities with the Elandsfontein Main Site. Remarkably, only 33% of all its species are extant. It has yielded many of the species characteristically associated with early *Homo* in southern Africa, but predates the *Homo* sites where extant species appear in greater proportions.

Elandsfontein

The Elandsfontein Main Site ushers in archaic Homo sapiens with the "Saldanha" skull. The fossils include the last evidence of some key Plio-Pleistocene boundary species including Antidorcas recki, Theropithecus oswaldi, and Metridiochoerus andrewsi, as well as the last appearance of the genus Megantereon, thus attesting to its great age. However, it is the first site to have extant species represented in over half of the timesensitive fauna.

Cave of Hearths

The Acheulian levels near the bottom of the Cave of Hearths deposits in the Makapansgat valley, yielded a mandible and a radius of a human that have been referred by Tobias (1971) to Homo sapiens rhodesiensis. The relative position in time of this Early Stone Age (ESA) site is difficult to ascertain. Cornelia, Elandsfontein, and Cave of Hearths have "Upper Acheulian" (Butzer et al., 1974; Mason, 1988), but the latter site certainly postdates Cornelia and Elandsfontein on the basis of the fauna. The seriation results suggest, as did Cooke (1988), that the faunal assemblage is typical of the Middle Stone Age (MSA) sites. It has very close affinities to the MSA site of Equus Cave and is consequently seriated in variable positions following Elandsfontein. As the site of Equus Cave has apparent fully modern Homo sapiens (Grine and Klein, 1985), it fits that Cave of Hearths, with ESA artifacts and archaic *Homo sapiens*, should be older.

Florisbad

Florisbad comes out in the seriation in variable positions following Elandsfontein, with a strong similarity to Equus Cave. As with the other Late Pleistocene MSA sites, the faunal seriation alone can tell us little about its relative age. However, the MSA deposits of Florisbad has five extinct species (Table 4), and sometimes seriates before the ESA deposits of Cave of Hearths, suggesting that it may be older than some of the other MSA sites. Kuman and Clark (1986) estimate the age of the Florisbad cranium and associated fauna to be in the range of 100,000 to 200,000 years old.

Equus Cave

Faunal remains from Equus Cave show that a somewhat more lush local environment once existed in what is now the desert margin at Taung. The site has yielded only three extinct species: Antidorcas bondi, which lasts until at least 38,000 B.P. at Border Cave (Klein, 1977); Equus capensis, the large equid which survives to the very end of the Pleistocene; and Megalotragus priscus, a giant alcelaphine last seen at 16,000 BP (Klein, 1980). The faunal seriation thus tells us little about the place in time of Equus Cave; the presence of MSA artifacts, however, places it prior to about 40,000 BP (Klein et al., 1991).

Klasies River Mouth

Although Klasies is an important MSA site with some evidence for Homo sapiens in a Late Pleistocene context, the faunal analysis of the hominid-associated MSA I and MSA II levels gives few clues as to its relative age. The site bears its strongest resemblance to Equus Cave, but has only one extinct species, Pelorovis antiquus, which went extinct between 12,000 and 10,000 BP (Klein, 1980). Attempts have been made to seriate the total faunal assemblage from the cave complex (Binford, 1986), but these studies have been shown to be problematic (Thackeray, 1986, 1987). Despite such uncertainty, there is little question that the lower-most deposits are on the order of 100,000 years, perhaps between 80,000 and 115,000 years old (Deacon and Geleijnse, 1988). Thus Florisbad, Klasies River Mouth, and perhaps Equus Cave, fall within a relatively short time range during which whatever mammalian faunal trends may have

existed could not be picked up by the seriation technique.

SUMMARY AND CONCLUSIONS

The faunal seriation of southern African cave site assemblages has proven to be a useful tool. Although it confirms the general validity of the faunal spans devised in the 1960s by Ewer and Cooke, greater resolution within those spans is now possible. Moreover, as with fossil species, the naming of faunal spans can obscure the evolutionary continuity of many mammals through time, so a sequence of individual sites is preferable.

On the basis of the total analysis, both with the logistical seriation of time-sensitive mammals and subjective scrutiny of the data, the following sequence of sites is most likely: Makapansgat Member 3, Makapansgat Member 4, Taung Dart deposits, Sterkfontein Member 4 and Taung Hrdlička deposits, Sterkfontein Member 5 (in part) and Kromdraai B, Kromdraai A and Swartkrans Member 1, Swartkrans Member 2, Swartkrans Member 3, Plovers Lake, Cornelia, Elandsfontein Main Site, Cave of Hearths Acheulian levels, Florisbad and Equus Cave and Klasies River Mouth.

An important caveat emerges from analysis of the FRI values across the sites. For most of the sites at which there is more than one deposit or "member," the FRI values for each unit are highest for other units of the same site. This is true of Makapansgat, Sterkfontein, Kromdraai, and Swartkrans, and thus seems to be more than coincidental. At each site there are a number of possible explanations including (1) temporal continuity across adjacent members, (2) similar external and internal environments yielding taphonomic consistency, and/or (3) inadvertent mixing of the fossils and breccias from separate deposits. The need to discern among these options highlights the importance of continuing field work and analysis.

The site sequence proposed here may change for some of the closely related sites, as the seriation for those sites is sensitive to the inclusion or exclusion of particular sites and species. The overall picture, however, is consistent for most sites and certainly the proposed sequence may be used as a working framework for the interpretation of evolutionary patterns among the mammal species and as corroborating evidence for independent geochronological and biochronological techniques. Ongoing research at Sterkfontein, Taung, and Plovers Lake, as well as recently initiated excavations at Gladysvale, Makapansgat, and Kromdraai, undoubtedly will yield evidence that can refine the picture of Neogene mammalian evolution and the place of hominids in it.

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