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Evolution of the modern baboon (*Papio hamadryas*): A reassessment of the African Plio-Pleistocene record



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ABSTRACT

Baboons (*Papio hamadryas*) are among the most successful extant primates, with a minimum of six distinctive forms throughout Sub-Saharan Africa. However, their presence in the fossil record is unclear. Three early fossil taxa are generally recognized, all from South Africa: *Papio izodi*, *Papio robinsoni* and *Papio angusticeps*. Because of their derived appearance, *P. angusticeps* and *P. robinsoni* have sometimes been considered subspecies of *P. hamadryas* and have been used as biochronological markers for the Plio-Pleistocene hominin sites where they are found.

We reexamined fossil *Papio* forms from across Africa with an emphasis on their distinguishing features and distribution. We find that *P. robinsoni* and *P. angusticeps* are distinct from each other in several cranial features, but overlap extensively in dental size. Contrary to previous assessments, no diagnostic craniomandibular material suggests these two forms co-occur, and dental variation at each site is comparable to that within *P. h. ursinus*, suggesting that only one form is present in each case. *P izodi*, however, may co-occur with *P. robinsoni*, or another *Papio* form, at Sterkfontein Member 4.

P izodi appears more primitive than *P. robinsoni* and *P. angusticeps*. *P. robinsoni* is slightly distinct from *P. hamadryas* subspecies in its combination of features while *P. angusticeps* might be included within one of the modern *P. hamadryas* varieties (i.e., *P. h. angusticeps*). No definitive *Papio* fossils are currently documented in eastern Africa until the Middle Pleistocene, pointing to southern Africa as the geographic place of origin for the genus. These results have implications for Plio-Pleistocene biochronology and baboon evolution.

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1. Introduction

The savannah baboons of the genus *Papio* are among the most well-known and successful extant primates, with a minimum of six recognizable populations distributed throughout Africa outside of the central forest area, as well as in southern Arabia (Thorington and Groves, 1970; Szalay and Delson, 1979; Jolly, 1993,

* Corresponding author. E-mail address: cgilbert@hunter.cuny.edu (C.C. Gilbert). 2001; Groves, 2001; Frost et al., 2003; Grubb et al., 2003; Fleagle, 2013; see Fig. 1). Despite their evolutionary success and wide distribution across modern African ecological communities, the origins of the genus in the fossil record are not clear. Current molecular and morphological evidence suggests that, among living African papionins, *Papio* is closely related to *Theropithecus*, *Lophocebus*, and *Rungwecebus* (Disotell et al., 1992; Disotell, 1994, 2000; Harris and Disotell, 1998; Fleagle and McGraw, 1999, 2002; Tosi et al., 1999, 2003; Davenport et al., 2006; Gilbert, 2007, 2013; Olson et al., 2008; Burrell et al., 2009; Zinner et al., 2009; Gilbert et al., 2009a, 2011; Roberts et al., 2010), and



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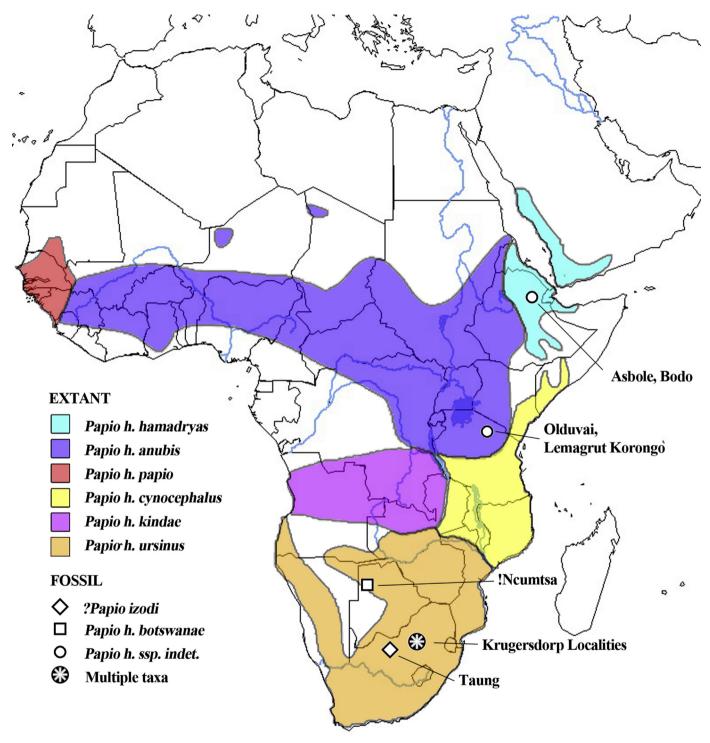


Figure 1. Map of Africa illustrating the geographic distribution of extant and fossil *Papio* populations. Krugersdorp localities include Sterkfontein, Swartkrans, Kromdraai, Bolt's Farm, Cooper's A-D, Gladysvale, Drimolen, Malapa, Haasgat, and Skurweberg.

within this group, the most recent analyses suggest a closer relationship between *Papio* and *Lophocebus*, with *Theropithecus* at the base of this clade (Perelman et al., 2011; Springer et al., 2012; Guevara and Steiper, 2014; Pugh and Gilbert, in press). The position of *Rungwecebus* is controversial, being most recently reconstructed as the sister taxon to *Papio* in molecular studies (Davenport et al., 2006; Olson et al., 2008; Burrell et al., 2009; Zinner et al., 2009; Roberts et al., 2010), yet most similar to *Lophocebus* in morphological comparisons (Jones et al., 2005;

Davenport et al., 2006; Singleton, 2009; Singleton et al., 2010; Gilbert et al., 2011a; Gilbert, 2013). Thus, the combination of these data sources implies a close relationship among these three taxa pending additional data.

While *Rungwecebus* is unknown in the fossil record, the earliest specimens of *Theropithecus* are dated to at least 4.2 Ma (Frost, 2001a; Harris et al., 2003; Jablonski et al., 2008; Frost et al., 2014; Frost et al., in revision; Gilbert and Frost, personal obs.). Undoubted *Lophocebus* specimens first appear in the fossil record

by approximately 2.0 Ma (Leakey and Leakey, 1976; Frost, 2001a; Jablonski et al., 2008; Gilbert, 2013) and possibly as early as 3.5 Ma (Harrison and Harris, 1996). Therefore, fossil specimens attributable to the genus *Papio* should also be present in the fossil record by at least 2.0 Ma, and potentially well into the Pliocene.

In fact, numerous taxa in the African Plio-Pleistocene have been assigned to the genus *Papio* over the last century, but many of them have proven to be more properly assigned to other genera such as Parapapio, Theropithecus, Soromandrillus and Gorgopithecus (see Szalay and Delson, 1979; Eck and Jablonski, 1984, 1987; Delson and Dean, 1993; Jablonski, 2002; Jablonski and Frost, 2010; Gilbert, 2013; Gilbert et al., 2016a). Currently, there are at least four named and widely recognized fossil Papio taxa (Papio izodi, Papio [hamadryas] angusticeps, Papio [hamadryas] robinsoni, and Papio hamadryas botswanae) from the African Plio-Pleistocene record, with other specimens placed in Papio sp. indet. that may represent one or more additional taxa. In this paper, we review the complicated and confused taxonomy and fossil record of Papio with particular focus on South Africa and, based on updated morphological analyses, propose a revised taxonomy of the genus.

1.1. History of Papio fossil record

Baboons are distinct and iconic primates that have been familiar to humans for centuries (Morris, 2013). Perhaps because of their distinctiveness, large, generalized monkeys with long rostra have been referred to as "baboons" for much of the last 150 years (e.g., see Szalay and Delson, 1979; Jablonski, 2002; and references therein). With increasing resolution of papionin systematics, it is now clear that there are several large-bodied, long-faced papionin genera. Their taxonomy has a complex history with interpretations varying from author to author (e.g., Haughton, 1925; Remane, 1925; Gear, 1926; Broom, 1936, 1940; Dietrich, 1942; Freedman, 1957; Jolly, 1965, 1967; Hill, 1967; Leakey, 1969; Szalay and Delson, 1979; Iwamoto, 1982; Delson, 1984; Eck and Jablonski, 1984, 1987; McKee, 1993; Delson and Dean, 1993; Frost, 2001a, 2007a; 2007b; Frost and Delson, 2002; Jablonski et al., 2008; Williams et al., 2012; Gilbert, 2013).

As there is little agreement on the taxonomy of extant Papio populations (see Thorington and Groves, 1970; Jolly and Brett, 1973; Szalay and Delson, 1979; Jolly, 1993, 2003; Groves, 2001; Grubb et al., 2003; Frost et al., 2003; Zinner et al., 2013), a brief explanation of the taxonomy used in this paper is warranted. Even though there is a significant amount of morphological and behavioral variation within and between the various populations of extant Papio, for the purposes of this paper we favor recognizing the six commonly recognized populations at the subspecies level within a single species, Papio hamadryas (i.e., P. h. ursinus, P. h. cynocephalus, P. h. kindae, P. h. anubis, P. h. hamadryas, P. h. papio). Given the clinal pattern of cranial variation (e.g., Frost et al., 2003), the clinal pattern of mtDNA variation (e.g., Newman et al., 2004; Burrell, 2008; Zinner et al., 2009, 2011), and the extensive hybridization between populations observed in the wild (e.g., Jolly et al., 2011), using the Biological Species Concept (BSC) it seems that a single species is warranted. Furthermore, there is little doubt that many of the extant subspecies display an overall level of craniodental similarity that would be difficult to distinguish at the species level in the fossil record (e.g., Jolly, 1993). Most relevant for the current discussion, from a paleontological view, this taxonomic scheme also has the advantage of easily distinguishing between modern forms of Papio (i.e., P. hamadryas subspecies) and more distinctive fossil forms which can be recognized more readily at the species level.

On the other hand, at least two of us (CCG, KDP) prefer the Phylogenetic Species Concept (PSC), which is more widely accepted

in current primate studies (e.g., see Groves, 2001, 2014; Fleagle, 2014; Jolly, 2014; Louis and Lei, 2014; Rylands and Mittermeier, 2014; Silcox, 2014; Tattersall, 2014; Yoder, 2014; Zimmerman and Radespiel, 2014; Zinner and Roos, 2014) and has the advantage of relying on consistent phenotypic (in this case, morphological) differences rather than reproductive isolation as a criterion of separation. Given the widespread documentation of hybridization among commonly recognized primate species (e.g., see Tung and Barreiro, 2017 and references therein), some even representing different genera (most relevant in this case, Papio x Rungwecebus and Papio x Theropithecus; see Davenport et al., 2006; Olson et al., 2008; Burrell et al., 2009; Zinner et al., 2009, 2018; Roberts et al., 2010; Tung and Barreiro, 2017), it seems more and more difficult to rely on reproductive isolation as the defining characteristic of a species or evolutionary coherent lineage/population. More directly relevant from a paleontological perspective, it is impossible to know whether one fossil population was reproductively isolated from another, making any paleontologist reliant on morphological features to diagnose species anyway. Thus, for those who prefer the PSC in the study of Papio taxonomy, each extant baboon population is recognized as its own distinct species: Papio anubis, Papio cynocephalus, P. hamadryas, Papio kindae, P. papio, and Papio ursinus. In the fossil record, under the PSC scheme, a fossil baboon that is considered a part of the modern radiation would still be ranked at the species level, while under the BSC scheme it would simply be another subspecies of P. hamadryas.

A brief breakdown of the morphological features and current distribution of named fossil *Papio* species is as follows (see also Figs. 2–4):

<u>1.1.1. Papio izodi</u> *P. izodi* was first named by Gear (1926) based on material from Taung. For the next three decades, this and subsequent material from Taung was variously assigned to *Papio* and *Parapapio* and for a short time even synonymized with *Parapapio antiquus* (Broom, 1936, 1940). Freedman (1957, 1963, 1965) reviewed the material from Taung and allocated all specimens with a perceived anteorbital drop to *P. izodi*, an approach that has generally been followed ever since. A more detailed discussion of *P. izodi* early taxonomic history is given by Freedman (1957). Currently, *P. izodi* is recognized from the Plio-Pleistocene sites of Taung (e.g., Gear, 1926; Freedman, 1957; Szalay and Delson, 1979; Delson, 1984, 1988; Gilbert, 2013) and Sterkfontein Members 2 and 4 (e.g., Eisenhart, 1974; Delson, 1984, 1988; McKee, 1993; Pickering et al., 2004; Heaton, 2006; Gilbert, 2013).

Relative to other *Papio* taxa, *P. izodi* is small-to-medium sized and displays molars and orbits that are large relative to cranial size, a relatively short (anteroposteriorly) and broad snout, a relatively dorso-ventrally short neurocranium, and a relatively dorsoventrally short malar region (Figs. 2–4; Freedman, 1957; Delson, 1988; McKee, 1993; Gilbert, 2013; Gilbert et al., 2015). In addition, this taxon displays shallow maxillary fossae, a variable anteorbital drop, weak to absent mandibular corpus fossae, and weak development of the maxillary ridges (in males and females) compared to most other *Papio* taxa (Figs. 2–4; Freedman, 1957; McKee, 1993; Gilbert, 2013; Gilbert et al., 2015). Similar to other *Papio* species, *P. izodi* specimens typically exhibit a prominent glabella, pinched temporal lines, downturned nuchal lines, a moderate supraorbital torus, and a moderate ophryonic groove or post-glabellar depression (Figs. 2–4; Freedman, 1957; Gilbert, 2007, 2013).

In a recent phylogenetic analysis, Gilbert (2013) hypothesized that *P. izodi* may in fact be a stem African papionin, largely based on the distinctive and primitive characters listed above (e.g., shallow facial fossae, weak development of maxillary ridges in males, weak to absent mandibular corpus fossae, and variable anteorbital drop). Therefore, *P. izodi* may warrant generic separation from the other

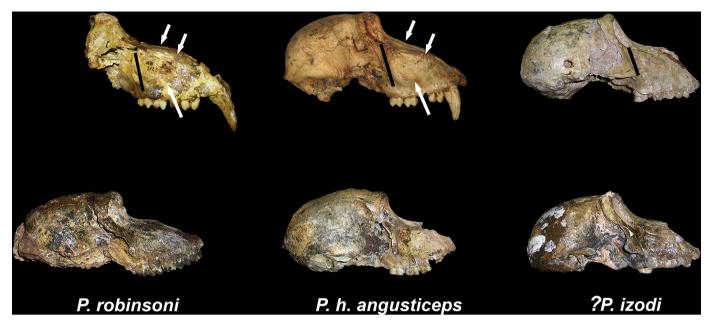


Figure 2. Major morphological differences between specimens of *P. robinsoni* (left), *P. h. angusticeps* (center), and *?P. izodi* (right) in lateral view. Top row = male specimens, bottom row = female specimens. *P. robinsoni* males represented by SK 555; *P. h. angusticeps* males represented by CO 100; and *?P. izodi* represented by SWP Uncatalogued Cranium from Member 2. *P. robinsoni* females represented by UCMP 56797; *P. h. angusticeps* represented by UCMP 56767; and *?P. izodi* represented by TP-10 (ex-UCMP 56605). Specimens are approximately to scale. Differences between taxa are most obvious among male specimens. In males, note the difference in the height of the maxillary ridges and of the maxillary fossae (white arrows), and relative malar height (black bars). In contrast to *P. robinsoni* and *P. h. angusticeps*, *?P. izodi* males do not possess definitive maxillary ridges or maxillary fossae. Many of the same differences are also seen to a lesser extent among the female specimens.

Papio taxa, and we formally recognize this uncertainty from this point forward with the taxonomic convention *?P. izodi.*

1.1.2. Papio robinsoni Freedman (1957) named a large series of partial crania, jaws and more fragmentary material from Swartkrans, Swartkrans II, Cooper's, Kromdraai, Bolt's Farm, Gladysvale and Skurweberg as a new species Papio robinsoni. This taxon has been described from a number of sites including Swartkrans Members 1-3, Swartkrans II, Skurweberg, Cooper's A, Kromdraai A, Bolt's Farm, Drimolen Main Quarry, and possibly Sterkfontein Member 4 (Freedman, 1957; Eisenhart, 1974; Szalay and Delson, 1979; Delson, 1984, 1988; McKee et al., 1995; Keyser, 2000; Adams et al., 2016). Dentally and cranially, P. robinsoni is most similar in size to modern P. hamadryas ursinus and P. hamadryas anubis, but Freedman (1957) differentiated P. robinsoni from modern P. hamadryas ssp. by features such as a more flattened muzzle dorsum with nasals lying inferior to the maxillary ridges in lateral view, a high frequency of the maxillae meeting in the midline and covering the nasal bones in males, a relatively shorter snout. relatively large P⁴s, more rounded maxillary ridges and definitive, but shallow-moderate depth (i.e., weaker) facial fossae (Figs. 2-4). Like other Papio taxa, P. robinsoni exhibits a definitive anteorbital drop, flattened muzzle dorsum, a prominent glabellar/supraorbital region, pinched temporal lines, downturned nuchal lines, and an ophryonic groove or depression behind the orbits (Figs. 2-4). Delson (Szalay and Delson, 1979, 1984, 1988) has argued that the distinctive features displayed by P. robinsoni can be found within the large amount of variation encompassing the extant Papio hamadryas subspecies, albeit in different frequencies, and on this basis preferred to recognize the taxon as P. h. robinsoni, an arrangement followed by Frost (2007a,b), Williams et al. (2012), and Gilbert (2013). In this paper, we use the P. robinsoni convention, and evaluate its morphological distinctiveness from P. hamadryas in the analyses below.

1.1.3. Papio angusticeps On the basis of a partial cranium and other material from Kromdraai A, Broom (1940) originally described this series as a new species of Parapapio. Freedman (1957) transferred the type material along with additional specimens from Kromdraai, Cooper's, and Minaar's Cave to the genus Papio based on the clear presence of an anteorbital drop and muzzle morphology. While all subsequent authors have agreed that this material belongs in the genus Papio there has been debate about its specific distinction from ?P. izodi (Szalay and Delson, 1979; Jablonski, 2002; Heaton, 2006; Gilbert, 2008; Jablonski and Frost, 2010). Others have noted shared similarities between the P. angusticeps material and modern P. hamadryas, especially the small-bodied P. h. kindae, and therefore separated it from ?P. izodi (Delson, 1984, 1988; McKee, 1993; Williams et al., 2012; Gilbert, 2013; Gilbert et al., 2015). In fact, some have recognized it as a subspecies of P. hamadryas (implied by Delson, 1984, 1988; and Frost, 2007a,b; first formalized by Williams et al., 2012; Gilbert et al., 2015), a decision we follow here from this point forward. For a more detailed discussion of the early taxonomic history of P. h. angusticeps, see Freedman (1957).

P. h. angusticeps (Broom, 1940) has been recognized at Kromdraai A, Kromdraai B, Cooper's A, Haasgat, Gladysvale, and Malapa (e.g., Freedman, 1957; Szalay and Delson, 1979; Delson, 1984, 1988; McKee and Keyser, 1994; McKee et al., 1995; Gilbert, 2013; Gilbert et al., 2015). As discussed above, in most respects, *P. h. angusticeps* is extremely similar to modern *P. hamadryas* sspp., possessing a definitive anteorbital drop, moderate-to-deep maxillary fossae, maxillary ridges in males and females, a long and narrow muzzle, a relatively tall malar region, a prominent glabella and supraorbital region, pinched temporal lines, downturned nuchal lines, and a moderate ophryonic groove/postglabellar depression (Figs. 2–4; Freedman, 1957; Delson, 1984, 1988; McKee, 1993; Gilbert, 2013; Gilbert et al., 2015). Freedman (1957) suggested that it possesses weaker maxillary ridges than extant *P. hamadryas*, but we disagree.

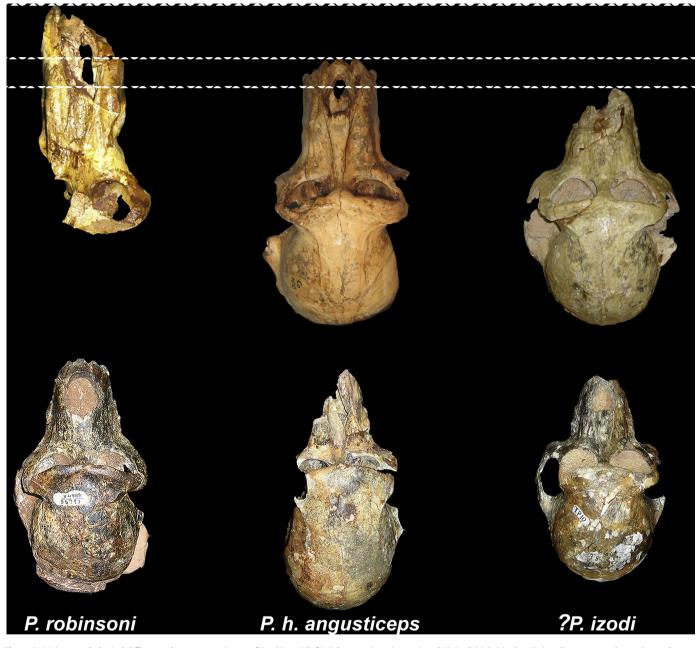


Figure 3. Major morphological differences between specimens of *P. robinsoni* (left), *P. h. angusticeps* (center), and *?P. izodi* (right) in dorsal view. Top row = male specimens, bottom row = female specimens. *P. robinsoni* males represented by SK 555; *P. h. angusticeps* males represented by CO 100; and *?P. izodi* represented by SWP Uncatalogued Cranium from Member 2. *P. robinsoni* females represented by UCMP 56797; *P. h. angusticeps* represented by UCMP 56767; and *PP. izodi* represented by TP-10 (ex-UCMP 56605). Specimens are approximately to scale. Differences between taxa are most obvious among male specimens. In males, note the difference in the relative breadth and length of the snout (length from orbitale inferior to [estimated] prosthion highlighted for each species with dotted lines). *?P. izodi* males have relatively shorter and broader snouts compared to *P. robinsoni* and *P. h. angusticeps*. Many of the same differences are also seen to a lesser extent among the female specimens.

<u>1.1.4. Papio spelaeus</u> This species was based on a single large male cranium encrusted in calcite matrix of unknown provenance first described by Broom (1936). Subsequent authors (Freedman, 1957; Szalay and Delson, 1979) have considered it to be of Holocene or at least nearly Holocene age and recognized its similarity to modern chacma baboons. We also recognize it as a junior synonym of *P. hamadryas ursinus*, and it will not be considered further here.

<u>1.1.5. Papio hamadryas botswanae</u> P. h. botswanae is currently known from a single cranium recovered from a Middle Pleistocene cave system in the !Ncumtsa (Koanaka) Hills, Botswana (Williams

et al., 2012). In cranio-dental size it is smaller than most *P. h. ursinus* and *P. h. anubis* males, similar to those of *P. h. hamadryas* and *P. h. papio*, but larger than *P. h. kindae*. While within the range of cranial shape variation of extant *P. hamadryas* populations, 3D cranial geometric morphometrics demonstrated the uniqueness of this specimen relative to the extant *P. hamadryas* subspecies, with *P. h. botswanae* displaying an extremely prominent supraorbital region, wide neurocranium and a relatively short and broad muzzle for a baboon of its size. Its muzzle is moderately klinorynch, but less so than that of *P. h. ursinus*, the modern subspecies with the most klinorynch rostra (Williams et al., 2012).

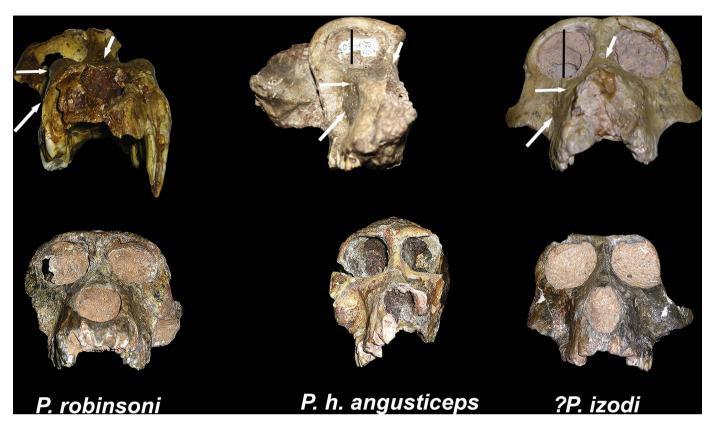


Figure 4. Major morphological differences between specimens of *P. robinsoni* (left), *P. h. angusticeps* (center), and *?P. izodi* (right) in frontal view. Top row = male specimens, bottom row = female specimens. *P. robinsoni* males represented by SK 555; *P. h. angusticeps* males represented by HGD 1249; and *?P. izodi* represented by SWP Uncatalogued Cranium from Member 2. *P. robinsoni* females represented by UCMP 56797; *P. h. angusticeps* represented by UCMP 56767; and *?P. izodi* represented by TP-10 (ex-UCMP 56605). Specimens are approximately to scale. Differences between taxa are most obvious among male specimens. In males, note the differences in orbit height/size (white/black bars), and in maxillary fossae development, maxillary ridge development, and height of the nasals relative to the maxillary ridges (white arrows). The orbits of *?P. izodi* are relatively larger than seen in *P. robinsoni* and *P. h. angusticeps*. As seen in lateral view and, again, in contrast to *P. robinsoni* and *P. h. angusticeps*, *?P. izodi* males do not possess definitive maxillary ridges or maxillary fossae. Many of the same differences are also seen to a lesser extent among the female specimens.

1.2. Goals of this study

The three fossil taxa from South Africa discussed above have been argued to be early members of the genus Papio. However, the morphological diagnoses of these taxa, summarized above, have not been reassessed in a number of years despite great changes to our understanding of papionin phylogeny and evolution (see above). Thus, it is possible that an updated analysis of these fossil Papio populations may help refine and revise our understanding of baboon evolution and taxonomy. For example, as alluded to above, Gilbert (2013) recently suggested that ?P. izodi from Taung and Sterkfontein retains many plesiomorphic features that may require taxonomic reassignment to a genus other than Papio. Both *P. robinsoni* and *P. h. angusticeps*, on the other hand, have at times been considered as subspecies of the extant species (Szalay and Delson, 1979; Delson, 1988; Frost, 2007a; Gilbert et al., 2015), and both were thus claimed to represent the first appearance of the modern form. Because of their derived appearance, P. h. angusticeps and P. robinsoni have further been used as important biochronological markers for the Plio-Pleistocene sites at which they are found, and they have been suggested to co-occur at Kromdraai A and B and Cooper's A (Freedman, 1957; Delson, 1984, 1988; McKee et al., 1995; Heaton, 2006; Jablonski and Frost, 2010).

In order to test these taxonomic and distributional hypotheses, a broader and more comprehensive review of extant and fossil *Papio* material is needed. Our goals are to 1) clarify which fossil populations can be referred to any specific named taxa; 2) offer a revised morphological diagnosis of fossil *Papio* species and the genus *Papio* based on our updated hypodigms; and 3) review the resulting distribution of fossil *Papio* species and populations in the Plio-Pleistocene fossil record. This has implications for our understanding of the origins and evolution of the genus, its first appearance in the fossil record, its biogeography, its taxonomy, and the biochronology of the sites where it occurs.

2. Materials and methods

In 2012-2016, we reviewed all major Plio-Pleistocene collections containing African cercopithecoids with the explicit goal of refining the taxonomy at each site and, ultimately, providing an updated biochronological analysis relative to the landmark studies of Delson (1984, 1988) on the basis of our results. Fossil specimens examined during this study (and in some cases previously by Delson) were housed at: the Ditsong National Museum, Pretoria (TMP, formerly Transvaal Museum); the Council for Geosciences, Pretoria (CGS); the Bernard Price Institute, Johannesburg (BPI); the University of the Witswatersrand Anatomy Department, Johannesburg (UW-AD); the Evolutionary Studies Institute, Johannesburg (ESI); the Iziko South Africa Museum, Cape Town (SAM); the National Museum of Tanzania, Dar es Salaam (NMT); the National Museums of Kenya, Nairobi (KNM); the National Museum of Ethiopia, Addis Ababa (NME); the Natural History Museum, London (NHM-UK); the Museum für Naturkunde, Berlin (MB); the Bayerische Staatssammlung für Paläontologie und Geologie, München, Germany (BSPG [formerly BSM]); and the University of California Museum of Paleontology, Berkeley (UCMP).

Thus, we examined nearly all of the fossil Papio material that was available at the time, employing typical comparative morphological methods. Specimens were qualitatively and quantitatively evaluated for key morphological features (Freedman, 1957; Delson, 1975, 1988; Szalay and Delson, 1979; Frost, 2001a,b; Fleagle and McGraw, 2002; Gilbert, 2007, 2013). Digital calipers were used to collect standard craniometric measurements, and digital photographs were taken of the more complete specimens whenever possible. Qualitative characters were scored according to the criteria described in Gilbert (2013). Comparative measurements and reference qualitative comparisons with extant taxa were taken from specimens housed at the American Museum of Natural History, New York (AMNH) as well as from Frost (2001a), Gilbert (2013), and the PRIMO online database (http://primo.nycep.org). For the full list of samples, characters, and measurements used, see Tables 1–7 and the references therein.

Univariate and multivariate statistical analyses were performed in PAST v3.14 (Hammer et al., 2001) and SPSS v23.0.0.2 (IBM Corp.). To investigate ranges of variation in fossil Papio populations across fossil sites, we compared our dental measurements in the form of premolar and molar area boxplots from the upper and lower dentitions of the fossil specimens to a modest sample of a living *P. hamadryas* subspecies, *P. h. ursinus* (n = 120-152 depending on tooth position; see Tables 1 and 2 for summary information and Supplementary Online Material [SOM] Tables S1-S4 for a complete list of extant and fossil Papio specimens and measurements). P. h. *ursinus* is an appropriate comparison in this case because it is similar in size to both P. h. angusticeps and P. robinsoni (slightly larger than *P. angusticeps*, very similar in dental size to *P. robinsoni*). thereby controlling for any significant size effects, and it is found in the same geographical region as the fossil populations in question, thereby controlling for any strong latitudinal or biogeographical effects. We reasoned that if the ranges found in the fossil samples were similar to, or less than, that observed in a modern subspecies, then we have no basis for definitively recognizing multiple Papio taxa in the absence of more diagnostic craniodental material. We also assessed the range of variation found across fossil and extant taxa using our new classification system (i.e., P. robinsoni, P. h. angusticeps, compared to P. h. ursinus). If the range of variation in the fossil taxa is again comparable to that observed in *P. h. ursinus*, this would suggest that our revised classification system does not introduce excessive variation despite time-averaging and including material across multiple sites in each fossil taxon. In addition to univariate comparisons, multivariate analyses were performed in relation to the 3D geometric morphometric analysis described in more detail below.

2.1. Geometric morphometrics

To quantitatively assess the overall cranial shape of fossil *Papio* taxa we collected 3D landmarks following the protocol of Frost et al. (2003) on 24 relatively complete fossil crania (Table 3). These were superimposed with a sample of 778 extant papionins from PRIMO (Table 4) in Morpheus (Slice, 2013) by Generalized Procrustes Analysis (GPA) to remove variation due to size, position, and orientation (Rohlf and Slice, 1990). Centroid size, the square root of the sum of the squared distance of each landmark to the centroid, was used as a measure of size (Bookstein, 1991). In order to minimize the effects of taphonomic deformation and impute bilateral landmarks missing from one side, all specimens were mirrored and averaged (Gunz et al., 2009).

Principal Components Analysis (PCA) was used as a data reduction and ordination technique to explore the distribution of specimens in shape space (Neff and Marcus, 1980; Manly, 1994). To maximize the number of specimens, remaining missing landmarks

were imputed by species-sex means where available. For the PCA, grand means were used to fill in any remaining landmarks. Previous studies of papionin cranial variation have demonstrated that allometry accounts for the largest share of total variance, and most of the shape differences were related to sex (e.g., Singleton, 2002; Frost et al., 2003). Therefore, following Frost et al. (2003) and Williams et al. (2012), we regressed GPA aligned coordinates against the natural log of centroid size and sex to produce adjusted landmark configurations (Frost et al., 2003). Discriminant function analysis on genera was performed on the GPA aligned coordinates with the extant papionins used as the model and fossil papionin specimens as unknowns. Cross-validation, where each known specimen was individually removed from calculation of the discriminant functions and then reclassified, was used to evaluate the reliability of the function.

2.2. Phylogenetic analysis

In addition to quantitative and qualitative craniometric comparisons, a 362 character phylogenetic analysis using parsimony, including 36 extant and fossil cercopithecoid species, was conducted in PAUP* 4.0b10 (Swofford, 2003). Relative to the previous analyses of Gilbert and colleagues (Gilbert and Rossie, 2007; Gilbert et al., 2009a, 2016a; Gilbert, 2013), the current analysis is distinct in that it considers extant species rather than genera as operational taxonomic units (OTUs), includes additional extant and fossil data from PRIMO and Frost (2001a), and includes the fossil taxon *P. robinsoni*. For full sample sizes by species, see Tables 5 and 6. In addition to the 318 craniodental characters listed in Gilbert (2013), 44 postcranial characters (22 for each sex) previously noted to be phylogenetically informative were also added to the current matrix (Table 7; see also Gilbert et al., 2016a, 2016b). The matrix used in this study is provided as a nexus file in the SOM.

Three separate analyses were performed. The first analysis, here termed "Morphology-only", constrained only the outgroup taxa (see below), thereby allowing all ingroup taxa to group freely. The second analysis, here termed "Molecular backbone P/L/R", incorporated the results of molecular studies, which consistently suggest a Cercocebus/Mandrillus (C/M) and Papio/Lophocebus/Rungwecebus/ Theropithecus (P/L/R/T) clade among extant African papionin taxa. Within the P/L/R/T clade, recent evidence suggests a closer relationship among P/L/R (see Introduction). Therefore, we imposed three molecular backbone/scaffold constraints within the ingroup at the genus level, to construct a tree consistent with these groupings, i.e., ((C,M),(T,(P,L,R))). Ingroup fossil taxa and extant species within genera were then allowed to float freely among these constraints. The third analysis, "Molecular backbone P/R", differs from "Molecular backbone P/L/R" in that only a sister relationship between Papio and Rungwecebus is also enforced within P/ L/R/T, as suggested by all nuclear DNA analyses including the kipunji (i.e., ((C,M),(T,L,(P,R)))). Again, ingroup fossil taxa and extant species within each genus were allowed to float freely among these constraints. In all analyses, a 10,000 replication, random addition sequence heuristic search was employed with Victoriapithecus, Allenopithecus, Parapapio lothagamensis, and Macaca constrained as successive outgroups to find the most parsimonious trees (MPTs). For clade support, a 1000 replication bootstrap procedure with replacement was conducted.

3. Results

3.1. Review of South African fossil Papio by site

Here we review the distribution of fossil *Papio* specimens at the main South African sites, following the listing of sites in Table 8 as

Taxon	P ⁴ BL	P ⁴ MD	M ¹ MBL	M ¹ DBL	$M^1 MD$	M ² MBL	M ² DBL	$M^2 MD$	M ³ MBL	M ³ DBL	M ³ MD
?Papio izodi	8.3 n = 10 (7.2–9.5)	6.4 n = 12 (5.0-7.2)	9.6 n = 9 (7.6-10.6)	9.2 n = 8 (8.8-9.9)	9.6 n = 13 (8,2-10.7)	11.3 n = 12 (9.8-12.5)	$ 10.4 \\ n = 13 \\ (9.1-11.4) $	$ 11.6 \\ n = 15 \\ (10.8-12.9) $	$ 11.4 \\ n = 11 \\ (9.9-12.3) $	9.9 n = 10 (8.7-10.7)	$ \begin{array}{r} 11.4 \\ n = 12 \\ (10.5 - 12.8) \end{array} $
Papio hamadryas angusticeps	8.2 n = 23 (6.7-9.1)	7.1 n = 25 (6.1-8.3)	9.5 n = 27 (7.4-11.0)	8.6 n = 28 (7.3–10.2)	10.3 n = 29 (8.0-12.3)	$ 11.2 \\ n = 27 \\ (9.4-12.9) $	$ 10.0 \\ n = 28 \\ (7.8-11.6) $	$ \begin{array}{r} 12.3 \\ n = 30 \\ (11.0-14.4) \end{array} $	11.1 n = 24 (8.4-12.8)	9.1 n = 24 (6.7-11.2)	$ \begin{array}{r} 12.2 \\ n = 26 \\ (10.0 - 14.4) \end{array} $
Papio robinsoni	9.4 n = 25 (8.3-10.9)	7.8 n = 26 (6.2-9.4)	10.5 n = 33 (9.0-12.1)	9.4 n = 29 (7.9–10.9)	$ 10.9 \\ n = 32 \\ (8.8-13.4) $	$ 11.9 \\ n = 41 \\ (10.0-13.7) $	10.8 n = 38 (8.9–13.2)	13.2 n = 42 (10.6-15.7)	$ 12.1 \\ n = 30 \\ (10.2-14.7) $	9.8 n = 29 (7.7-11.9)	$ \begin{array}{r} 13.2 \\ n = 28 \\ (11.1 - 15.5) \end{array} $
Papio cf. robinsoni Swartkrans Member 2	9.1 n = 1 (9.1–9.1)	7.5 n = 1 (7.5-7.5)	$ \begin{array}{r} 10.1 \\ n = 1 \\ (10.1 - 10.1) \end{array} $	9.1 n = 1 (9.1–9.1)	$ \begin{array}{r} 11.1\\n=1\\(11.1-11.1)\end{array} $	$ 11.8 \\ n = 1 \\ (11.8-11.8) $	10.5 n = 1 (10.5-10.5)	$ \begin{array}{r} 13.4 \\ n = 1 \\ (13.4 - 13.4) \end{array} $	$ 11.6 \\ n = 1 \\ (11.6-11.6) $	9.6 n = 1 (9.6-9.6)	$ 13.2 \\ n = 1 \\ (13.2-13.2) $
Papio cf. robinsoni Kromdraai B	9.1 n = 2 (8.3-9.8)	8.2 n = 3 (7.1–9.0)	10.8 n = 1 (10.8-10.8)	9.7 n = 1 (9.7–9.7)	$ \begin{array}{r} 11.6 \\ n = 1 \\ (11.6-11.6) \end{array} $	12.0 n = 2 (11.3-12.7)	10.9 n = 2 (10.4-11.3)	$ \begin{array}{r} 14.0 \\ n = 2 \\ (13.1-15.0) \end{array} $	10.1 n = 2 (8.4-11.7)	7.3 n = 2 (5.2–9.4)	$ 11.2 \\ n = 2 \\ (10.0-12.4) $
Papionini gen. et sp. indet. Swartkrans Member 3	9.1 n = 1 (9.1–9.1)	7.0 n = 1 (7.0-7.0)	10.4 n = 1 (9.8–10.9)	9.2 n = 1 (8.8-9.5)	10.0 n = 1 (9.5–10.5)	$ 11.9 \\ n = 1 \\ (11.5-12.7) $	$ \begin{array}{r} 11.1 \\ n = 1 \\ (10.3 - 12.3) \end{array} $	13.1 n = 1 (11.7-14.2)	$ 12.0 \\ n = 1 \\ (11.3-12.9) $	10.5 n = 1 (9.8–11.4)	13.5 n = 1 (12.2-14.8)
Papionini gen. et sp. indet. Cooper's D			9.0 n = 3 (8.5-9.4)	8.8 n = 3 (8.5-9.2)	$ 11.5 \\ n = 3 \\ (10.6-13.0) $	8.8 n = 1 (8.8-8.8)	$ \begin{array}{r} 10.0 \\ n = 1 \\ (10.0-10.0) \end{array} $	$ \begin{array}{r} 14.0 \\ n = 1 \\ (14.0-14.0) \end{array} $	$ 11.4 \\ n = 2 \\ (11.0-11.8) $	10.0 n = 2 (9.7-10.2)	14.0 n = 2 (12.8-15.2)
Papio sp. indet. Swartkrans II	8.2 n = 1 (8.2-8.2)	7.0 n = 1 (7.0-7.0)	9.2 n = 1 (9.2–9.2)	8.5 n = 1 (8.5-8.5)	9.8 n = 1 (9.8–9.8)	10.0 n = 1 (10.0-10.0)	10.2 n = 1 (10.2-10.2)	12.5 n = 1 (12.5-12.5)	10.9 n = 1 (10.9-10.9)	9.9 n = 1 (9.9–9.9)	
Papio sp. indet. Sterkfontein Member 4/4?	9.4 n = 5 (8.7-10.5)	7.4 n = 5 (6.4-8.5)	$ \begin{array}{r} 10.9 \\ n = 4 \\ (10.6-11.6) \end{array} $	$ \begin{array}{r} 10.4 \\ n = 4 \\ (10.0-10.9) \end{array} $	$ \begin{array}{r} 10.5 \\ n = 5 \\ (10.2 - 10.8) \end{array} $	$ 12.8 \\ n = 6 \\ (11.8-13.4) $	$ 12.1 \\ n = 5 \\ (11.6-12.6) $	12.9 n = 6 (11.8–13.5)		$ \begin{array}{r} 11.2 \\ n = 4 \\ (9.7-11.9) \end{array} $	
Papio hamadryas botswanae	8.2 n = 1 (8.2-8.2)	6.7 n = 1 (6.7-6.7)	9.5 n = 1 (9.5–9.5)	9.4 n = 1 (9.4–9.4)	10.5 n = 1 (10.5-10.5)	12.5 n = 1 (12.5-12.5)		$ \begin{array}{r} 12.1 \\ n = 1 \\ (12.1 - 12.1) \end{array} $	$ 12.2 \\ n = 1 \\ (12.2-12.2) $	$ \begin{array}{r} 10.1 \\ n = 1 \\ (10.1 - 10.1) \end{array} $	
Papio hamadryas ursinus	8.7 n = 127 (7.0-10.3)	7.9 n = 127 (6.1–9.8)	10.0 n = 152 (8.9-12.0)	9.3 n = 152 (8.0-11.0)	$ 11.2 \\ n = 154 \\ (8.5-13.1) $	11.9 n = 141 (9.5-13.7)	$ 11.0 \\ n = 139 \\ (9.0-12.9) $	13.6 n = 141 (10.7-15.5)	$ 12.4 \\ n = 120 \\ (10.3-14.2) $	10.7 n = 120 (8.2-13.1)	$ 14.1 \\ n = 122 \\ (11.4-16.4) $

 Table 1

 Summary statistics for the upper dentition of extant and fossil Plio-Pleistocene Papio species in South Africa.

Notes: For each taxon, the mean (top), sample size (middle), and range (bottom) are given for each tooth position. For individual specimen measurements, see SOM Tables S1-S4.

Taxon	P ₄ BL	P ₄ MD	M ₁ MBL	M ₁ DBL	M ₁ MD	M ₂ MBL	M ₂ DBL	M ₂ MD	M ₃ MBL	M ₃ DBL	$M_3 MD$
	_	_	_	_	_	_	10.2	12.6	11.8	_	15.7
?Papio izodi	_	-	-	-	-	-	n = 1	n = 2	n = 1	-	n = 2
-	-	_	-	-	_	_	(10.2–10.2)	(11.8–13.4)	(11.8–11.8)	_	(15.5–15.8
	6.7	7.8	8.1	8.2	9.8	9.9	9.6	11.8	10.4	9.1	14.8
Papio hamadryas angusticeps	<i>n</i> = 31	n = 32	<i>n</i> = 34	<i>n</i> = 36	<i>n</i> = 36	<i>n</i> = 33	<i>n</i> = 32	<i>n</i> = 36			<i>n</i> = 32
	(4.7–7.6)	(6.4–9.4)	(6.7–9.6)	(6.6–9.6)	(7.8–12.3)	(8.0-11.8)	(7.6–10.7)	(9.1–13.5)	(9.0–12.6)	(8.0–10.8)	(13.1–19.2
	7.5	9.0	8.3	8.6	10.3	10.6	10.1	12.8	11.0	9.5	16.7
Papio robinsoni	n = 26	n = 26	n = 14	<i>n</i> = 15	n = 18	n = 22	<i>n</i> = 23	n = 27			n = 24
	(6.0 - 9.0)	(7.4 - 10.4)	(7.1–9.3)	(7.7 - 9.7)	(8.2–12.6)	(8.8–12.1)	(8.3–11.2)	(11.1–15.5)	$\begin{array}{cccc} 11.8 & - \\ n = 1 & - \\ (11.8-11.8) & - \\ 10.4 & 9.1 \\ n = 34 & n = 30 \\ (9.0-12.6) & (8.0-10.8) \end{array}$	(13.6–18.8	
Papio cf. robinsoni	7.1	9.6	_	_	_	9.6	9.6	12.7	10.5	9.2	16.4
Swartkrans Member 2	n = 1	n = 1	-	-	-	n = 1	n = 1	n = 1			n = 2
Swartkrans Weinber 2	(7.1 - 7.1)	(9.6–9.6)	-	_	-	(9.6 - 9.6)	(9.6 - 9.6)	(12.7 - 12.7)	(10.0–11.1)	(9.1–9.2)	(15.5–17.3
Papio cf. robinsoni	7.6	8.9	8.3	8.6	11.2	10.4	9.7	12.3	10.3	9.5	15.3
Kromdraai B	n = 5	n = 5	n = 7	n = 7	n = 7	n = 6	n = 5	n = 7	n = 2	n = 3	<i>n</i> = 3
KIOIIIdiaa b	(7.0-8.1)	(8.5–9.4)	(7.4-8.8)	(7.9–9.5)	(10.5-12.5)	(9.4–11.5)	(8.8–10.5)	(10.0-14.7)	(10.1–10.5)	(10.1–10.5) (8.6–10.0)	(15.2–15.5
Papionini gen. et sp. indet.	7.7	9.6	8.5	9.1	10.9	10.7	10.4	13.6	10.3	9.2	16.2
Swartkrans Member 3	n = 4	n = 4	n = 2	n = 2	n = 2	n = 2	n = 2	n = 2	n = 6	n = 6	n = 6
Swaltklans Member 5	(6.8-8.1)	(7.7–10.3)	(8.0–9.1)	(8.5–9.7)	(9.6-12.2)	(9.3–12.1)	(8.9–11.9)	(12.0-15.2)	(9.1-11.8)	$\begin{array}{c} -\\ -\\ -\\ -\\ -\\ -\\ -\\ -\\ -\\ -\\ -\\ -\\ -\\ $	(14.3–18.9
Denianini sen et en indet	7.4	9.9	8.2	8.0	11.2	9.2	8.8	13.3	_	_	_
Papionini gen. et sp. indet. Cooper's D	n = 2	n = 2	<i>n</i> = 3	<i>n</i> = 3	<i>n</i> = 3	n = 2	n = 2	n = 2	_	_	_
cooper's D	(7.1–7.7)	(9.5-10.3)	(7.5–9.4)	(7.2–9.1)	(10.5-12.2)	(9.1–9.2)	(8.0–9.6)	(12.0-14.6)	-	-	-
	7.4	8.7	_	_	_	11.0	10.4	12.5	11.7	10.9	18.2
Papio sp. indet.	n = 1	n = 1	_	_	_	n = 1	n = 1	n = 1	n = 1	n = 1	n = 1
Swartkrans II	(7.4–7.4)	(8.7-8.7)	-	-	-	(11.0-11.0)	(10.4–10.4)	(12.5 - 12.5)	(11.7 - 11.7)	(10.9–10.9)	(18.2–18.2
Dents on indet	7.3	8.0	8.8	9.3	9.8	10.5	10.6	12.5	10.5	10.0	16.5
Papio sp. indet.	<i>n</i> = 3	n = 3	n = 2	n = 3	<i>n</i> = 3	n = 4	<i>n</i> = 3	n = 4	n = 2	n = 2	<i>n</i> = 3
Sterkfontein Member 4/4?	(6.7 - 7.6)	(7.3-8.9)	(8.5–9.1)	(9.2 - 9.4)	(9.0–10.3)	(10.0-11.0)	(9.6-11.2)	(11.4–13.2)	(10.1–10.9)	(9.4–10.5)	(15.9–17.
	-	_	7.3	7.9	9.6	9.7	9.4	11.4	9.4	8.1	13.6
Papio hamadryas botswanae	_	_	n = 1	n = 1	n = 1	n = 1	n = 1	n = 1	n = 1	n = 1	n = 1
• •	_	_	(7.3–7.3)	(7.9–7.9)	(9.6–9.6)	(9.7–9.7)	(9.4–9.4)	(11.4–11.4)	(9.4–9.4)	(8.1-8.1)	(13.6–13.0
	7.0	8.9	8.2	8.6	10.7	10.4	9.9	13.1	11.3	10.2	16.9
Papio hamadryas ursinus	<i>n</i> = 127	<i>n</i> = 127	<i>n</i> = 151	n = 152	<i>n</i> = 154	n = 140	<i>n</i> = 138	n = 140	n = 122	n = 121	<i>n</i> = 123
	(5.7 - 8.0)	(7.0 - 10.6)	(6.4 - 9.4)	(6.8 - 9.9)	(8.0 - 12.4)	(8.7 - 12.5)	(7.9 - 11.6)	(10.1 - 15.3)	(9.2 - 13.0)	(8.1 - 11.6)	(14.3–19.

Notes: For each taxon, the mean (top), sample size (middle), and range (bottom) are given for each tooth position. For individual specimen measurements, see SOM Tables S1-S4.

Table 3
Fossil specimens included in a 3D Geometric Morphometric Analysis of Papio cranial shape with DFA results.

Taxon	Institution	Specimen#	Sex	Site	DFA result
P. h. angusticeps	UW-AD	GV 4040	Female	Gladysvale	Papio (100%)
P. h. angusticeps	TMP	KA 194	Female	Kromdraai A	Papio (63%) Parapapio (37%)
P. h. angusticeps	TMP	CO 135a	Female	Coopers A	Papio (99%) Mandrillus (1%)
P. h. angusticeps	TMP	CO 101	Female	Coopers A	Papio (92%) Dinopithecus (6%) Mandrillus (2%)
P. h. angusticeps	UCMP	56767	Female	Bolt's Farm Pit 6	Papio (54%) Theropithecus (46%)
P. h. angusticeps	TMP	CO 100	Male	Coopers A	_
P. h. angusticeps	TMP	HGD 600	Male	Haasgat	_
P. h. angusticeps	TMP	HGD 606	Male	Haasgat	_
P. h. botswanae	BNMM	FC 346	Male	!Ncumtsa	_
P. robinsoni	TMP	SK 562	Female	Swartkrans Mb. 1	Papio (100%)
P. robinsoni	UCMP	56797	Female	Bolt's Farm Pit 23	Papio (84%) Theropithecus (16%)
P. robinsoni	TMP	SK 560	Male	Swartkrans Mb. 1	_
P. robinsoni	TMP	SK 555	Male	Swartkrans Mb. 1	_
P. robinsoni	BPI	M3147	Unknown	Skurweberg	Papio (100%)
?Papio izodi	UW-AD	TP 11	Female	Taung	Procercocebus (62%) Papio (36%) Parapapio (1%) Theropithecus (1%)
?Papio izodi	UW-AD	TP 10	Female	Taung	Papio (71%) Parapapio (22%) Procercocebus (6%)
?Papio izodi	UW-AD	TP 4	Female	Taung	Papio (51%) Procercocebus (43%) Parapapio (5%)
?Papio izodi	TMP	STS 262	Female	Taung	Soromandrillus (60%) Papio (25%) Macaca (14%)
?Papio izodi	TMP	T 13	Female	Taung	Soromandrillus (100%)
?Papio izodi	UW-AD	TP 12	Male	Taung	_
?Papio izodi	UW-AD	T89-11-1	Male	Taung	_
?Papio izodi	TMP	T 10	Male	Taung	_
?Papio izodi	UWMA	SWP 29	Female	Sterkfontein Member 4	_
Papio sp. indet	TMP	SK II 25	Female	Swartkrans II	_

Notes: DFA Result column indicates taxon that the individual specimen was classified to along with the posterior probability of that classification in parentheses. - indicates that the specimen could not be included in the DFA analysis due to its incompleteness.

Table 4

Extant sample included in a 3D Geometric Morphometric Analysis of *Papio* cranial shape.

Taxon	Females	Males	Both
Papio hamadryas anubis	59	124	183
Papio hamadryas cynocephalus	9	25	34
Papio hamadryas hamadryas	4	30	34
Papio hamadryas kindae	22	19	41
Papio hamadryas papio	1	16	17
Papio hamadryas ursinus	82	116	198
Cercocebus agilis	9	10	19
Cercocebus atys	3	3	6
Cercocebus torquatus	11	20	31
Lophocebus albigena	20	30	50
Mandrillus sphinx	14	27	41
Mandrillus leucophaeus	18	36	54
Macaca sylvanus	13	15	28
Theropithecus gelada	13	29	42
Totals	278	500	778

closely as possible for ease of reference, where the different sites are organized by the species present. In each case, we briefly note the most important specimens upon which our species identifications for that site are based (see also Figs. 2–4; SOM Figs. S1–S4).

<u>3.1.1. Sterkfontein Member 2</u> Although still formally unpublished, *?P. izodi* has been previously recognized at Sterkfontein Member 2 by Pickering et al. (2004) and Heaton (2006). We had the opportunity to study some of the Member 2 material in 2009

(CCG, access courtesy of R. Clarke) and 2012 (CCG and SRF, courtesy of R. Clarke), and we confirm this identification here. Among the cercopithecoid material from Member 2 are at least four partial crania that unambiguously document the presence of *?P. izodi* (Table 8; SOM Fig. S1; Heaton, 2006: Figs. 4–7). These partial crania, particularly the females, are so similar to the type series from Taung as to leave no doubt to their taxonomic affinity (SOM Fig. S1). Thus, these crania display relatively large orbits, relatively large teeth, moderate to absent maxillary ridges, moderately-developed to absent maxillary fossae, and a relatively shorter and broader rostrum compared to other *Papio* species. In addition, the anteorbital drop in these specimens is variably developed. A more detailed analysis of these specimens awaits their formal description and introduction into the literature.

<u>3.1.2. Sterkfontein Member 4</u> *?P. izodi* has previously been documented at Sterkfontein Member 4 by Eisenhart (1974), Delson (1984, 1988), and McKee (1993) in the form of a subadult female specimen, STS 262. Because fossils deriving from the Sterkfontein Type Site (STS) can be of uncertain provenience (although generally assumed to be derived from Member 4), it is also important to note that we recognize at least one more secure specimen, SWP 29a+b, a subadult partial face and mandible of a female from Member 4, as *?P. izodi* on the basis of its apparent anteorbital drop, relatively large orbits, relatively large teeth, and relatively short, rounded rostrum with no maxillary ridges and weakly excavated fossae.

Sample sizes for extant taxa included in cladistic analysis.

Taxon	Quantitative character sample size (Mean, Median, Mode, Range)	Qualitative character sample size (Mean, Median, Mode, Range)
Allenopithecus nigroviridis Males	4.8, 5, 5, 0–5	5.9, 6, 6, 0–7
Allenopithecus nigroviridis Females	4.8, 5, 5, 0-7	6.4, 7, 7, 0–7
Cercocebus agilis Males	10.0, 8, 0, 0–23	14.2, 16, 16, 0–16
Cercocebus agilis Females	6.6, 8, 3, 0–13	11.5, 13, 13, 0–13
Cercocebus atys Males	5.6, 4, 11, 0–12	1.9, 2, 2, 0–4
Cercocebus atys Females	9.1, 5, 0, 0–21	1.8, 2, 2, 0–2
Cercocebus chrysogaster Males	3.3, 0, 0, 0–9	5.6, 6, 6, 0–6
Cercocebus chrysogaster Females	2.3, 0, 0, 0-6	3.6, 4, 4, 0-4
Cercocebus torquatus Males	16.5, 20, 20, 0–32	30.9, 32, 33, 0–35
Cercocebus torquatus Females	13.1, 16, 18, 0–18	12.9, 14, 15, 0–16
Lophocebus albigena Males	18.1, 18, 18, 0-32	29.5, 31, 32, 0–33
Lophocebus albigena Females	15.6, 17, 17, 0–36	30.1, 35, 36, 0–36
Lophocebus aterrimus Males	1.5, 1, 1, 0–29	26.9, 29, 29, 0–29
Lophocebus aterrimus Females	2.2, 2, 2, 0-22	19.7, 22, 22, 0–22
Macaca sp. Males	20.5, 19, 20, 6-79	72.5, 78, 79, 0-80
Macaca sp. Females	19.0, 18, 19, 0–73	60.3, 68, 73, 0-74
Mandrillus leucophaeus Males	24.1, 25.5, 31, 0-35	13.6, 15, 15, 0–17
Mandrillus leucophaeus Females	14.6, 17, 18, 0–20	17.5, 20, 20, 0–21
Mandrillus sphinx Males	6.7, 6, 6, 4–18	11.4, 12, 12, 0–14
Mandrillus sphinx Females	2.9, 2, 2, 0–15	7.9, 9, 9, 0–10
Papio hamadryas anubis Males	9.0, 8, 7, 0–39	33.8, 36, 39, 0–39
Papio hamadryas anubis Females	8.9, 10, 11, 2–17	13.5, 16, 17, 0–17
Papio hamadryas cynocephalus Males	11.0, 13, 13, 0–17	13.7, 15, 15, 0–15
Papio hamadryas cynocephalus Females	6.9, 8, 9, 0–11	5.2, 6, 6, 0–6
Papio hamadryas hamadryas Males	1.2, 1, 0, 0–13	12.1, 13, 13, 0–13
Papio hamadryas hamadryas Females	0.5, 0, 0, 0–2	1.4, 2, 2, 0–2
Papio hamadryas kindae Males	4.3, 6, 0, 0–13	11.1, 13, 13, 0–13
Papio hamadryas kindae Females	4.0, 6, 0, 0–17	13.8, 15, 17, 0–17
Papio hamadryas papio Males	1.6, 1, 1, 0–10	9.4, 10, 10, 0–10
Papio hamadryas papio Females	0.0, 0, 0, 0–1	0.9, 1, 1, 0–1
Papio hamadryas ursinus Males	3.1, 4, 0, 0–36	32, 34.5, 36, 0–37
Papio hamadryas ursinus Females	2.1, 3, 3, 0–11	8.1, 9, 11, 0–11
Theropithecus gelada Males	17.7, 19, 21, 0–22	16.6, 17, 17, 0–17
Theropithecus gelada Females	16.9, 20, 21, 0–22	5.4, 6, 6, 0–6
Males average	9, 9, 9, 1–25	20, 22, 22, 0–23
Females average	8, 8, 8, 0–18	13, 15, 15, 0–16

Notes: Sample sizes are listed for specimens identifiable to sex used in character analysis. For each taxon, measurements and qualitative character state assignments were made, supplemented with additional data from the major sources listed in Tables 6–8. For a full list of characters and character states, see Gilbert (2013), Gilbert et al. (2016a,b), and Table 7.

A limited number of specimens assumed to be from Member 4 are consistent with more derived Papio taxa such as P. robinsoni and P. h. angusticeps, but require more complete and diagnostic material to be certain. SWP 1290 and SWP 1230 are maxillae displaying definitive maxillary fossae inconsistent with Parapapio and ?P. izodi. The coordinates given for each of these specimens correspond to Member 4 according to Kuman and Clarke (2000), but the UW-AD catalog lists them as from Member 5, leading to an uncertain provenience. STS 387a is a right maxillary fragment with a clear and well-defined maxillary fossa, also consistent with derived Papio taxa, but neither Parapapio nor typical ?P. izodi. Tooth length is also within the range of *P. robinsoni* and *P. h. angusticeps* (see boxplots Figs. 5–9). STS 358 is a large male partial mandible, highly crushed and distorted. It appears to have a shallow, but clear, corpus fossa, again more consistent with derived Papio taxa than Parapapio or ?P. izodi. Because these last two specimens derive from the old STS collections, their exact provenience is not entirely secure.

Two specimens are more securely placed in Member 4 but are less diagnostic as to (sub)species. SWP 31 (STW 31) is a female partial face from Member 4 that has been previously identified as *P. robinsoni* (SOM Fig. S4; see also Eisenhart, 1974: Figs. 323-324; Delson, 1984, 1988). The right maxilla shows a moderately excavated but well-defined maxillary fossa, beginning at about M¹ and extending posteriorly to the zygoma, which it invades very slightly. The fossa is not as deep as those of some *P. robinsoni* and *P. h. angusticeps* specimens, but there is considerable distortion due to superioinferior plastic deformation. An anteorbital drop appears to

be present as well, but it is hard to be certain given distortion. The teeth of SWP 31 are large and the malar height is taller than those of female *?P. izodi* specimens, closer to that seen in females of extant *Papio*. This specimen is perhaps most similar to *P. robinsoni* females. Although larger than most, it is, however, within the range for females of *?P. izodi* and *P. h. angusticeps* as well.

SWP 35 (STW 35) is a small fragment of a left maxilla from Member 4 preserving a P^4 and a small area of the lateral wall of the rostrum (SOM Fig. S4; see also Eisenhart, 1974: Figs 321–322). It shows a clear fossa over the M¹ partial alveolus that is more extensive than any observed in *?P. izodi* or *Parapapio* (SOM Fig. S4). Furthermore, the P⁴ is considerably larger than the known *?P. izodi* range (see measurements in SOM Table S3). Given its fragmentary nature, we conservatively assign this fossil to *Papio* sp. indet., along with the other specimens listed above.

It should also be noted that at least three Member 4 specimens (SWP 1180, SWP 1217, and SWP 1348) previously cataloged as *P. robinsoni* are more properly identified as Papionini gen. et sp. indet. due to their lack of diagnostic morphology (e.g., no facial fossae, maxillary ridges, or anteorbital drop is preserved; therefore, they could be either *Papio* or *Parapapio*).

In summary, at Sterkfontein Member 4, *?P. izodi* is clearly present. In addition, there is a second more derived *Papio* taxon present, but it is unclear whether this is *P. robinsoni*, *P. h. angusticeps*, or another derived form. While two of these specimens are known to be from Member 4, others of these more derived specimens could be from either Member 4 or Member 5.

Sample for fossil taxa included in cladistic analysis.

Fossil Taxon/OTU	Sample Size (Males, Females)	Key specimens	Source
Dinopithecus ingens	(4, 7)	SB 7, SK 401, SK 542a, SK 546, SK 548, SK 553, SK 554, SK 574, SK 599, SK 600, SK 603	Freedman, 1957; Gilbert, 2013; this study
Gorgopithecus major	(6, 4)	KA 150, KA 153, KA 154, KA 192, KA 524/676, KA 605, KA 944, KA 1148, SK 604a, OLD 1962/S.196, FLK NNI 1011	Freedman, 1957; Gilbert et al., 2016a; this study
Lophocebus sp. nov.	(6, 8)	KNM-ER 594, KNM-ER 595, KNM-ER 827, KNM-ER 898, KNM-ER 965, KNM-ER 1661, KNM-ER 3090, KNM-ER 6014, KNM-ER 6063, KNM-ER 18922, KNM-ER 40476, KNM-ER 44260, KNM-ER 44262, KNM-ER 44317	Jablonski et al., 2008; Gilbert, 2013; this study
Papio hamadryas angusticeps	(9, 9)	CO 100, CO 101, CO 102, CO 115/103, CO 134B/D, CO 135A, KA 156, KA 161, KA 165, KA 166, KA 167C, KA 168, KA 188, KA 194, KB 94, GV 4040, UW 88-886, UCMP 56767	Freedman, 1957; McKee and Keyser, 1994; Gilbert et al., 2015; this study
?Papio izodi	(3, 9)	SAM 11728, SWP Un 2, T13, T89-11-1, TP4/M681/AD946, TP7/M684/AD992, TP10, TP11, TP12, UCMP 125854, UCMP 125855, UCMP 125856	Freedman, 1957; Freedman, 1961; Freedman, 1965; Heaton, 2006; Gilbert, 2013; this study
Papio robinsoni	(8, 15)	SK 406, SK 407, SK 408, SK 409, SK 416, SK 421, SK 549, SK 555, SK 557, SK 558, SK 560, SK 562, SK 565, SK 566, SK 571B, SK 602, SK 3211B, SB 2, M3147, UCMP 56797, UCMP 56786, BF 38	Freedman, 1957; Freedman, 1965; Freedman and Brain, 1977; this study
Soromandrillus quadratirostris	(4, 5)	NME-USNO, NME Omo 47-1970-2008, NME L 4-13b NME Omo 42-1972-1, NME Omo L-185-6, NME Omo 75N (71)-C2, DGUNL LEBA02, DGUNL LEBA03, DGUNL LEBA06	Iwamoto, 1982; Eck, 1977; Eck and Jablonski, 1984; Delson and Dean, 1993; Gilbert et al., 2009b; Gilbert, 2013; this study
Parapapio ado	(2,4)	BMNH M14940, EP 1579/98, LAET 74-242/243/244, LAET 75-483, LAET 75-1209, LAET 78-5269, MB MA 42444/42445/42458	Leakey and Delson, 1987; Harrison, 2011; this study
STS 254, STS 255, STS STS 337, STS 338, STS		M202/MP2, M211/MP11, M2961, M2962/MP76, M2978/MP92, M3037, M3067, STS 254, STS 255, STS 258, STS 264, STS 267, STS 297, STS 331, STS 332, STS 335, STS 337, STS 338, STS 339, STS 360, STS 363, STS 378A, STS 379, STS 390A, STS 393, STS 396A, STS 397, STS 409, STS 411A/B, STS 469, STS 534, STS 542, STS 562, STS 564, BF 43	Freedman, 1957; Freedman, 1960; Maier, 1970; Freedman and Stenhouse, 1972; Eisenhart, 1974; Gilbert, 2013; this study
Parapapio jonesi	(4, 12)	AL 363-15, AL 363-1, M215/MP15, M218/MP18, M3051/MP 165, STS 250, STS 284, STS 313, STS 355, STS 372, STS 547, STS 565, SWP (STW) 27, SWP 389, SWP 1728, SWP 2947	Freedman, 1957; Freedman, 1960; Maier, 1970; Freedman and Stenhouse, 1972; Eisenhart, 1974; Freedman, 1976; Frost and Delson, 2002; Gilbert, 2013; this study
"Parapapio" lothagamensis	(5, 2)	KNM-LT 419, KNM-LT 448, KNM-LT 449, KNM-LT 23065, KNM-LT 23091, KNM- LT 24111, KNM-LT 24136	Leakey et al., 2003; this study
Parapapio whitei	(5, 5)	M3072, MP221, MP223, STS 259, STS 266, STS 352, STS 359, STS 374, STS 389, STS 563	Freedman, 1957; Maier, 1970; Freedman, 1976; Freedman and Stenhouse, 1972; Eisenhart, 1974; Gilbert, 2013; this study
Pliopapio alemui	(4, 6)	AME-VP-1/64, ARA-VP 1007, ARA-VP 1723, ARA-VP 1/73, ARA-VP 1/133, ARA- VP 1/563, ARA-VP 1/1006, ARA-VP 1/2553, ARA-VP 6/437, ARA-VP 6/933	Frost, 2001b; Frost et al., 2009; Gilbert, 2013; this study
Procercocebus antiquus	(5, 14)	SAM 4850, SAM 5356, SAM 5364, M3078, M3079, T11, T14, T17, T18, T20, T25, T21, T89-154, T88-17, TP8, TP9, TP13, UCMP 56624, UCMP 56653, UCMP 56694, UCMP 56821/125956	Gilbert, 2007; Gilbert, 2013; this study
Theropithecus baringensis	(2, 0)	KNM-BC 2, KNM-BC 1647	Leakey, 1969; Delson and Dean, 1993; Eck and Jablonski, 1984; this study
Theropithecus brumpti (12, 6) NME L17-45, NME 32-154, NME L32-155, NME L122-34, NL L345-3, NME L345-287, NME L576-8, KNM-TH 46700, K L345-3, NME L345-287, NME L576-8, KNM-TH 46700, K		NME L17-45, NME 32-154, NME L32-155, NME L122-34, NME L338Y-2257, NME L345-3, NME L345-287, NME L576-8, KNM-TH 46700, KNM-WT 16749, KNM- WT 16806, KNM-WT 16808, KNM-WT 16828, KNM-WT 16888, KNM-WT 17571, KNM-WT 17555, KNM-WT 17560, KNM-WT 39368CX	Eck and Jablonski, 1987; Harris et al., 1988; Jablonski et al., 2008; Gilbert et al., 2011b; this study
Theropithecus oswaldi darti			Eck, 1993; Eck and Jablonski, 1987; Freedman, 1957; Maier, 1970; Maier, 1972; this study
Victoriapithecus macinnesi	(4, 2)	KNM-MB 18993, KNM-MB 21027, KNM-MB 27876, KNM-MB 29100, KNM-MB 29158, KNM-MB 31281	Benefit, 1987; Benefit, 1993; Benefit and McCrossin, 1991; Benefit and McCrossin, 1997; this study

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Notes: Sample sizes are listed for key specimens, identifiable to sex, used in character analysis. Measurements and character state assignments were made and supplemented with additional data from the major sources listed for each taxon.

Postcranial characters added to the Gilbert	(2013) morphological character matrix.
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	Character description	Reference
A1	Vertebral Number: quantitative; modal number of thoracic and lumbar vertebrae	Williams (2012)
S1	Scapular Index : quantitative; Max. height \times 100/Max. length	Fleagle and McGraw (2002)
H1	Deltoid Plane Width: quantitative; deltoid plane width x 100/humeral head width	Fleagle and McGraw (2002)
H2	Brachialis Flange Width: quantitative; Max width of the brachialis flange \times 100/Geometric Mean	Gilbert et al. (2016a)
H3	Supinator Crest Length : qualitative; supinator crest height; 0 = not extended, 1 = polymorphic, 2 = extended	Fleagle and McGraw (2002)
H4	Distal Articulation Breadth: quantitative; anterior articular width x 100/biepicondylar width	Fleagle and McGraw (2002)
H5	Medial Trochlear Lip : qualitative; 0 = normal, 1 = polymorphic, 2 = prominent	Fleagle and McGraw (2002)
H6	Olecranon Fossa Width: quantitative; olecranon fossa breadth × 100/Geometric Mean	Gilbert et al. (2016a)
H7	Lateral Margin of Olecranon Fossa: qualitative; $0 = normal$, $1 = prominent$	Fleagle and McGraw (2002)
H8	Olecranon Fossa Depth: qualitative; 0 = normal, 1 = polymorphic, 2 = deep	Gilbert et al. (2016a)
H9	Olecranon Fossa Shape: qualitative; 0 = rounded, 1 = polymorphic, 2 = triangular	Gilbert et al. (2016a)
H10	Distal AP Depth : quantitative; max AP depth of distal articulation × 100/Gmean	Gilbert et al. (2016a)
H11	Medial Epicondyle Orientation : qualitative; $0 = \langle 45^{\circ}, 1 = 45{-}60^{\circ}, 2 = \rangle 60^{\circ}$	Harrison (1989) Frost and Delson (2002)
U1	Coronoid Width : quantitative; coronoid width \times 100/articular notch width	Fleagle and McGraw (2002) Harrison (1989) Guthrie (2011)
U2	Interosseous Crest: qualitative; 0 = strong, 1 = weak	Fleagle and McGraw (2002)
R1	Radial Shaft Shape : qualitative; $0 =$ triangular, $1 =$ rounded	Fleagle and McGraw (2002)
MC1	Relative MC I Length : quantitative; $0 = $ short, $1 = $ intermediate, $2 = $ long compared to MC II-V	Harrison (1989) Frost and Delson (2002) Guthrie (2011)
I1	llium Breadth : quantitative; ilium min. width \times 100/max. acetabular diameter	Fleagle and McGraw (2002)
I2	Gluteal Tuberosity : qualitative; 0 = prominent, 1 = weak	Fleagle and McGraw (2002)
F1	Patellar Groove, Medial Lip : qualitative; 0 = weak/round, 1 = moderate	Fleagle and McGraw (2002)
F2	Patellar Groove, Lateral Lip : qualitative; 0 = weak, 1 = moderate, 2 = strong/sharp	Fleagle and McGraw (2002)
T1	Tibial Compression : quantitative; mid AP diameter × 100/mid ML diameter	Fleagle and McGraw (2002)

Notes: Samples for extant taxa as given in the references listed. For any given character where character states were consistent among extant species within sampled genera, the genus average was used for any unsampled extant species within a given genus. For available postcranial characters, *Allenopithecus* was scored on the basis of 3 males (1 adult, 2 subadults with most epiphyses fused) and 1 female. Fossil taxa sampled include adult *T. darti, T. brumpti, Pp. jonesi, Pr. antiquus, P. izodi, Pp. ado, "Pp". lothagamensis,* and *Victoriapithecus*. All postcranial characters were treated as ordered.

<u>3.1.3. Taung</u> The type series of *?P. izodi* documents this taxon at Taung. The best specimens are listed in Table 8.

<u>3.1.4. Malapa</u> UW-88-886 establishes *P. h. angusticeps* as the only non-hominin primate at the site, with a date range of ca. 2.4–2.0 Ma (see Gilbert et al., 2015).

<u>3.1.5. Haasgat</u> McKee and Keyser (1994) and Adams (2012) recognized *P. h. angusticeps* at Haasgat, as we do here. Many specimens show the diagnostic morphology of *P. h. angusticeps*, including deep maxillary and mandibular corpus fossae, marked maxillary ridges, and rostra that are narrower and less flattened than in most *Papio* (see also SOM Fig. S2). In addition, they are also smaller, on average, than *P. robinsoni*. See McKee and Keyser (1994: Figs. 2–4) and Adams (2012: Fig. 3) for more complete descriptions. The best specimens exhibiting clear and diagnostic morphology are listed in Table 8.

<u>3.1.6. Kromdraai A</u> Previous analyses have suggested that both *P. robinsoni* and *P. h. angusticeps* are found at Kromdraai A (Freedman, 1957; Delson, 1984, 1988; McKee et al., 1995; Heaton, 2006). *P. h. angusticeps* is clearly established by the type specimen KA 194, a partial female cranium (Broom, 1940; SOM Fig. S2). However, while studying the entire Kromdraai A sample in 2012, we noticed that there are no specimens preserving diagnostic features unique to *P. robinsoni* and/or exclusive of *P. h. angusticeps*. The small number of *P. robinsoni* specimens previously recognized by Freedman (1957) and Delson (1984,

1988) were diagnosed based on their general *Papio* features and large size compared to the rest of the sample. In fact, at all sites where *P. h. angusticeps* and *P. robinsoni* have been previously argued to co-occur, we were struck by the observation that only one of the two taxa is ever represented by relatively complete cranial material, while the other is typically recognized on the basis of fragmentary fossils judged to be too large (*P. robinsoni*) or too small (*P. h. angusticeps*) to fit with the better preserved specimens. In the absence of more definitive evidence, we recognize only one variable *Papio* species at Kromdraai A, *P. h. angusticeps* (see also metric analysis below, Figs. 5–9).

<u>3.1.7. Cooper's A</u> This sample includes Cooper's A-E in the old collection at the Ditsong National Museum, and these are all equivalent to Cooper's A in the new system (C. Steininger, pers. comm.). The bulk of this material was collected from dump heaps at a limestone mine, making exact provenience uncertain (Folinsbee and Reisz, 2013; C. Steininger, pers. comm.). In any case, the predominant papionin taxon present among the Cooper's A collection is *P. h. angusticeps*, which is represented by a number of good cranial specimens preserving features such as well-defined and deep maxillary fossae, well-defined maxillary ridges, and small premolars (Table 8; Figs. 2–3; SOM Fig. S2). While both Freedman (1957) and Delson (1984, 1988) again recognized some fragmentary specimens at Cooper's A as *P. robinsoni*, we feel they are all comfortably accommodated within a single population

Diagnostic specimens establishing the presence of *Papio* taxa by site.

Taxon	Site	Diagnostic/Best specimen(s)	Brief description(s)	Reference
	Sterkfontein	SWP 2948 SWP 2953	Subadult female partial cranium Partial cranium missing most of rostrum; sex unknown	Heaton (2006) Heaton (2006)
	Member 2	SWP Uncatalogued SWP 2946	Male partial cranium Female partial cranium	Heaton (2006) Heaton (2006)
	Sterkfontein Member 4	SWP 29a+b STS 262	Subadult female partial face and mandible Subadult female partial cranium	Heaton (2006) Eisenhart (1974); Delson (1984, 1988); McKee (1993)
		UW-AD 922/TP-7	Female partial cranium with dentition (TYPE)	Gear (1926); Jones (1937); Freedman (1957)
PP. izodi		SAM 11728 SAM 5357	Female partial cranium with dentition Cranial endocast with partial mandible and dentition	Freedman (1961) Freedman (1957)
	Taung	TP-10/UCMP 56605 TP-4/AD 946 TP-12/UCMP 56652	Female partial cranium with dentition Female partial face with mandible and dentition Male partial cranium with dentition	Freedman (1961) Freedman (1957) Freedman (1965)
		T89-11-1 UCMP 125856 T13	Male partial cranium with dentition Male partial cranium with dentition ?Female cranium with dentition	this study Delson (1988); Gilbert (2013) Freedman (1957)
		T10 UCMP 125854 UCMP 125855	Subadult cranium with dentition Female partial cranium with dentition Female partial cranium with dentition	Freedman (1957); this study Delson (1988); Gilbert (2013) Delson (1988); Gilbert (2013)
Papio cf. izodi	Gladysvale	GV 4340	Female partial cranium, no dentition	Berger et al. (1993); this study
	Malapa	UW-88-886	Male partial cranium	Gilbert et al. (2015)
		HGD 1249 HGD 600 HGD 605 HGD 603	Male partial right face with dention Male muzzle with dentition Partial maxilla with dentition Left male maxilla with dentition	McKee and Keyser (1994); Adams (2012 McKee and Keyser (1994); Adams (2012 McKee and Keyser (1994); Adams (2012 McKee and Keyser (1994); Adams (2012
	Haasgat	HGD 606 HGD 644 HGD 601 HGD 1246 HGD 1243 HGD 618	Male maxilla with dentition Male left maxilla with dentition Subadult female partial cranium with dentition Female partial mandible with dentition Female partial mandible with dentition Male partial mandible with dentition	McKee and Keyser (1994); Adams (2012 McKee and Keyser (1994); Adams (2012
		HGD 615 HGD 626	Female partial mandible with dentition Female partial mandible with dentition	McKee and Keyser (1994); Adams (2012 McKee and Keyser (1994); Adams (2012
Dh. sussisting	Kromdraai A	KA 194	Female partial cranium with dentition (TYPE)	Broom (1940); Freedman (1957)
P. h. angusticeps		CO 100	Partial male cranium with dentition	Freedman (1957)
	Cooper's A	CO 101 CO 103/115 CO 134B-D	Partial female cranium with dentition Male mandible with dentition Subadult male partial maxilla and mandible with dentition	Freedman (1957) Freedman (1957) Freedman (1957); this study
		CO 135 CO 102	Partial female cranium with dentition Partial muzzle with dentition	Freedman (1957) Freedman (1957)
	Gladysvale	GV 4040	Partial female cranium with dentition	this study
	Bolt's Farm Pit 6	UCMP 56767 UCMP 56766 UCMP 56768 UCMP 56772 UCMP 56769 UCMP 177639 UCMP 56771	Female partial cranium with dentition Male crushed muzzle with dentition Male partial mandible with dentition Male partial mandible with dentition ?Female partial mandible with dentition Female partial mandible with dentition	Freedman (1965); this study Freedman (1965); this study Freedman (1965); this study Freedman (1965); this study Freedman (1965); this study this study Freedman (1965); this study
		DN 2160 DN 541 DN 403 DN 528	Male partial cranium with dentition Male crushed face with dentition Male palate with dentition Male right face and muzzle with dentition	Adams et al. (2016); this study Adams et al. (2016); this study Adams et al. (2016); this study Adams et al. (2016); this study
P. robinsoni	Drimolen Main Quarry	DN 363 DN 2344 DN 2162 DN 2345 DN 1158 DN 1158	?Female crushed partial cranium with dentition Female partial crushed cranium with dentition Male mandibular corpus with dentition Female partial mandible with dentition Right mandibular fragment with dentition Left mandibular fragment with dentition	Adams et al. (2016); this study Adams et al. (2016); this study
		SK 555 SK 560 SK 602 SK 14006 SK 557 SK 558 SK 558	Male muzzle with dentition (TYPE) Male muzzle with dentition Male crushed face with dentition Crushed face with M3 Female partial cranium with dentition Female partial cranium with dentition	Freedman (1957) Freedman (1957) Freedman (1957) Freedman and Brain (1977) Freedman (1957) Freedman (1957) Freedman (1957)
	Swartkrans Member 1	SK 562 SK 566 SK 14083	Female maxilla and muzzle with dentition Female compressed muzzle with dentition Male left mandibular corpus and ramus with dentition	Freedman (1957) Freedman (1957) Freedman and Brain (1977)

(continued on next page)

Table 8 (continued)

Faxon	Site	Diagnostic/Best specimen(s)	Brief description(s)	Reference
		SK 406 SK 408 SK 407 SK 421 SK 571a SK 14156	Male partial mandible with dentition Male partial mandible with dentition Female mandibular corpus with dentition Female partial mandible with dentition Female partial mandible with dentition Female partial mandible with dentition	Freedman (1957) Freedman (1957) Freedman (1957) Freedman (1957) Freedman (1957) Freedman and Brain (1977)
	Skurweburg	SB 2 M3147	Crushed male partial cranium Partial muzzle with dentition	Freedman (1957) this study
	Bolt's Farm Pit 23	BF 38 UCMP 56797	Male muzzle and palate with dentition Female partial cranium with dentition	Freedman (1965); this study Freedman (1965); this study
	Swartkrans Member 2	SKX 814	Subadult female palate and partial muzzle with dentition	Delson (1988); this study
Papio cf. robinsoni	Sk 406Male partial mandible with dentitionSk 407Female partial mandible with dentitionSk 407Female partial mandible with dentitionSk 421Female partial mandible with dentitionSk 571aFemale partial mandible with dentitionSk 14156Female partial mandible with dentitionSkurweburgSB 2M3147Partial muzzle with dentitionBolt's Farm Pit 23BF 38UCMP 56797Female partial cranium with dentitionSwartkransSKX 814Member 2KB 1456Kromdraai BKB 1456Ka 196/KB 94Mandibular fragment with dentitionBolt's Farm Pit 10?UncataloguedPartial maxilla with dentitionSwartkrans IISK 1125Skartkrans IISK 1125Skartkrans IISK 1126SterkfonteinSWP 31SterkfonteinSTS 387aMaxillary fragment with dentitionMember 4 or 5SWP 1230SterkfonteinSTS 387aMakillary fragment with dentitionSterkfonteinSWP 1230Partial mandible with dentitionSterkfonteinSWP 1232Male partial mandible with dentitionSterkfonteinSWP 1232Female partial mandible with dentitionSterkfonteinSWP 1232Female partial mandible with dentitionSterkfonteinSWP 1232Female partial mandible with dentitionSterkfonteinSWP 1232SterkfonteinSWP 1232Male partial mandible with dentition<	Delson (1988); this study Delson (1988); this study Freedman (1957); this study Freedman (1957); this study		
	Bolt's Farm Pit 10?	Uncatalogued	Partial maxilla with dentition	this study
	Swartkrans II			Freedman (1957); this study Freedman (1957); this study
				Eisenhart (1974); this study Eisenhart (1974); this study
Papio sp. indet.	specimen(s) Specimen(s) Sk 406 Male partial mandible with dentition SK 408 Male partial mandible with dentition SK 407 Female mandibluar corpus with dentition SK 421 Female partial mandible with dentition SK 421 Female partial mandible with dentition SK 571a Female partial mandible with dentition Sk 14156 Female partial mandible with dentition Skurweburg SB 2 Crushed male partial mandible with dentition Bolt's Farm Pit 23 BF 38 UCMP 56797 Female partial renaium Swartkrans SKX 814 Member 2 Subadult female palate and partial muzzle with dentition Member 2 KB 1456 Subadult female palate with dentition KB 5230 Krondraai B KB 5230 KA 196/KB 94 Mandibular fragment with dentition Bolt's Farm Pit 107 Uncatalogued Partial maxilla with dentition SK K1 125 Sterkfontein SWP 31 Sterkfontein SWP 31 Sterkfontein SWP 31 Membe	Eisenhart (1974); this study Eisenhart (1974); this study this study Heaton (2006); this study		
SK 408Male partial mandible with de SK 407SK 407Female mandibular corpus with SK 421SK 421Female partial mandible with d SK 421SK 421Female partial mandible with d SK 571aSkurweburgSB 2 M3147Bolt's Farm Pit 23BF 38 UCMP 56797Bolt's Farm Pit 23BF 38 UCMP 56797Papio cf. robinsoniSwartkrans Member 2Swartkrans Member 2SKX 814 KB 1456Subadult female palate and partial rami dentitionPapio cf. robinsoniKromdraai B SwartkransPapio sp. indet.Swartkrans II Sterkfontein Member 4Sterkfontein Member 4SWP 31 SWP 1230Papio sp. indet.Sterkfontein Sterkfontein SWP 1230Sterkfontein Member 5SWP 1232 SWP 1232Male partial mandible with de Member 5Sterkfontein SWP 1012Female partial mandible with de Member 4Sterkfontein Member 5Sterkfontein SWP 1012Female partial mandible with de Member 5	Male partial mandible with dentition	this study		
		SWP 1012	Female partial mandible with dentition	this study

Notes: "Reference" column lists other references which discuss the specimen in question. If the specimen in question has been reassigned to a different taxon in this study, the original reference is given in addition to "this study". If the specimen in question was not directly mentioned by the original reference (only the taxon), the original reference is also given in addition to "this study". Finally, specimens formally assigned to a taxon for a first time in this study are listed as "this study".

(Figs. 5–9). There is no relatively complete cranial material that is diagnostically *P. robinsoni* at Cooper's A, in contrast to the material found at Swartkrans and Skurweberg.

<u>3.1.8. Gladysvale *P. h. angusticeps* is represented at Gladysvale by GV 4040, a female partial cranium with partial dentition. Interestingly, there is also a possible female partial cranium (GV 4340) with large orbits, slight maxillary fossae, a possible anteorbital drop, and slight to absent maxillary ridges; these features are most consistent with *?P. izodi* (see also Berger, 1993; Berger et al., 1993). However, the specimen lacks any dentition and is not complete enough to be certain (we consider it to be cf. *?P. izodi* here). Furthermore, the time averaging and geological complexity present between the various internal and external deposits at Gladysvale make it impossible to determine whether GV 4040 and GV 4340 come from the same level or whether they represent two different depositional events at the site (see also Berger and Tobias, 1994; Lacruz et al., 2002; Pickering et al., 2007).</u>

3.1.9. Bolt's Farm Pit 6 Following Freedman (1965), Delson (1984, 1988) previously recognized only *P. robinsoni* at Bolt's Farm Pit 6 and Pit 23. While we concur with this assessment of Pit 23, at Pit 6 we find the *Papio* material more consistent with the features diagnostic for *P. h. angusticeps* and we re-assign the *Papio* material present at this site to that subspecies. There are a number of good specimens of *P. h. angusticeps* present at Pit 6, including: UCMP 56767, a female cranium very similar to KA 194 with narrow muzzle, deep maxillary fossae invading the infraorbital plate, relatively small orbits, peaked nasals well above the muzzle dorsum, maxillae that do not meet in the midline,

and well-defined maxillary ridges (better developed than seen in P. robinsoni UCMP 56797 from Pit 23; see Figs. 2-4); UCMP 56766, a male crushed muzzle/rostrum with very deep and welldefined maxillary fossae invading the infraorbital plate, a muzzle which appears to have probably been narrower than that of *P. robinsoni*, and a P^4 that is not noticeably large (see Figs. 5, 7; SOM Tables S3–S4); UCMP 56768, a male partial mandible with P₄ qualitatively small, mandibular corpus fossae clearly present under P_4/M_1 contact and deep on left hand side; and UCMP 56772, a male partial mandible with partial dentition and very deep mandibular corpus fossa under P_4/M_1 on the left side. Other diagnostic specimens of P. h. angusticeps from Pit 6 are listed in Table 8. In addition, UCMP 56771 is a female complete mandibular dentition incompletely prepared from surrounding matrix and consistent in size with P. h. angusticeps but with no mandibular morphology preserved. Given that there is no evidence of dental variation beyond that expected in a single population of modern P. h. ursinus, we again recognize only one Papio taxon at Bolt's Farm Pit 6, P. h. angusticeps.

3.1.10. Drimolen Main Quarry *P. robinsoni* is reported at Drimolen Main Quarry by Keyser (2000) and subsequently confirmed by Adams et al. (2016) and our own observations here (access courtesy of C. Menter). While the specimens await formal description and publication, they display characteristic *P. robinsoni* features such as large size, a definitive anteorbital drop, a squared-off "boxy" rostrum, qualitatively large premolars, and weak-to-moderately developed maxillary and mandibular fossae (SOM Fig. S3). They seem to differ from the Swartkrans and Skurweberg

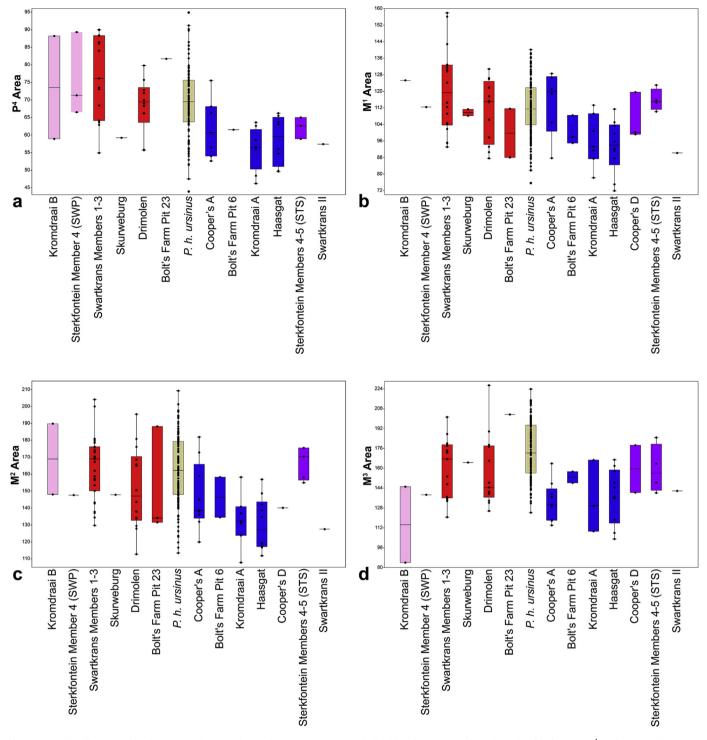


Figure 5. Boxplots illustrating dental size ranges (expressed as tooth area = maximum mesiodistal length x maximum buccolingual width) for upper P⁴s and upper molars among Plio-Pleistocene South African sites and a sample of extant *P. h. ursinus* specimens. Pink indicates those populations designated as *P. cf. robinsoni*, red indicates those populations designated as *P. nobinsoni*, blue indicates those populations designated as *P. nobinsoni*, blue indicates those populations designated as *P. nobinsoni*, blue indicates those populations designated as *P. nobinsoni*, see Tables 1 and 2 and SOM Tables S1–S4.

populations in that the maxillae do not seem to contact in the midline on the rostrum, but only two specimens are really assessable for this feature (DN 2160, 541) and so it is possible that this feature is present at Drimolen Main Quarry, just in lower frequency. The most definitive specimens are listed in Table 8.

with several other good specimens that help define the morphological features of the species (Figs. 2–4; SOM Fig. S3). The best specimens are listed in Table 8.

3.1.11. Swartkrans Member 1 The type specimen of *P. robinsoni* and muzzle of flattened muz

<u>3.1.12. Skurweberg</u> *P. robinsoni* is clearly documented at Skurweberg by the male crushed partial cranium SB 2, and the partial face and muzzle of unknown sex, M3147. Both specimens display a flattened muzzle dorsum, clear maxillary ridges (more rounded in

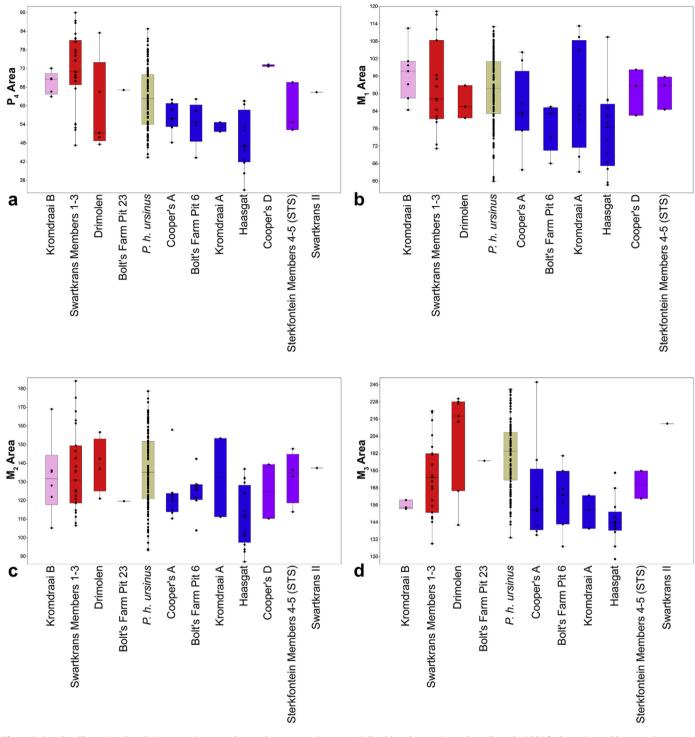


Figure 6. Boxplots illustrating dental size ranges (expressed as tooth area = maximum mesiodistal length x maximum buccolingual width) for lower P₄s and lower molars among Plio-Pleistocene South African sites and a sample of extant *P. h. ursinus* specimens. Pink indicates those populations designated as *P. cf. robinsoni*, red indicates those populations designated as *P. nobinsoni*, blue indicates those populations designated as *P. nobinsoni*, blue indicates those populations designated as *P. nobinsoni*, blue indicates those populations designated as *P. nobinsoni*, see Tables 1 and 2 and SOM Tables S1–S4.

SB 2, sharper in 3147), clear maxillary fossae that are not as deep as those of *P. h. angusticeps*, and maxillae that override the nasals and meet in the midline.

3.1.13. Bolt's Farm Pit 23 P. robinsoni is present at Bolt's Farm Pit 23

based on the presence of BF 38 (adult male muzzle and palate

preserving partial dentition) and UCMP 56797 (adult female partial

cranium with partial dentition), as previously noted by Freedman (1976) and Delson (1984, 1988) (Figs. 2–4).

<u>3.1.14. Swartkrans Member 2</u> A number of fragmentary specimens are present that are most consistent with *P. robinsoni*, but lack enough of the cranium and dentition to be certain. In particular, the large size and shallow-to-moderate maxillary fossa preserved in subadult female SKX 814 is a good match for SK 562 from

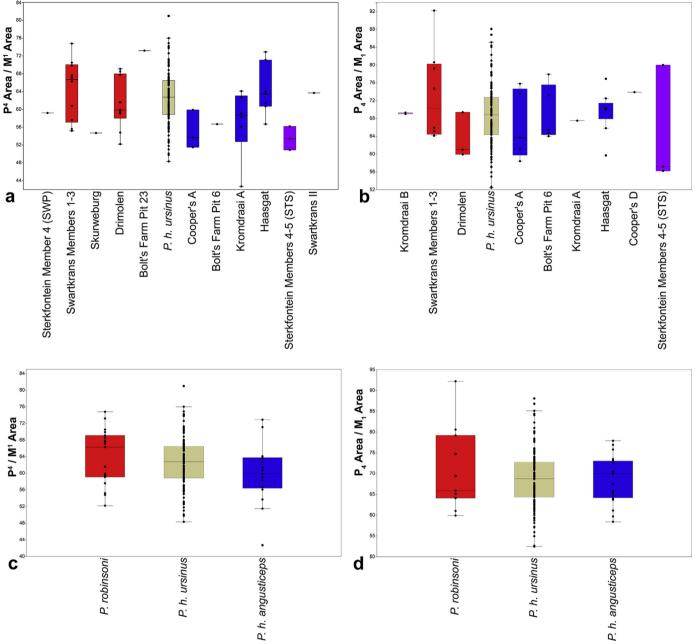


Figure 7. Boxplots illustrating dental size ranges for upper and lower P4/M1 ratios (expressed as [P4 area/M1 Area] x 100) among Plio-Pleistocene South African sites and among Plio-Pleistocene South African *Papio* taxa compared with a sample of extant *P. h. ursinus* specimens. Pink indicates those populations designated as *P. cf. robinsoni*, red indicates those populations designated as *P. robinsoni*, blue indicates those populations designated as *P. h. angusticeps*, purple indicates those populations designated as *Papio* sp. indet., and gold indicates extant *P. h. ursinus*. For samples by tooth position, see Tables 1 and 2 and SOM Tables S1–S4.

Member 1. Overall, the material is sufficient to suggest it is most similar to *P. robinsoni* among known papionin taxa, and we conservatively assign it to *P.* cf. *robinsoni* until more complete and definitive fossils come to light.

<u>3.1.15. Kromdraai B</u> Previous authors generally also recognized both *P. h. angusticeps* and *P. robinsoni* from Kromdraai B, at least tentatively (Delson, 1984, 1988; McKee et al., 1995; Heaton, 2006). There are a few specimens at Kromdraai B that are referable to the genus *Papio*, most consistent with *P. robinsoni*. KB 5416, a subadult female palate with partial dentition, is large and displays maxillary fossae that are shallow to moderate. KB 5230 is a large female mandible with shallow mandibular corpus fossae and a large P₄. KA 196 and KA 197, mandibular fragments described as "Parapapio" coronatus by Freedman (1957) but noted to be most similar to Papio, are considered Papio specimens here.¹ These specimens have since been renumbered KB 94 and KB 104, respectively, implying a revised association with KB deposits rather than those from KA. Without more definitive craniofacial material, it is not possible to assign these Papio specimens to any species with certainty. Overall, however, they are more consistent with *P. robinsoni* and we assign them to *P. cf. robinsoni* here. As with Kromdraai A, the Papio dental material at Kromdraai B is

¹ The holotype of *Parapapio coronatus* Broom and Robinson, 1950 - KA 195, since renumbered to KB 122 - is most likely a junior synonym of *Cercopithecoides* (Szalay and Delson, 1979; Anderson et al., 2015).

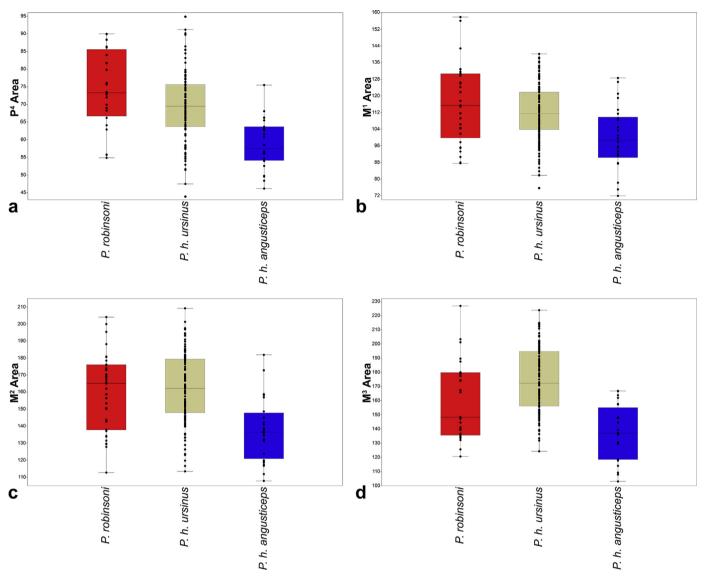


Figure 8. Boxplots illustrating dental size ranges (expressed as tooth area = maximum mesiodistal length x maximum buccolingual width) for upper P⁴s and upper molars among Plio-Pleistocene South African *Papio* taxa and a sample of extant *P. h. ursinus* specimens. Red indicates those populations designated as *P. robinsoni*, blue indicates those populations designated as *P. h. angusticeps*, and gold indicates extant *P. h. ursinus*. For samples by tooth position, see Tables 1 and 2 and SOM Tables S1–S4.

consistent with a single population (Figs. 5–9), and we therefore do not find any convincing evidence at this time suggesting both *P. robinsoni* and *P. h. angusticeps* are present at Kromdraai B.

<u>3.1.16.</u> Bolt's Farm Pit 10? An uncatalogued right maxillary fragment with M^{2-3} that may be from Pit 10 is similar in dental size to *P. robinsoni*. The zygomatic arch is present, and its anterior face is invaded by a clear but shallow fossa, although it is hard to gauge definitively because of damage and glyptol glue. This specimen is incompatible with *Parapapio* because of the presence of the fossa and its larger size. Overall morphology is most consistent with *P. robinsoni*, and we recognize it here as *P. cf. robinsoni*.

<u>3.1.17. Swartkrans II</u> Both Freedman (1957) and Delson (1984) recorded *P. robinsoni* as present at the Swartkrans II locality on the basis of the female partial cranium SK II 25. The specimen preserves much of the left side of the face and neurocranium, parts of the right side of the face, and partial dentition. While undoubtedly a member of the genus *Papio*, the female SK II 25 is somewhat smaller than most *P. robinsoni*, it possesses a narrower muzzle, and displays smaller premolars. In fact, SK II

25 is also reminiscent of the P. h. angusticeps females at Cooper's A but slightly larger in size. In addition to SK II 25, the male Papio mandible SK II 26 displays very distinct and deep corpus fossae in contrast with the all of the P. robinsoni and P. cf. robinsoni mandibular material at Swartkrans Members 1-3 and Drimolen Main Quarry. Overall, the observable craniodental features are most consistent with extant Papio and P. h. angusticeps. However, the dental size of the Swartkrans II population, particularly SK II 26, is at the high end of the range of P. h. angusticeps and most similar to P. robinsoni and extant Papio. If SK II 26 is included in the P. h. angusticeps hypodigm, it would have the largest P_4 in the sample and be larger in M_3 size than all other known males except one (CO 134B/D) (Fig. 6). However, the total P. h. angusticeps size range would still not be larger than in extant P. ursinus or even P. robinsoni (Fig. 6). Put another way, SK II 26 would be among the largest known P. angusticeps individuals, but not extend the known size range of P. h. angusticeps in an unreasonable way. In total, this combination of size (more similar to P. robinsoni) and morphology (most similar to P. h. angusticeps) makes a definitive

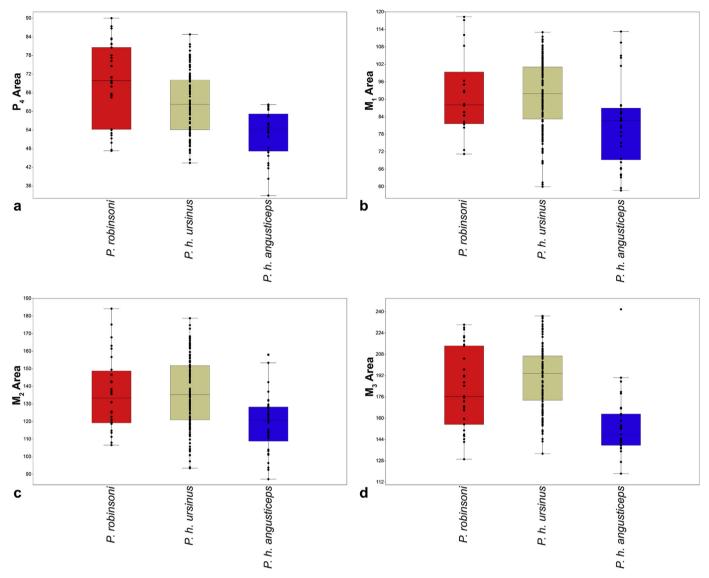


Figure 9. Boxplots illustrating dental size ranges (expressed as tooth area = maximum mesiodistal length x maximum buccolingual width) for lower P₄s and lower molars among Plio-Pleistocene South African *Papio* taxa and a sample of extant *P*. *h. ursinus* specimens. Red indicates those populations designated as *P. robinsoni*, blue indicates those populations designated as *P. h. angusticeps*, and gold indicates extant *P. h. ursinus*. For samples by tooth position, see Tables 1 and 2 and SOM Tables S1–S4.

species attribution difficult for the SK II series. We therefore assign this material to *Papio* sp. indet.

<u>3.1.18. Sterkfontein Members 5 and 6</u> As noted above, some specimens assumed to be from Member 4 may derive from Member 5 instead (see also Heaton, 2006). There is also a male partial mandible (SWP 1232) from Member 5 displaying a long P₃ honing flange and definitive mandibular corpus fossae, typical of *Papio*. From Member 6, SWP 1012 represents a female partial mandible with deep corpus fossae below the P₃-M₁, again consistent with *Papio*. While Heaton (2006) argued that the *Papio* material from Member 5 represents *P. robinsoni*, we find none of the specimens diagnostic. Additional material is necessary to confirm which species may be present at Members 5-6. We therefore recognize these specimens as *Papio* sp. indet.

<u>3.1.19. Swartkrans Member 3</u> There are a number of specimens, more fragmentary than those in Member 2, that are consistent with *P. robinsoni* in size, but potentially consistent with other *Papio* taxa or other papionin species as well. Specimens previously attributed to *P. robinsoni* from Member 3 by Delson (1988) do not

preserve enough diagnostic cranial morphology to be certain. Therefore, we consider this population best allocated to Papionini gen. et sp. indet. until more complete and diagnosable material is recovered.

<u>3.1.20. Cooper's D</u> A small collection of isolated teeth from Cooper's D were described as cf. *Papio* sp. (Folinsbee and Reisz, 2013). Because this material is fragmentary, preserving no diagnostic features of any fossil or extant *Papio* species, and is also within the size range of *P. h. angusticeps*, *P. robinsoni*, and other papionin taxa, we conservatively assign the previously recognized *Papio* material at Cooper's D to Papionini gen. et sp. indet.

3.2. Review of East African Plio-Pleistocene Papio-bearing sites

In addition to the South African sites above, there are several in eastern Africa where *Papio* has been reported. These generally fall into two categories, middle and later Pleistocene material that is modern in overall appearance and assigned either to *Papio* sp. or one of the extant varieties (e.g., *P. hamadryas* sspp., *P. h.* cf. anubis, etc.), or earlier Plio-Pleistocene forms that were initially identified as *Papio*, but subsequently allocated to other genera or are too fragmentary to identify to genus with confidence (Szalay and Delson, 1979; Jablonski, 2002; Jablonski and Frost, 2010).

3.2.1. Middle and later Pleistocene Among the more complete specimens of this group is a well-preserved partial cranium of a juvenile male (BSPG, 1931 II 26) from Bed IV or above at Olduvai that has been referred to Papio sp. and P. cf. hamadryas sspp. (Remane, 1925; Dietrich, 1942; Leakey, 1971; Szalay and Delson, 1979; Frost, 2014: Fig. 7). Even though it is a juvenile with the M² beginning to erupt, the specimen shows a clear anteorbital drop, maxillary fossae and ridges, and a flattened muzzle dorsum. A left maxilla with M^3 and left male mandibular symphysis with $C-P_4$ from Lemagrut Korongo and an M² in a small maxillary fragment from Laetoli (probably the Ngaloba Beds) that preserve clear facial fossae and other diagnostic features of *Papio* have been variously referred to P. cf. neumanni, P. hamadryas cf. anubis, and Papio sp. (Dietrich, 1942; Leakey and Delson, 1987; Harrison, 2011). A series of diagnostic facial material, including relatively complete male and female maxillae and mandibles, from Asbole in the Afar region of Ethiopia dated to 600 Ka was assigned to *P. hamadryas* ssp. indet. by Frost and Alemseged (2007). Morphologically the specimens are consistent with both P. h. anubis and P. h. hamadryas, but generally on the smaller end of the extant size range. Here we refer all this material from Olduvai, Laetoli, Lemagrut Korongo, and Asbole to P. hamadryas ssp. indet. Isolated teeth from the ca. 500 Ka Dawaitoli Formation of the Middle Awash, Ethiopia, and Members J - K of the Shungura Formation. Ethiopia have also been assigned to Papio sp. or P. hamadarys sspp., but are not diagnostic beyond being papionins similar in size to extant P. hamadryas (Eck, 1976, 1977; Kalb et al., 1982; Frost, 2001a; Frost, 2007a, 2007b). Overall, this material suggests that baboons similar to the extant varieties were widespread in eastern Africa by the Middle Pleistocene. Their relationship to Papio fossils from South Africa is unclear.

3.2.2. Earlier Plio-Pleistocene Several earlier Plio-Pleistocene specimens have been assigned to Papio, often as new species, but these have been subsequently allocated to other genera. Two partial crania and several other fragments from Olduvai Bed I originally described as cf. Papio sp. nov. by Leakey and Leakey (1976) have more recently been assigned to Gorgopithecus major (Gilbert et al., 2016a). Leakey and Leakey (1976) also tentatively allocated a maxilla and three partial mandibles from the Upper Burgi and Okote Members to cf. Papio sp. Frost (2001a) suggested this material could represent either a species of Papio or late occurring Dinopithecus quadratirostris (since transferred to Soromandrillus quadratirostris [Gilbert, 2013]), and thus left it as Papionini gen. et sp. indet. C. Jablonski et al. (2008), on the other hand, placed two of these (KNM-ER 143 and KNM-ER 144) in Parapapio sp. indet. A. and two (KNM-ER 141 and KNM-ER 142) in *Parapapio* sp. indet. B. A right dP⁴ and distal humerus from Laetoli tentatively designated cf. Papio sp. by Leakey and Delson (1987) has been reallocated, along with a more recently discovered male right upper canine, to Papionini gen. et sp. indet. by Harrison (2011).

Leakey (1969) described a partial skull from the Baringo Chemeron Formation dated to 3.2 Ma as the type specimen of a new species, *Papio baringensis*. Since then, however, a number of studies have suggested that this specimen is morphologically most similar to *Theropithecus*, and it has been tentatively reassigned there by most authors (e.g., Eck and Jablonski, 1984, 1987; Delson and Dean, 1993; Frost, 2001a; Frost and Delson, 2002; Jablonski, 2002; Jablonski and Frost, 2010; Gilbert, 2013).

Finally, sizeable numbers of Plio-Pleistocene specimens from Omo (Usno Formation and Members A through G of the Shungura Formation) were previously referred to *Papio* sp. by Eck (1976, 1977).

These largely consisted of isolated teeth, but more diagnostic material that spans Members D through lower G includes two female partial crania, a male rostrum, and several mandibular and maxillary fragments that clearly indicate a large long-faced papionin. Delson (1984), Delson and Dean (1993), and Frost (2001a) referred these more complete specimens along with the holotype cranium of *Papio quadratirostris* from the Usno Formation (Iwamoto, 1982), and a number of others from the Humpata Plateau in Angola, to *Dinopithecus*, but these authors also recognized *Dinopithecus* as a subgenus of *Papio.*² More recently, Gilbert (2013) transferred this material to a new genus, *Soromandrillus*, a decision we follow here. Other specimens from Omo Shungura not listed in the *Soromandrillus* hypodigm are best left as Papionini gen. et sp. indet. although many of the isolated teeth may well be *S. quadratirostris*.

3.3. Morphological analysis of revised hypodigms

To test the robusticity of our revised hypodigms for the named South African *Papio* taxa as well as our hypothesis that *P. robinsoni* and *P. h. angusticeps* do not co-occur at Kromdraai A or elsewhere, we conducted several descriptive statistical analyses. For comparison and in order to make the fewest possible changes relative to previous taxonomic schemes, all material previously reported as "*Papio*" for each site where *P. robinsoni*, *P. h. angusticeps*, or both had been argued to occur was lumped together into one population for analysis (see SOM Tables S1–S4 for a complete list of specimens and measurements). Our results are presented in Figures 5–9 and suggest that, with very few exceptions, the range of variation found at each site is no greater than that of extant *P. h. ursinus* collected in the past century or so. Likewise, when data across sites are combined into paleotaxa, the range of variation is again similar to that observed in *P. h. ursinus*.

In a few instances, outliers do exist. However, in these cases we note that other large papionins are present at the site in question, or surrounding sites, and the allocation of these specimens to Papio is only tentative. In other words, almost all outlier specimens are lacking in sufficiently diagnostic morphology to confirm their attribution to Papio, and therefore could represent other papionin taxa, such as Gorgopithecus or Dinopithecus. For example, SK 590 from Swartkrans Member 1 is a large maxillary fragment preserving a large M¹ and M² that metrically fall within the Gorgopithecus range as well as the P. robinsoni range. Likewise, SK 419 and SK 445 preserve large M₁-M₂ from Swartkrans Member 1 that are both within the *Gorgopithecus* range as well. KA 163 preserves a dP_4 -M₁, with the M₁ being in the *Gorgopithecus* size range. HGD 660 is identified as an isolated M₁, and its measurements are within the Gorgopithecus size range, but it is also possible this specimen is an isolated M₂ instead. Finally, while CO 134 B/D exhibits a very large M₃, the other teeth in this subadult male mandible are not outliers and the M₃ may also be slightly larger due to the fact that it is unworn. There are other large teeth included in our sample that similarly overlap the range of multiple large papionin taxa, and as noted above, it is also possible that a few isolated teeth are mis-identified to position. In an effort to be conservative, we included as many specimens previously attributed to Papio as we could for each site, but it may ultimately be more reasonable to refer to some of the outlier specimens as Papionini gen. et sp. indet., particularly since some of them overlap the size ranges of *G. major* and *Dinopithecus ingens*, which are also known to co-occur with P. robinsoni or P. h. angusticeps

² Eck and Jablonski (1984, 1987) transferred the Usno Fm. holotype to *Ther-opithecus quadratirostris* and Jablonski (1994, 2002) considered the Angolan and Shungura specimens to be *T. baringensis.* We strongly disagree, as detailed in the cited publications.

(Freedman, 1957; Szalay and Delson, 1979; Delson, 1984, 1988; Jablonski, 2002; Jablonski and Frost, 2010).

Based on the above reassessment of fossil Papio species distributions and the revised hypodigms resulting from our study of the material, we also conducted a 3D geometric morphometric analysis of all three South African fossil Papio taxa relative to other papionins, particularly extant P. hamadryas ssp (Frost et al., 2003). As with previous analyses of papionin crania, allometry was the largest factor contributing to the shape variation in our sample with PC1 explaining 61% of total variance and being highly correlated with the natural log of centroid size ($r^2 = 0.87$). The second component accounted for 9.6% of the sample variance and largely separates male Mandrillus from other papionins (Fig. 10). The third component accounted for 6.0% of variance and seriated Theropithecus, Macaca, Papio, Cercocebus and Mandrillus, and Lophocebus from most positive to most negative (Fig. 10). Most of the fossil Papio specimens fell within the 95% confidence ellipse for extant Papio, but several of the P. robinsoni specimens and one Papio sp. (TMP SK II 25) fell somewhat outside. All but the last specimen are within the convex hull for extant Papio.

The discriminant function analysis was highly accurate among the model sample, with the cross-validation correctly classifying 98.9% of specimens. All of the *P. h. angusticeps* and *P. robinsoni* specimens that could be included were classified within extant *Papio* (Table 3). Only two of the *?P. izodi* specimens, however, were classified as *Papio*, whereas two were also classified as *Soromandrillus* and one as *Procercocebus* (Table 3). Interestingly, both *Soromandrillus* and *Procercocebus* retain primitive papionin craniodental morphologies in addition to the shared derived features linking them with extant *Cercocebus* and *Mandrillus* (e.g., shallow maxillary fossae, relatively shallow to absent mandibular corpus fossae, straight nasal bones/nasal profile; see Gilbert, 2007, 2013), perhaps indicating that *?P. izodi* shares primitive papionin shape retentions with these taxa as well.

Finally, we also performed a three-part 362 character cladistic analysis on all extant and fossil African papionins at the species level, including all three South African fossil Papio taxa, and treating all extant Papio species/subspecies populations as individual OTUs. In the first analysis, we enforced no topological constraints on the ingroup (i.e., Morphology-only) and recovered six most parsimonious trees (MPTs). The majority-rule consensus suggests that P. h. angusticeps and P. robinsoni are both likely to be nested within the modern P. hamadryas radiation (Fig. 11). However, the extant P. hamadryas taxa are reconstructed as a paraphyletic group in all MPTs and P. robinsoni is outside of the crown P/L/T/R clade in two of the six MPTs. Thus, P. robinsoni is also outside of the P. hamadryas + Lophocebus + Rungwecebus group (along with P. h. papio) in the strict consensus tree (Fig. 11; see SOM for all MPTs). ?P. izodi is hypothesized as a stem P/L/T/R taxon in all MPTs, never being found within the modern P. hamadrayas radiation. P. h. angusticeps is reconstructed as the sister taxon to P. h. kindae in all MPTs, in support of Delson's (1984, 1988) observations. In the second analysis enforcing the ((C,M),(T,(P,L,R))) backbone, ?P. izodi is again outside of the crown P/L/T/R clade in all trees, but P. h. angusticeps and P. robinsoni are always within the P. hamadryas clade, with P. h. angusticeps reconstructed as the sister taxon to a P. robinsoni+P. h. hamadryas clade at the base of the P. hamadryas radiation (Fig. 12; see SOM for all 14 MPTs). In the third analysis, enforcing the ((C,M),(T,L,(P,R))) backbone, P. h. angusticeps and P. robinsoni are again always within the P. hamadryas clade, in the same relative positions as in the second analysis (Fig. 13). ?P. izodi is again never found within the crown P/L/T/R clade, instead being reconstructed as a late-occurring stem P/L/T/R taxon (Fig. 12; see SOM for all 12 MPTs). Among all analyses, bootstrap support for any clades within the larger *P/L/T/R* clade was low, except in the cases of *Theropithecus* and extant *Lophocebus* (Figs. 11–13).

In total, our results suggest that, of the four widely recognized fossil *Papio* taxa in the southern African Plio-Pleistocene fossil record, *?P. izodi* appears the most morphologically primitive and distinct from the extant population, *P. robinsoni* specimens are most often within the modern range of morphological variation, but also sometimes on the periphery or just outside the range of variation seen in extant populations, and *P. h. angusticeps* as well as *P. h. botswanae* are always included within the extant range of morphological variation. Based on these results, we offer revised diagnoses of these fossil species as well as the genus *Papio* below.

4. Systematic paleontology

Class MAMMALIA Linnaeus, 1758. Order PRIMATES Linnaeus, 1758. Infraorder CATARRHINI E. Geoffroy, 1812 Superfamily CERCOPITHECOIDEA Gray, 1821 Family CERCOPITHECIDAE Gray, 1821 Subfamily CERCOPITHECINAE Gray, 1821 Tribe PAPIONINI Burnett, 1828

Papio Erxleben, 1777

(= or including: *Cercopithecus* Erxleben, 1777, in part. *Cynocephalus* Geoffroy and Cuvier, 1795 (not of Boddaert, 1768). *Simia* (*Chaeropithecus*) Gervais, 1839; Senechal, 1839. *Choeropithecus* Blainville, 1839. *Hamadryas* Lesson, 1840 (not of Hübner, 1806). *Mandrillus* Milne-Edwards, 1841 (not of Ritgen, 1824). *Cercopithecus* Linnaeus, 1758: Peters, 1853, in part. *Choiropithecus* Reichenbach, 1862. *Theropithecus* I. Geoffroy, 1843: Reichenbach, 1863, in part (fide Napier, 1981, p. 80). *Comopithecus* Allen, 1925. *Parapapio* Jones, 1937: Broom, 1940, in part; Freedman, 1957, in part. *Papio* (*Chaeropithecus*) Gervais, 1839: Ellerman, Morrison–Scott and Heyman, 1953; Szalay and Delson, 1979, in part.)

Type species: Papio hamadryas (Linnaeus, 1758).³

Other included species: ?P. izodi Gear, 1926, P. robinsoni Freedman, 1957.

Generic diagnosis: This diagnosis is modified from those of Freedman (1957), Szalay and Delson (1979) and Frost (2007a) on the basis of our above results. *Papio* is a genus of medium to large sized papionin monkeys that is most obviously

³ The authorship and type species of *Papio* are part of a complicated history, most of which was laid out by Delson and Napier (1976, 1977). In brief, Linnaeus (1758) named *Simia hamadryas* with reference to "Alp. aegypt. 248", and *Simia sphinx* citing a number of images of a short-tailed robust monkey (either a mandrill or a drill). In 1766 (12th edition), Linnaeus named *Simia cynocephalus* with reference to Brisson and to Jonstonus (1650, Fig. 59). Buffon (1766) discussed several "baboons", with illustrations of the "grand papion", interpreted as the Guinea baboon of reddish-brown color; and the greenish-yellow "petit papion", probably an olive baboon. He also discussed the mandrill, with illustrations of both male and female individuals.

Müller (1773) used the name *Papio* at the generic level for *S. sphinx* Linnaeus, now defined as the mandrill (lectotype designated by Delson and Napier, 1976). Then Erxleben (1777) used *Papio* for several Linnaean species including *Simia sphinx*, for which he mentioned both mandrills and savannah baboons (such as the two papions of Buffon); his description was of a Guinea baboon with dark-reddish fur. Desmarest (1820) named *Cynocephalus papio* for the Guinea baboon (*Cynocephalus* Geoffroy and Cuvier, 1795, for baboons is preoccupied by *Cynocephalus* Boddaert, 1768 for colugos). Delson and Napier (1976) petitioned the International Commission for Zoological Nomenclature to decide whether to use *Papio* for mandrills or baboons, and the ICZN decided for the latter alternative, upholding common usage over absolute priority (International Commission for Zoological Nomenclature, 1982).

Therefore, the type species of *Papio* Erxleben, 1777 is *P*. (ex-*C*.) *papio* (Desmarest, 1820), if all baboon varieties are considered as full species. If they are considered subspecies, as we do here, the senior synonym and nomen becomes *P*. (ex-*S*.) *hamadryas* (Linnaeus, 1758), as is usually indicated.

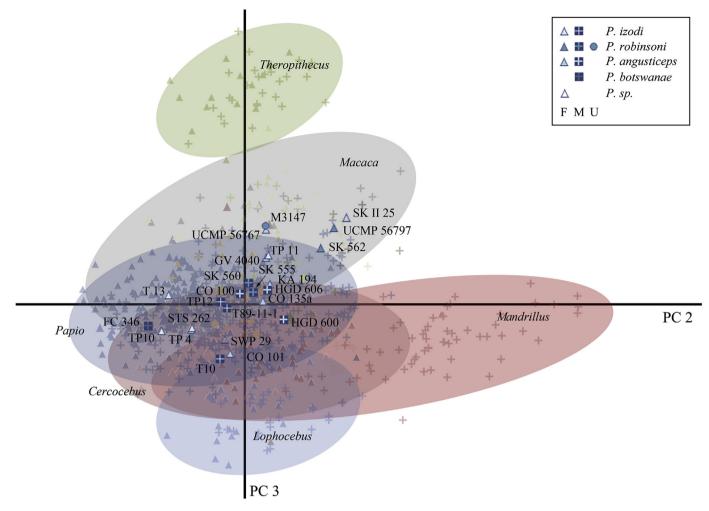


Figure 10. PCA of 3D GM analysis. Plot of PC 2 vs. PC 3 with *Papio* fossils shown and labeled (PC 1 excluded do to its strong correlation with centroid size). 95% confidence ellipses are shown for extant genera; from top to bottom: green = *Theropithecus*, gray = *Macaca*, blue = *Papio*, pink = *Cercocebus*, red = *Mandrillus*, light blue = *Lophocebus*. Note that several of the *P. robinsoni* specimens and one *Papio* sp. (TMP SK II 25) fall somewhat outside of the 95% confidence ellipse for extant *Papio*, but only SK II 25 is completely outside the convex hull for extant *Papio*. See also results of Discriminant Functions Analysis (DFA) in Table 3.

distinguished from other large papionins such as Mandrillus, Soromandrillus, Theropithecus, Dinopithecus, Gorgopithecus, and *Paradolichopithecus* by the combination of definitive maxillary fossae, mandibular corpus fossae, a flattened to slightly "tented" muzzle dorsum with relatively vertical sides in both males and females, and definitive maxillary ridges, especially in males. It possesses a marked anteorbital drop, which is distinct from Parapapio, Lophocebus, Rungwecebus, Cercocebus, Procercocebus, and most Macaca. Glabella and the supraorbital region is more prominent than in most papionins, but less so than in Theropithecus. The "squared" or slightly "tented" muzzle in crosssection is different from T. (Theropithecus), Parapapio, Pliopapio, Gorgopithecus, Paradolichopithecus, Macaca (other than the Sulawesi species), Lophocebus, Procercocebus, and Cercocebus, but shared with T. (Omopithecus), Dinopithecus, Mandrillus, and Soromandrillus. The cranial vault commonly lacks a sagittal crest, or if one is present, it is found only in the occipital region near inion. The temporal lines are typically "pinched" (converging medially along the margin of the supraorbital torus and then taking a sharp turn posteriorly; see Gilbert, 2007), particularly in males, and inion is usually inferiorly directed, unlike in Mandrillus, Soromandrillus, Cercocebus, and Procercocebus. The molars are more straight sided and the crowns less flaring than those of Mandrillus, Lophocebus, and Cercocebus, but more sloping than those of *Theropithecus*. The P4 is not typically enlarged relative to the M1 as it is in *Mandrillus*, some *Soromandrillus*, and *Cercocebus*. The incisors are relatively large compared to the molars (which is typical of many papionins), but larger than those of *Parapapio*, *Paradolichopithecus*, and *Theropithecus* and smaller than those of *Lophocebus* and *Cercocebus*. The postcranium is known only for the extant species and is more terrestrially adapted than other cercopithecids besides *Theropithecus*. Unlike *Theropithecus*, the hand of *Papio* does not display noticeable lengthening of manual digit ray I or shortening of manual digital ray II.

Papio hamadryas (Linnaeus, 1758).

(=: Simia hamadryas Linnaeus, 1758; Simia cynamolgus Linnaeus, 1758; Cynocephalus wagleri Agassiz, 1828; Cynoscephalus hamadrias Rüppell, 1835 (lapsus); Hamadryas chaeropithecus Lesson, 1840; Theropithecus nedjo Reichenbach, 1863 (fide Napier, 1981, p. 80); Hamadryas aegyptiaca Gray, 1870; Papio arabicus Thomas, 1899; Papio brockmanni Elliot, 1909; Papio hamadryas abissinicus Roth, 1965; Papio hamadryas sudanensis Roth, 1965. Including [synonyms of other subspecies, indicated as a, c, p, k, u]: Simia cynocephalus Linnaeus, 1766 (Papio cyanocephalus: Smithers, 1966; lapsus); Simia sphinx Erxleben, 1777 (not of Linnaeus, 1758) [p]; Simia porcaria Boddaert, 1787 (not of Brünnich,

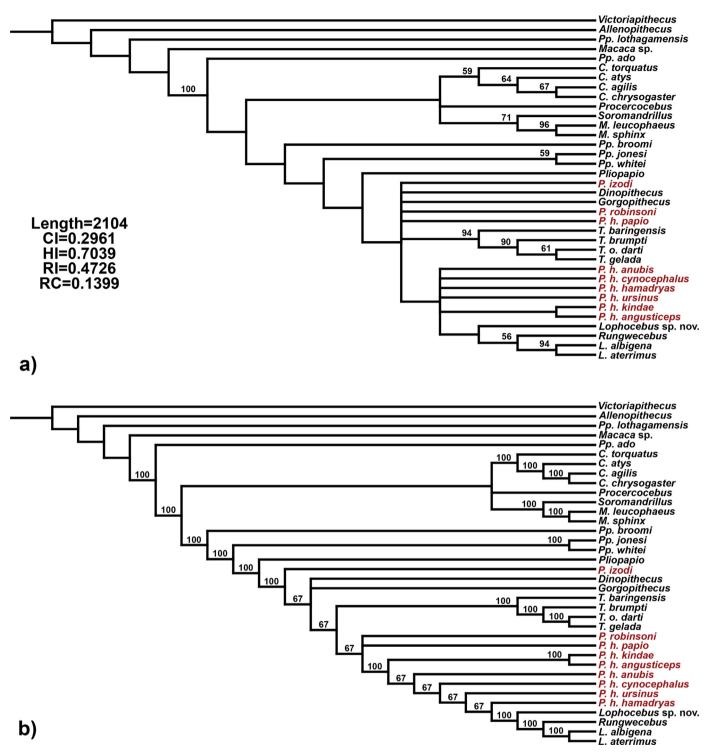


Figure 11. a) Strict consensus tree from the morphology-only analysis. Numbers above branches indicate bootstrap support over 50% for any clades. b) Majority-rule consensus tree from this analysis. Numbers above branches represent the percentage of MPTs in which a given clade is found. For matrix and individual MPTs, see SOM.

1782) (Papio porcarius E. Geoffroy, 1812; emendation) [u]; Simia (Cercopithecus) hamadryas ursinus Kerr, 1792; Simia (Papio) variegata Kerr, 1792 [suppressed by ICZN 1970, may be sphinx Linnaeus, 1758 or cynocephalus Linnaeus, 1766]; Simia basiliscus Schreber, 1800 [c]; Simia sublutea Shaw, 1800 [c]; Simia sphingiola Hermann, 1804 [u]; Papio comatus E. Geoffroy, 1812 [u]; Cynocephalus papio Desmarest, 1820; Cynocephalus babouin Desmarest, 1820 [c] (Cynocephalus babuin: Jardine, 1833; Papio babuini: Roth, 1965; lapsi); Cynocephalus antiquorum Schinz, 1821 [c]; Cynocephalus capensis Smith, 1826 [u]; Simia anubis Lesson, 1827; Cynocephalus thoth Ogilby, 1838 [c] (Papio toth: Matschie, 1898; lapsus); Cynocephalus olivaceus I. Geoffroy, 1851 [p]; Papio rubescens Temminck, 1853 [p]; Cercopithecus ochraceus Peters, 1853 [c] (P. cynocephalus ochraeus: Roth, 1965; lapsus?); Cercopithecus flavidus Peters, 1853 [c]; Cynocephalus choras Ogilby, 1838 [a]; Cynocephalus doguera Pucheran and Schimper, 1856 [a]; Cynocephalus langheldi Matschie, 1892 [c]; Papio thoth ibeanus Thomas, 1893 [c]; Papio pruinosos Thomas,

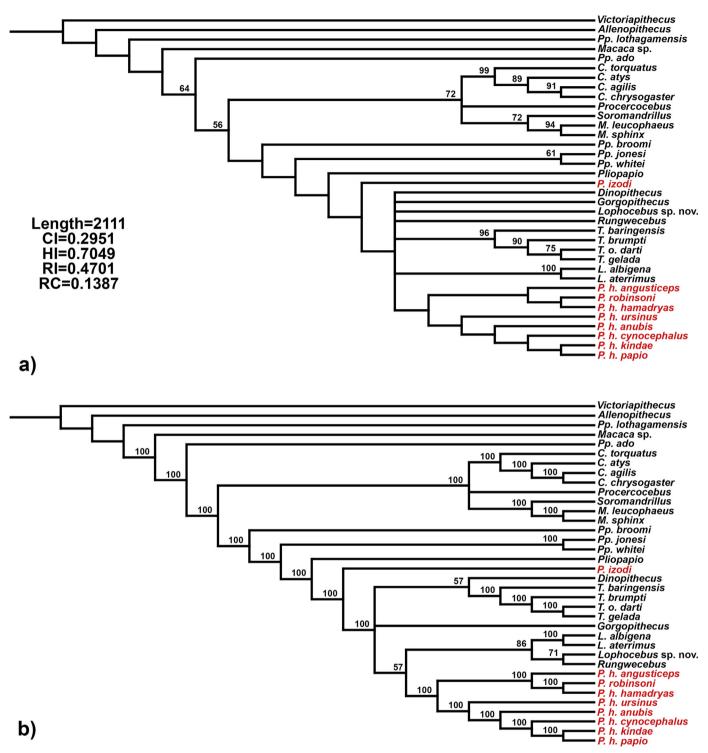


Figure 12. a) Strict consensus tree from the (*Papio, Lophocebus, Rungwecebus*) [((*C*,*M*),(*T*,(*P*,*L*,*R*)))] molecular backbone analysis. Numbers above branches indicate bootstrap support over 50% for any clades. b) Majority-rule consensus tree from this analysis. Numbers above branches represent the percentage of MPTs in which a given clade is found. For matrix and individual MPTs, see SOM.

1897 [a]; Papio neumanni Matschie, 1897 [a]; Papio heuglini Matschie, 1898 [a]; Papio yokoensis Matschie, 1900 [a]; Papio lydekkeri Rothschild, 1902 [a]; P. anubis olivaceus de Winton, 1902 [a] (not of I. Geoffroy, 1851); Papio furax Elliot, 1907 [a]; Papio strepitus Elliot, 1907 [c]; Papio nigeriae Elliot, 1908 [a]; Papio tesselatum Elliot, 1909 [a]; Papio porcarius griseipes Pocock, 1911 [u]; Papio lestes Heller, 1913 [a]; Papio vigilis Heller, 1913 [a]; Papio sylvestris Lorenz, 1915 [a]; Papio werneri Wettstein, 1916 [a]; Papio graueri Lorenz, 1917 [a]; Choiropithecus rhodesiae
Haagner, 1918 [c]; P. kindae Lönnberg, 1919; Papio porcarius orientalis Goldblatt, 1926 [u]; Papio porcarius occidentalis Goldblatt, 1926 [u]; Cynocephalus transvaalensis Zukowsky, 1927 [u]; Papio cynocephalus jubilaeus Schwarz, 1928 [c]; Papio porcarius nigripes Roberts, 1932 [u]; Papio porcarius ngamiensis Roberts, 1932 [u]; Papio porcarius chobiensis Roberts, 1932 [u]; Parapapio angusticeps Broom, 1940; Papio comatus ruacana Shortridge,

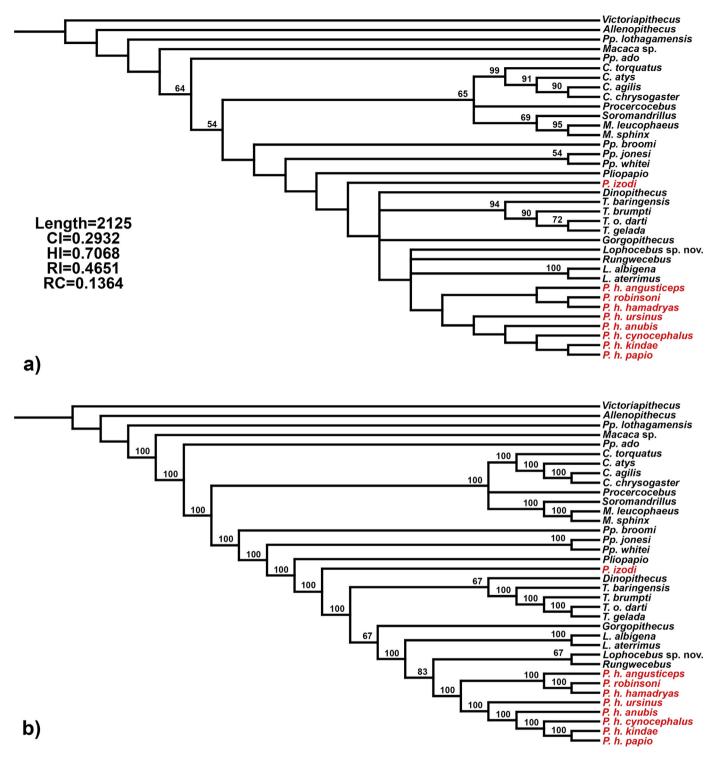


Figure 13. a) Strict consensus tree from the (*Papio, Rungweeebus*) [((*C*,*M*),(*T*,*L*,(*P*,*R*)))] molecular backbone analysis. Numbers above branches indicate bootstrap support over 50% for any clades. b) Majority-rule consensus tree from this analysis. Numbers above branches represent the percentage of MPTs in which a given clade is found. For matrix and individual MPTs, see SOM.

1942 [u]; Papio ruhei Zukowsky, 1942 [c]; Papio angusticeps (Broom, 1940): Freedman, 1957; Papio doguera tibestianus Dekeyser and Derivot, 1960 [a]; P. ursinus chacamensis Roth, 1965 [u] (nomen nudum); Papio anubis niloticus Roth, 1965 [a] (nomen nudum); Papio izodi Gear, 1926: Szalay and Delson, 1979, in part; Papio hamadryas angusticeps (Broom, 1940): Williams et al., 2012; Papio hamadryas botswanae Williams et al., 2012.) Type specimen: illustration cited by Linnaeus from "Alp. aegypt. 248"

Included subspecies: *P. h. hamadryas* (Linnaeus, 1758), *P. h. cynocephalus* (Linnaeus, 1766), *P. h. ursinus* (Kerr, 1792), *P. h. papio* (Desmarest, 1820), *P. h. anubis* (Lesson, 1827), *P. h. kindae* Lönnberg, 1919; *P. h. angusticeps* (Broom, 1940), *P. h. botswanae* Williams et al., 2012.

Specific diagnosis: Differs from *?P. izodi* in the presence of deeper maxillary fossae and mandibular corpus fossae, more pronounced maxillary ridges, a definitive anteorbital drop, a relatively longer, narrower and more flattened muzzle dorsum, a relatively taller malar region, orbits and molars that are smaller relative to overall cranial size, and overall larger size (other than for *P. h. kindae*). Differs from *P. robinsoni* in that the nasals are prominent above the maxillary ridges in lateral view, the maxillary and mandibular corpus fossae are often deeper and more clearly defined, and the P⁴ is smaller.

Papio robinsoni Freedman, 1957.

(= or including: *Parapapio* sp. Robinson, 1952; *Parapapio whitei* Broom, 1940: Freedman, 1957, 1976, in part; *Papio hamadryas robinsoni*: Szalay and Delson, 1979). Type Specimen: TMP SK 555.

Specific Diagnosis:

Differs from *P. hamadryas* and *?P. izodi* in the possession of larger upper and lower P4s (on average), a flatter muzzle with the nasals positioned below the maxillary ridge in lateral view (particularly in males), shallower maxillary and mandibular corpus fossae (on average), and a high incidence of the maxillae meeting in the midline of the face before nasion. Further differs from *?P. izodi* in overall larger size, relatively smaller orbits and molars, a definitive anteorbital drop, better developed facial fossae, better developed maxillary ridges, and a relatively taller malar region.

?Papio izodi Gear, 1926 (new combination).

(= or including: *Papio antiquus* Haughton, 1925, in part (*Papio africanus*: Gear, 1926, lapsus: Broom, 1934; in part); *Papio izodi* Gear, 1926; *Parapapio izodi* (Gear, 1926): Broom, 1940, in part; *Parapapio antiquus* (Haughton, 1925): Broom, 1948, in part, Freedman, 1957, in part; *Papio antiquus* Haughton, 1925: Gear, 1958, in part; *Papio wellsi* Freedman, 1961; *P. whitei* Broom, 1940: Freedman, 1965, in part.

Type Specimen: UW-AD 992 lectotype (Jones, 1937).

Specific Diagnosis: Differs from *P. hamadryas* and *P. robinsoni* in possessing variably developed maxillary ridges (i.e., absent to moderately developed in males and females), variably developed maxillary fossae (absent to moderately developed in males and females), weak to absent mandibular corpus fossae (in both sexes), a relatively shorter and broader rostrum, a more weakly developed to absent anteorbial drop, a relatively short malar region, and relatively large orbits and molars. Differs further from *P. robinsoni* in smaller cranial size. Differs further from *P. hamadryas* by displaying a relatively large dentition.

5. Discussion

After consideration of the relevant morphology, we conclude that *?P. izodi* is not definitively a member of the genus *Papio* (possibly necessitating a new genus), and that *P. robinsoni* and *P. h. angusticeps* are successively derived towards extant *Papio* populations, with *P. h. angusticeps* almost certainly a member of the modern species and possibly *P. robinsoni* as well. The inclusion of *?P. izodi* within the genus *Papio* would require a morphological expansion of the genus to include features such as relatively large orbits (and molars), variable facial fossae, the expression of weak to no maxillary ridges in males and females, a short malar region, and the variable expression of an anteorbital drop (see also Gilbert, 2013; Gilbert et al., 2015). We interpret many of these features as more "primitive" and distinct compared to those expressed by modern *Papio*, and our comprehensive morphometric and cladistic analyses also support this interpretation. At the same time, *?P. izodi* is generalized enough to

have potentially given rise to *Papio* sensu stricto, and it does variably express a number of cranial features consistent with the extant genus relative to the more primitive *Parapapio* species. In addition, the only known postcranial specimen attributable to *?P. izodi* from Taung (a partial humerus) also displays features consistent with extant *Papio* (Gilbert et al., 2016b). Thus, while it seems clear that *?P. izodi* is derived relative to other fossil African papionin taxa (such as *Parapapio*), it is not clear whether it is a member of the genus *Papio*, a generalized member of the crown *P/L/R/T* clade, or simply a late occurring stem *P/L/R/T* taxon.

Another taxon that has previously been included within the genus *Papio*, most often as a subgenus, is *Dinopithecus*. Our cladistic analysis, as well as those by Gilbert (2008, 2013) and Gilbert et al. (2016a), strongly suggests that *Dinopithecus* lies outside of the genus *Papio* and finds the evidence for *Dinopithecus* as a crown vs. stem member of the *P/L/R/T* clade equivocal. Similar to *?P. izodi, Dinopithecus* displays weak facial fossae, which we interpret as a conservative retention. It is true that what is preserved of the male cranium SK 599 looks very similar to those regions of a modern *P. hamadryas* specimen, particularly in dorsal view, but differences exist in the basicranium (see Gilbert, 2013), and there is too little morphology present to know just how similar or different the entire skull would have been. Thus, we recognize *Dinopithecus* as a separate genus from *Papio*, pending additional fossil evidence.

The other taxa discussed here are undoubtedly closely related to the modern baboon P. hamadryas, and therefore members of the genus Papio, but seemingly differ in degree in terms of their morphological similarity to the extant populations. On the basis of our analyses, we formally recognize *P. h. angusticeps* as a clear member of the modern radiation as initially suggested by Delson (1984, 1988) and supported by Frost (2007a, 2007b) and Gilbert et al. (2015) (see diagnosis above). All morphometric and cladistic analyses place the various P. h. angusticeps specimens within the modern P. hamadryas radiation, and there is little justification for its continued separation at the specific level if the BSC is being used. *P. robinsoni*, on the other hand, is slightly different from the modern taxa in terms of shape, and in the Morphology-only trees it falls outside of the rest of the modern Papio taxa in one-third of the MPTs (2 out of 6 trees; see SOM). However, it is classified as a member of extant Papio in our morphometric DFA and placed within the monophyletic Papio clade in all MPTs when a molecular backbone is enforced, illustrating its overall similarity to the modern taxa. Again, if a biological species concept is being used, one could probably make a good argument for placing *P. robinsoni* within the modern species (i.e., P. h. robinsoni), but it could be equally argued that specific rank should be maintained (P. robinsoni). Here we favor the latter taxonomic arrangement to reflect its possibly more primitive status compared to P. h. angusticeps and its consistent expression of morphological features found at much lower frequencies in modern P. hamadryas. However, we also admit that a taxonomic scheme recognizing P. h. robinsoni is almost equally compelling given the results of our analyses, and it could be argued that P. h. ursinus is, in some ways, just as morphologically distinct from the other P. hamadryas subspecies as is P. robinsoni. Thus, we consider both P. robinsoni and P. h. robinsoni to be acceptable and, essentially, equivalent alternatives, with one of us (ED) preferring P. h. robinsoni.

Our revised classification has some repercussions for our understanding of the evolution of the genus *Papio*. We find no compelling craniodental evidence to support more than one definitive fossil *Papio* taxon at any site during the Pliocene or Early Pleistocene. *P. robinsoni* and *P. h. angusticeps* do not appear to cooccur at any South African Plio-Pleistocene site, and previous recognition of both taxa at Cooper's A and Kromdraai A and B was mainly based on large or small fragmentary specimens that do not expand the range of dental size variation beyond that expected for a single population of modern *P. h. ursinus* (Figs. 5–9), particularly when the effects of time-averaging are considered. Perhaps most convincingly, the size ranges of upper and lower P4s at each site are well within what one would expect in a modern *P. h. ursinus* population; because *P. robinsoni* has been specifically noted to possess larger premolars than other fossil *Parapapio* and *Papio* taxa (see above), one might expect to see an obvious bimodal distribution or abnormally large range if two taxa were present (Figs. 5–9).

In a few of our metric dental comparisons, there are one or two specimens that extend the range of the fossil populations significantly, usually in the larger size direction (e.g., Swartkrans M¹, Swartkrans M₁, M₂, Kromdraai A M₁, Haasgat M₁). In these cases, we again note that in an effort to be conservative, we followed previously published taxonomic IDs for Papio at all of the sites as much as possible (e.g., Freedman, 1957; Delson, 1984). However, particularly in the case of Swartkrans and Kromdraai A, there are other large papionins present that could account for some of the large isolated teeth and/or maxillary/mandibular fragments previously included in Papio, namely G. major and D. ingens (see also above). In fact, out of all specimens examined with known provenience, we excluded only two fossils from our analyses, both P₄'s previously referred to Papio sp. (SK 434 and SKX 35315/16). We excluded them because they were larger than the known P₄ range for G. major, making an attribution to any specific large papionin tenuous at best. It is quite possible that other fragmentary specimens may belong to other large papionin taxa at these sites as well (some perhaps previously unrecognized, e.g., at Haasgat), so the resulting lack of significant range extension beyond that seen in extant P. h. ursinus even with possible taxonomic mixing and time-averaging is particularly compelling.

The lack of co-occurrence between *P. h. angusticeps* and *P. robinsoni* is a notable conclusion of this study given that these taxa have been argued to co-occur for 60 years since the landmark study by Freedman (1957) (see Table 9 for revised distribution of fossil *Papio* taxa). However, multiple hypotheses can be envisioned for why these two taxa do not co-occur. For example, the lack of co-occurrence could be due to fluctuations in climate and environment resulting in a range shift among these fossil *Papio* populations. A modern analog might be considered between adjacent

P. hamadryas populations: if climate change resulted in the breakdown of a river or forest barrier, for instance, it is guite possible that P. h. ursinus and adjacent P. h. cynocephalus ranges might also change slightly to the north or south, respectively. Over thousands or hundreds of thousands of years and many climatic cycles, such events could happen multiple times. If some individuals were preserved as fossils during these events, in an intermediate zone of range overlap the result would be the alternation of *P. h. ursinus* and *P. h. cvn*ocephalus at different localities. Note that if these climatic fluctuations occurred on the level of thousands of years, this would appear as a geological instant in the fossil record, and such populations or sites would likely be indistinguishable from each other, thus accounting for differing taxa at sites of approximately the same age. Yet another hypothesis might involve range extensions with interbreeding and asymmetrical gene flow similar to what is thought to be occurring between P. h. anubis and P. h. cynocephalus populations in southern Kenya today, where P. h. anubis may be expanding its range into that of P. h. cynocephalus (Charpentier et al., 2012). Other baboon hybrid zones appear to be quite dynamic as well (Bergman et al., 2008; Jolly et al., 2011). If these populations were to be preserved as fossils, they would also preserve a chaotic pattern of occurrence. Regardless of the cause, if P. robinsoni and P. h. angusticeps really were allotaxa, it might argue for subspecific status for the former taxon as it would suggest they were ecologically exclusionary, whereas more distinct taxa might be better able to overlap.

Another interesting observation from the above morphological analyses is that, if truly reflective of the primitive condition for the genus *Papio*, the shared morphology of *P. robinsoni* and *P. h. angusticeps* suggests that *P. h. ursinus* is derived in its cranial morphology despite being consistently reconstructed as the first *P. hamadryas* subspecies to diverge from others in mitochondrialbased molecular phylogenetic analyses (Newman et al., 2004; Wildman et al., 2004; Zinner et al., 2009, 2013). *P. robinsoni* and *P. h. angusticeps* look morphometrically and qualitatively most similar to *P. h. anubis/P. h. hamadryas* and *P. h. cynocephalus/ P. h. kindae*, respectively, in features such as the less klynorhynch facial orientation (Frost et al., 2003; Leigh, 2006), overall broader cranium, and less peaked maxillary ridges. In fact, the distinctive *P. h. ursinus* morphology does not appear in the fossil record until

Table 9

Distribution of fossil Papio taxa across Plio-Pleistocene African sites. "T" = type locality of taxon.

	Papio robinsoni	Papio hamadryas angusticeps	Papio hamadryas botswanae	Papio hamadryas sspp.	?Papio izodi	Papio sp. indet.	Papionini gen. et sp. indet.	Total# of Papio species present
!Ncumtsa			Т					1
Asbole				Х				1
Bolt's Farm Pit 10	cf.							1
Bolt's Farm Pit 23	Х							1
Bolt's Farm Pit 6		Х						1
Cooper's "A"		Х						1
Cooper's D							Х	1
Drimolen	Х							1
Gladysvale		Х			cf.			2
Haasgat		Х						1
Kromdraai A		Т						1
Kromdraai B	cf.							1
Lemagrut Korongo				Х				1
Malapa		Х						1
Middle Awash, Dawaitoli Fm.				cf.				1
Olduvai Bed IV or above				Х				1
Skurweburg	Х							1
Sterkfontein Mb. 5						х		1
Sterkfontein Mb. 2					Х			1
Sterkfontein Mb. 4					Х	Х		2
Swartkrans II						Х		1
Swartkrans Mb. 1	Т							1
Swartkrans Mb. 2	cf.							1
Swartkrans Mb. 3							Х	1
Taung					Т			1

the late Pleistocene or Holocene (Freedman, 1965), Cladistically, P. robinsoni most often falls closest to P. h. papio or P. h. hamadryas, while P. h. angusticeps is sister to P. h. kindae or P. h. *hamadryas* + *P. robinsoni*, although the morphological characters uniting these taxa are unclear and P. h. papio and P. h. hamadryas represent the two extant taxa with the lowest sample sizes. Thus, some of the features uniting these taxa cladistically may be an artifact of coding issues resulting from low sample sizes and missing data as generally discussed in Gilbert (2013). Overall, the morphology of P. robinsoni and P. h. angusticeps suggest that, although P. h. ursinus appears to have been the first of the modern populations to diverge genetically, it does not retain the ancestral cranial morphology for extant baboons and, instead, is likely to be more derived in its cranial morphology compared to other extant Papio species/subspecies such as P. h. anubis, P. h. cynocephalus, and P. h. kindae. Based on their morphological similarity to P. robinsoni and P. h. angusticeps, we consider P. h. anubis and P. h. cynocephalus, in particular, as the extant taxa retaining the highest number of ancestral craniodental morphologies.

While we find no evidence that *P. robinsoni* and *P. h. angusticens* co-occur at any specific site, it is possible that ?P. izodi and an indeterminate Papio species both appear at Sterkfontein Member 4 (Table 9, Fig. 14). It is also possible that P. h. angusticeps and ?P. izodi co-occur at Gladysvale (Table 9, Fig. 14), although the provenance of the two specimens relative to each other is unclear. Given the presence of very large and derived Theropithecus dentition at Gladysvale, this implies at least part of this assemblage is younger than 1.5 Ma, and possibly even Middle Pleistocene (see also Berger and Tobias, 1994; Lacruz et al., 2002; Pickering et al., 2007), whereas the last appearance for *?P. izodi* would otherwise be Sterkfontein Mb. 4 or Taung. Therefore, in the absence of more definitive evidence regarding the exact provenance of the primate specimens and the span of time represented by the different deposits at Gladysvale, it seems probable that ?P. izodi and P. h. angusticeps do not co-occur.

Based on the distribution of *P. robinsoni* and *P. h. angusticeps* across South African sites (Table 9; Fig. 14), it appears that the first appearance datum (FAD) of modern baboons (*P. hamadryas* ssp.) is close to ~2 Ma and perhaps slightly before (~2.4–2.0), as evidenced

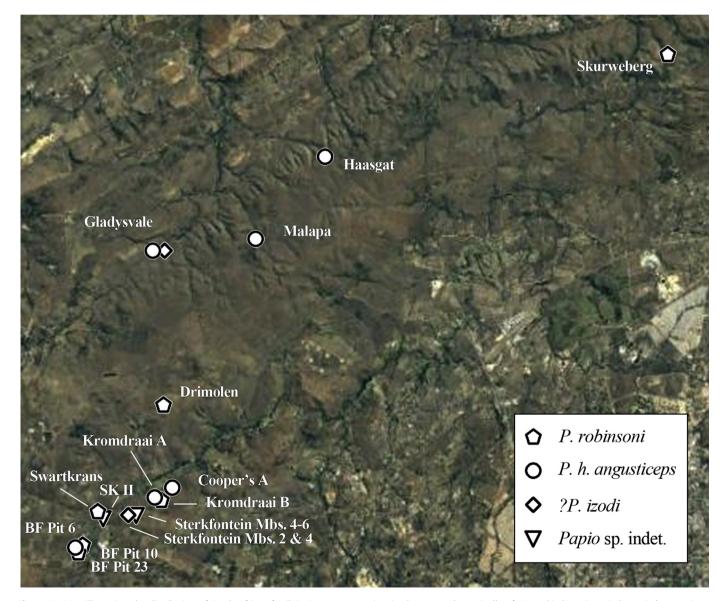


Figure 14. Map illustrating the distribution of South African fossil Papio taxa across sites in the Krugersdorp "Cradle of Humankind" region: circles = P. h. angusticeps, pentagons = P. robinsoni, diamonds = ?P. izodi, and triangles = Papio sp. indet.

by the oldest populations of *P. h. angusticeps* at Malapa and Haasgat (Adams et al., 2013; Herries et al., 2014; Gilbert et al., 2015). A broader definition of P. hamadryas including P. robinsoni might place the FAD slightly earlier still, ~2.5-2.0 Ma if P. robinsoni is confirmed as the derived Papio species at Sterkfontein Member 4 and depending on revision to the geochronology and biochronology at other sites where *P. robinsoni* is found, such as Drimolen Main Ouarry (e.g., see Adams et al., 2016) and Skurweberg. Interestingly, the FAD of both taxa overlaps the range of recent molecular clock estimates for the evolution of the extant P. hamadryas radiation, ~2.2–1.8 Ma (Newman et al., 2004; Wildman et al., 2004; Zinner et al., 2009, 2013), providing additional evidence that they are probably close to the origin of the modern species/subspecies (Gilbert et al., 2015). This FAD for modern Papio also parallels the FAD (approximately 2.0 Ma) of its close relative Lophocebus (at least that based on material with solid provenance) (Jablonski et al., 2008). Currently, no definitive fossil Papio specimens are known from eastern Africa until the Middle Pleistocene, which seems to suggest that the genus arose in southern Africa and subsequently migrated northwards.

If molecular divergence date estimates as well as the fossil record are accurate, there is good evidence to suggest that ~2.4–2.0 Ma is a reasonable FAD for the genus *Papio*, and this information may be useful in future biochronological analyses. In total, the current evidence suggests that unquestioned members of the genus *Papio* are not found in deposits much older than 2.0 million years, perhaps providing a maximum age-limit to new deposits where fossil *Papio* specimens are identified along with other fauna, such as hominins.

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