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THE NATURAL HISTORY OF MODEL ORGANISMS

Insights into the evolution of social systems and species from baboon studies

Abstract Baboons, members of the genus *Papio*, comprise six closely related species distributed throughout sub-Saharan Africa and southwest Arabia. The species exhibit more ecological flexibility and a wider range of social systems than many other primates. This article summarizes our current knowledge of the natural history of baboons and highlights directions for future research. We suggest that baboons can serve as a valuable model for complex evolutionary processes, such as speciation and hybridization. The evolution of baboons has been heavily shaped by climatic changes and population expansion and fragmentation in the African savanna environment, similar to the processes that acted during human evolution. With accumulating long-term data, and new data from previously understudied species, baboons are ideally suited for investigating the links between sociality, health, longevity and reproductive success. To achieve these aims, we propose a closer integration of studies at the proximate level, including functional genomics, with behavioral and ecological studies.

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Introduction Humans have been captivated by baboons for thousands of years: from ancient Egypt, where the god of wisdom, Thoth, was depicted with a baboon head, to the mid-19th century when Charles Darwin remarked, "He who understands baboon would do more towards metaphysics than Locke" *(Darwin, 1838).* At the beginning of the 20th century, the South African naturalist Eugene Marais provided one of the first detailed accounts of free-ranging baboons (*Marais, 1939*), and by the 1950s, baboons had become the subject of more systematic scientific enquiry, both in the field and in captivity. This was the decade that the American physical anthropologist Sherwood Washburn and his student Irven DeVore set out to investigate baboons in Kenya (*Vore and Washburn, 1961*). Washburn reasoned that these ground-living primates were a good model for early human adaptations because they evolved in African savannas alongside ancestral hominins. Meanwhile, increasing interest in the use of non-human primates as biomedical models for humans led to the funding in 1958 of a three-year proposal titled "Initiation and Support of Colony of Baboons" by the US National Institutes of Health, with the first group of baboons shipped to the United States from Kenya in 1960 (*VandeBerg, 2009*). Since then, research in captivity on
baboons as a biomedical model has been complemented by extensive fieldwork on baboon populations across Africa. Knowledge of the links between health and fitness in baboons under natural circumstances, including natural levels of genotypic and phenotypic variation, appears critical to put results from captive studies into context. An understanding of the evolution and life history of these animals in the wild also allows the scientific community to assess the validity of results derived from captive populations.

While the earlier field studies set out to uncover a baboon archetype, subsequent research has revealed that there is no such thing as “the baboon”. Indeed, many would argue that the value of this genus lies precisely in the substantial variation in the social systems, life histories and ecologies within and between the baboon species (see Box 1 for a glossary of specialist terms used in this article). Collectively, these characteristics make baboons an excellent model organism for investigating a range of fundamental biological processes, such as physiological and behavioral adaptation, hybridization and speciation with gene flow (Alfred and Baldwin, 2015). In this way, the baboon model constitutes an important complement to other mammalian model organisms, such as wild house mice (Mus musculus; Phifer-Rixey and Nachman, 2015) and deer mice (genus Peromyscus; Bedford and Hoekstra, 2015).

**Systematic classification and distribution**

Within the primate order, all extant baboons belong to the genus *Papio*. The genus is part of the tribe Papionini, within the family

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**Figure 1. Distribution of the six *Papio* species.** Species distributions are modified from Zinner et al. (2013). Male baboon drawings by Stephen Nash. Reprinted with permission from Fischer et al. (2017).

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Cercopithecidae. The fossil record and phylogeographic analyses indicate that baboons originated in southern Africa. Nuclear and mitochondrial estimates put the date of initial divergence of baboon lineages at 1.5–2.1 million years ago (Newman et al., 2004; Rogers et al., 2019; Zinner et al., 2013). At about the same time, during the Pleistocene epoch, baboons started to expand their range across sub-Saharan Africa into both northern and southern savannas.

Presently, six species are recognized: the chacma baboon (Papio ursinus), which is found in southern Africa; the yellow baboon (Papio cynocephalus), which inhabits large parts of eastern Africa; the Kinda baboon (Papio kindae), which is found in Zambia, eastern Angola, and southern DR Congo; the olive baboon (Papio anubis), whose distribution ranges from northern DR Congo to parts of Kenya and Tanzania across to Sierra Leone in the west and to Eritrea in the east; the Guinea baboon (Papio papio), which is found from Sierra Leone to Mauritania and Senegal; and the hamadryas baboon (Papio hamadryas), which inhabits parts of Eritrea, Ethiopia, Somalia and the south-western part of the Arabian peninsula (Figure 1). Hybrid zones are found where species’ distributions come into contact.

While the systematic grouping into taxa within the genus Papio is well accepted on both phenotypic and genetic evidence, the taxonomic ranking is disputed. According to the biological species concept (Mayr, 1963), all taxa would be considered one polytypic species (Papio hamadryas) because where they meet in the wild they interbreed freely, producing viable and fertile hybrid offspring. Given that the different taxa vary substantially in appearance, behavior, and the characteristics of their society, we follow the phylogenetic species concept (Cracraft, 1983), and refer to the different taxa as "species".

Figure 2. Illustration of key traits across baboon species. (A) Phenotypic variation between species. Pictures show adult males and females. (B) Crania of male baboons. (C) Sexual swellings of female baboon during peak estrus. Species are grouped by social organization (uni- and multi-level) and dispersal behavior (male- or female-biased dispersal): Images from Alexis Amann, Andrea Cardini, Sarah Elton, Julia Fischer, Courtney Fitzpatrick, James Higham, Megan Petersdorf, Joan Silk and Larissa Swedell. DOI: https://doi.org/10.7554/eLife.50989.004
Box 1. Glossary.

**Admixture**: Genetic admixture refers to the exchange of genetic information among two populations or taxa that had been reproductively isolated and which genetically diverged (see introgressive hybridization).

**Consortship**: Consortships occur when females are sexually receptive and involve a male and female pair who associate in close proximity, often mating repeatedly. Typically, male-female consort pairs travel, feed and rest together. Consortships can last for hours or days.

**Genetic architecture**: Refers to the underlying genetic basis of a phenotypic trait (morphological, physiological, behavioral) and the variation in the respective trait.

**Ghost lineage**: A term from paleontology and phylogenetics. It refers to a phylogenetic lineage that has no fossil record or living representatives, but is inferred to have existed, for example, by whole-genome analyses of related taxa.

**Hybridization**: The interbreeding between two differentiated populations, usually closely related species, resulting in the combination of genetic material from previously isolated gene pools.

**Introgression or introgressive hybridization**: Observed between species or between genetically well-separated populations. It refers to the movement of genes, or gene flow, from one species into the gene pool of another by the repeated backcrossing of interspecific hybrids with one of their parent species.

**Life history**: The life history of an organism is a characterization of its patterns of development, reproduction, aging and mortality. Key measures of primate life history include length of gestation, age at the first occurrence of menstruation, age at first reproduction, number of offspring per litter, number of births per year, interval between births and life span.

**Mating system**: The distribution of matings among sexually active individuals within a social unit. Primate species can be monogamous (mating occurs mostly between pair partners), polyandrous (one female mates with multiple males), polygynous (one male mates with multiple females), or polygynandrous (males and females have multiple mating partners).

**Mitochondrial and nuclear lineages**: Mitochondria, organelles of almost all eukaryotic cells, carry their own genome. In contrast to nuclear genomes, recombination of the mitochondrial genome is a rare event and, since mitochondria are almost exclusively inherited via the maternal lineage, nuclear and mitochondrial genetic lineages can experience independent evolutionary histories. This often results in discordant phylogenies when using sex chromosomes (gonosomal), non-sex chromosomes (autosomal) or mitochondrial markers. Even phylogenies based on different nuclear genes or parts of the nuclear genome can lead to some discordances. Nevertheless, one can use nuclear and mitochondrial lineages to infer different evolutionary events within the evolutionary history of a species.

**Phylogenetic species concept**: This concept defines a species as an irreducible group or cluster whose members are descendants from a common ancestor and who all possess a combination of certain defining derived traits known as apomorphies. Such groups are monophyletic (contrasted with paraphyletic or polyphyletic groups). Reproductive isolation is not a precondition for the definition of species. Since monophyletic groups are often nested, ranking a particular group as a species can be problematic.

**Social organization**: The number of individuals and the composition of a group, including when and where those individuals are distributed. Groups may for instance be stable or reveal a fission-fusion system where the group temporarily splits into smaller sub-groups. Baboon societies may be uni-level (individuals live in a stable group and generally roam together) or multi-level (groups consist of predictable sub-groups, which may in turn consist of smaller sub-groups). An important aspect of the social organization is the dispersal behavior, that is, which sex typically remains in the group into which it was born (i.e. its natal group) and which sex leaves the natal group to breed. In most mammals, females stay (female philopatry) and males leave (male dispersal), which is considered the ancestral state. The timing and type of dispersal has important consequences for the genetic structure of groups.
Morphology

All baboon species are anatomically and morphologically well adapted to a quadrupedal terrestrial lifestyle (Fleagle, 2013). They have dog-like muzzles and males have large canine teeth. Depending on the species, body mass varies between 17 and 30 kg for adult males, and between 10 and 15 kg for females, resulting in a sexual dimorphism in mass ranging between 1.55 and 2.20 (Anandam et al., 2013; Fischer et al., 2017; Swedell, 2011).

The species differ notably in their fur color, body size and sexually selected characteristics, such as the distinct capes which are most pronounced in Guinea and hamadryas baboons but also present in olive baboons. Females of all species develop sexual swellings of the anogenital region when they are fertile. These swellings change throughout the menstrual cycle, such that maximum swelling typically coincides with ovulation (Higham et al., 2008). The size and shape of the swellings varies considerably among species (Figure 2; Petersdorf et al., 2019).

Ecology

All baboon species are largely terrestrial during the day but retreat to sleeping trees or cliffs during the night. They exhibit great ecological flexibility, allowing them to occupy habitats including semi-deserts grasslands, woodland savannas, humid forests, and Afroalpine grasslands over 3,000 meters above sea level (Chala et al., 2019; Fuchs et al., 2018). Baboons eat a broad range of foods, although their diet mainly consists of plants, including fruit, seeds, leaves, and roots. They also eat insects and other arthropods and, occasionally, kill small antelopes, hares, rodents, birds and smaller monkeys (Goffe and Fischer, 2016; Swedell, 2011).

Phylogeography

Phenotypic differences between species are well recognized (Jolly, 1993), and based on their molecular phylogeny, baboons are generally split in two major groups: north and south (Dunn et al., 2013; Frost et al., 2003). However, genetic evidence reveals a complex evolutionary history of the genus Papio. Analysis of mitochondrial DNA yields a phylogeny that includes several major haplogroups or clades – groups of individuals who belong to a specific mitochondrial lineage. These haplogroups reflect the geographic origin of the respective specimens better than their external phenotypes or taxonomic classification, making species appear to be paraphyletic and polyphyletic when mapped onto the mitochondrial phylogeny (Zinner et al., 2009; Zinner et al., 2011).

Comparisons of whole genome sequences confirm the six baboon species taxonomy and suggest that the initial evolutionary divergence separated a southern lineage that ultimately produced Kinda, chacma and yellow baboons, from a northern lineage that produced olive, hamadryas and Guinea baboons (Rogers et al., 2019). Ancient hybridization events appear to have affected the genetic makeup of all species. For instance, Guinea baboons most likely experienced genetic admixture with a “ghost lineage” that is probably extinct, or that is at least not represented in the sample of genomes analyzed to date (Rogers et al., 2019).

Multiple episodes of admixture and introgression have been linked to climate change and range expansion (Rogers et al., 2019; Walker et al., 2017; Wall et al., 2016; Zinner et al., 2013). Similar evolutionary mechanisms, including gene transfer by introgressive hybridization, are now recognized to have influenced the evolution of Neanderthals, Denisovans and modern humans (Green et al., 2010; Prüfer et al., 2017; Prüfer et al., 2014; Reich et al., 2010; Ackermann et al., 2019). However, the absence of genomic data from
The six baboon species vary substantially in their social characteristics, including social organization, social style and mating patterns. Group size varies within and among species. In chacma, olive and yellow baboons, group size ranges from about a dozen up to roughly one hundred animals (Markham et al., 2015; Swedell, 2011), while hamadryas and Guinea baboons temporarily aggregate into groups of several hundreds of individuals (Patzelt et al., 2011; Swedell, 2013). Sex ratios in baboons vary too; some groups are fairly balanced, while adult females in other groups can outnumber adult males by about 10 to 1 (Swedell, 2011).

Chacma, olive, Kinda and yellow baboons – recently dubbed “COKY” baboons (Jolly, 2019) – live in multi-male-multi-female groups, in which related females constitute the stable core, while males leave the group they were born into and join another. Clear rank hierarchies among males and females can be discerned based on aggressive interactions, including threats, chases and physical aggression, as well as signals of submission. In females, related individuals (known as matrilines) typically occupy adjacent ranks. For female chacma, olive and yellow baboons, female kin constitute the most important social partners (Silk et al., 2017; Silk et al., 2010; Silk, 2003). In Kinda baboons, however, males are the most significant grooming partners for females (Petersdorf et al., 2019).

Females of all COKY species interact and mate with several males in the group. High-ranking males generally experience higher mating success than lower-ranking ones, though this rank-related mating skew is more pronounced in chacma baboons than in olive or yellow baboons (Bulger, 1993; Henzi and Barrett, 1999; Stådele et al., 2019). During female receptive periods, males aggressively guard their female mating partner, resulting in sexual “consortships” (Noé and Sluijter, 1990; Smuts, 1985). Consorts may last from several hours up to several days. Consort success (and thus mating success) is often related to male dominance status (Gesquiere et al., 2011). In yellow and olive baboons, however, male coalitions may be able to take the female away from a dominant male (Noé and Sluijter, 1990; Smuts, 1985).

Male competition and aggressiveness vary considerably between species. Infanticide is frequent in some populations of chacma baboons (Palombit et al., 2001), but rare in olive, yellow and hamadryas baboons (Lemasson et al., 2008; Swedell, 2011). Lactating females often form close ties to specific males, which are often the sires of their infants (Huchard et al., 2010; Moscovice et al., 2010; Nguyen et al., 2009; Stådele et al., 2019). These relationships appear to be an adaptation against infanticide by recent immigrant males (Palombit et al., 1997) and harassment by other group members (Alberts et al., 2003) as well as a form of paternal investment (Buchan et al., 2003; Huchard et al., 2013). Male alliances are absent in chacma baboons, while common in yellow and olive baboons (Noé and Sluijter, 1995). These differences in male competitive regimes are reflected in their dispersal behavior: male chacma baboons in the Okavango delta, for instance, do not emigrate from their natal group until after they are fully grown (Beehner et al., 2009), while male olive and yellow baboons often emigrate during adolescence (Alberts and Altmann, 1995; Packer et al., 1995).
Over the past decade, studies of Kinda baboons have broadened our knowledge of morphological and behavioral variation within the genus. Kinda baboons are smaller in body size, have reduced sexual dimorphism in body and canine size, and have larger relative testis volume, compared to other baboon species (Jolly, 2019). Kinda females exhibit small sexual swellings (Figure 2) and give inconspicuous calls (Petersdorf et al., 2019). Chacma females, in contrast, give loud copulation calls, which function to incite male-male competition (O’Connell and Cowlishaw, 1994).

Hamadryas baboons — in contrast to the COKY baboon species described above — live in a multi-level society with reproductive units, called “one male units” comprising one sexually active leader male, a variable number of females, and sometimes a follower male (Kummer, 1968). Associations between several one-male units constitute a clan (Abeglen, 1984); several clans and unaffiliated bachelor males form a band, the main ecological unit, and multiple bands coalesce at resources, especially sleeping sites, to form troops (Schreier and Swedell, 2009). Recent behavioral and genetic studies of hamadryas baboons show that leader and follower males tend to be maternally related, in line with the fact that they disperse only rarely. Females within a unit are also more likely to be related than expected by chance (Städele et al., 2016).

Guinea baboons also live in a multi-level society. Several units comprising a primary male, 1–6 females, young, and occasional secondary males make up parties, and 2 to 3 parties constitute a gang within a larger community (Fischer et al., 2017). Male Guinea baboons maintain strong bonds and a high degree of spatial tolerance (Fischer et al., 2017). Some, but not all males with strong bonds are highly related, suggesting that the existence of kin in the group promotes male tolerance (Patzelt et al., 2014). In striking contrast to most other baboon species, aggression between males is so rare that it is not possible to discern a dominance hierarchy with certainty (Dal Pesco and Fischer, 2018). Males engage in extended ritualized greetings that apparently function to reinforce delineations between parties and to test bonds between males (Dal Pesco and Fischer, 2018). Females freely transfer between units, parties and gangs. Female tenure with a given male may vary between weeks and years (Goffe et al., 2016). Both Guinea and hamadryas baboons exhibit female-biased dispersal (Kopp et al., 2015; Städele et al., 2015).

Note that many of the most significant differences in social behavior between species have been observed across different populations in multiple African sites, as well as in captivity. Thus, there is good evidence that the variation we describe here reflects true species differences and not just variation between populations. Yet, characterizing the variation within species in greater detail would be extremely valuable.

Despite the variation in social organization and aggresiveness between the different baboon species, there is very little variation in the vocal repertoires and call types within the genus (Hammerschmidt and Fischer, 2019). This suggests that the structure of vocal patterns is highly conserved. Because species vary in their aggressiveness and their propensity to affiliate, they also differ in the frequency with which they use signals that either relate to fighting ability or “benign intent”, respectively (Faraut et al., 2019; Fischer et al., 2017).

**Variation in social cognition**

Variation in social organization and in the nature and extent of competition over resources between baboon species is thought to result in differential selective pressure on social cognition (Amici et al., 2008; Aureli et al., 2008). To date, most of the work on baboon social knowledge has been done on chacma baboons that exhibit steep dominance hierarchies (known as despotism). A suite of studies by the American primatologists Dorothy Cheney and Robert Seyfarth and colleagues revealed that chacma baboons have sophisticated social knowledge (reviewed in Cheney and Seyfarth, 2008). For instance, the animals represent the nested hierarchical rank relationships of their group members (Bergman, 2003), track the consortship status of pairs in their group (Crockford et al., 2007), and selective deploy aid to unrelated individuals that were former grooming partners (Cheney et al., 2010).

Field playback experiments revealed that baboon species respond differently to social information. While the territorial chacma baboons respond strongly to apparent intruders (Kitchen et al., 2013), the spatially tolerant Guinea baboons paid more attention to vocalizations from co-resident group members compared to neighbors or strangers (Maciej et al., 2013). Similarly, chacma baboons respond strongly to simulated rank reversals.
(Bergman, 2003) or break-ups of existing consortships (Crockford et al., 2007), while Guinea baboon males were more interested in social information consistent with current social association patterns (Faraut and Fischer, 2019). The somewhat surprising responses of the Guinea baboons may be a result of the high gregariousness of the species, where deviant interaction patterns may initially be classified as “social noise” (Faraut and Fischer, 2019). In summary, these findings suggest that the content of what is represented, namely the associations between different individuals or their group memberships, appears to be relatively similar across the two species, while the value of different types of social information may vary substantially in relation to the type of society.

Sociality, health, aging and fitness
Over the past decade, baboon research has provided ground-breaking insights into the relationships between social status, social relationships, health and fitness measures such as offspring survival and longevity. Data from two long-term studies of baboon behavior and life history suggest that sociality enhances the fitness of females. For example, infants born to yellow baboon females who are more socially integrated have higher survival than infants of less social mothers (Archie et al., 2014; Silk, 2003); similar patterns are also seen in chacma baboons (Silk et al., 2010). As in many other primates, higher-ranking male baboons sire more offspring than other males (Altmann et al., 1996). Higher-ranking females have shorter periods before they resume menstrual cycling following birth (Gesquiere et al., 2018; Johnson, 2003; Packer et al., 1995; Smuts and Nicolson, 1989; Wasser et al., 2004), which may be linked to quicker restoration of positive energy balance (Gesquiere et al., 2018). Consistent with this, feeding on crops in olive baboons (Higham et al., 2009), or discarded human food for yellow baboons (Altmann et al., 1977), also leads to a quicker return to menstrual cycling and increases reproductive output.

A number of studies have investigated the proximate mediators of the relationship between behavior and fitness. In particular, many researchers have taken advantage of non-invasive ways to measure glucocorticoid hormones, a class of hormones known to mediate the energetic demands that accompany social and ecological challenges. Concentrations of glucocorticoid hormones increase during specific challenges that are known to threaten an individual’s fitness. For example, lactating chacma females that were at risk for infanticide because a new male immigrated into the group exhibited elevated glucocorticoid hormones compared to female counterparts that were not at risk (Beehler et al., 2005). Additionally, loss of a close female relative increases glucocorticoid concentrations, and this increase may be responsible for initiating a compensatory broadening and strengthening of female grooming networks (Engh et al., 2006).

Several studies have investigated the relationship between glucocorticoid concentrations, rank and social stability in male baboons. In a long-term study of yellow baboons, high-ranking males had lower glucocorticoid concentrations, regardless of hierarchy stability, while alpha males may experience higher concentrations than expected for their rank (Gesquiere et al., 2011). Nonetheless, high-ranking yellow baboon males get sick less often and heal from wounds faster, suggesting that these high-ranking males are in better health and do not suffer trade-offs from these extra demands (Archie et al., 2012). Higher-ranking chacma baboon males also had higher glucocorticoid concentrations (Bergman et al., 2005; Kalbitzer et al., 2015).

More recently, baboons were also established as a promising model for studying the impact of sexually transmitted diseases on mating behavior. Female olive baboons in a population infected with the bacterium Treponema pallidum, a subspecies of which causes syphilis in humans, copulated less with males showing clinical signs of infection (Paciência et al., 2019). These findings highlight how pathogens may impose important selective pressures in mate choice and ultimately social evolution.

Functional genomics
The addition of data on Kinda and Guinea baboons increases the value of the baboon as a model, as we begin to have data available for all baboon species. While one aim of future analyses will be to understand the sources of variation between species, documenting similarities is equally valuable. Technological developments in genomic sequencing (Robinson et al., 2019; Rogers et al., 2019), including from non-invasively collected samples such as feces (Perry et al., 2010; Snyder-Mackler et al., 2016), have brought genomics to the forefront of baboon behavioral studies (Tung et al., 2010). Given the close phylogenetic relatedness
of the six baboon species, variation in key aspects of social behavior, and the presence of hybrids displaying intermediate phenotypes within hybrid zones, investigation of causal pathways from genotype to phenotype seems particularly promising within the baboon model (Bergey et al., 2016; Jolly et al., 2008).

Formerly, research into primate behavioral genetics focused on identifying a few small specific functional polymorphisms in sequence or length, and on linking these to phenotypic variation (e.g., Kalbitzer et al., 2016). However, such studies are likely to overestimate the effect of one single aspect of genetic variation. With genotyping of single nucleotide polymorphisms (SNPs), and whole-genome sequencing, primate field studies are beginning to explore the wider genomic architecture that underlies variation in social behavior (Rogers, 2018). As well as whole genome sequence data, researchers now have access to annotations for protein coding genes and transcriptomes (Robinson et al., 2019; Rogers et al., 2019; Vilgalys et al., 2019). We therefore expect an exponential increase in the number and diversity of available genomes, which will facilitate the investigation of the basis of baboon adaptations and adaptive flexibility.

In conjunction with research on other model organisms, such as deer mice (Bedford and Hoekstra, 2015), such studies provide fundamental insights into the foundation of natural variation and adaptation in socially living mammals.

### Baboons in the Anthropocene

Baboons allow us to study the effects of accelerating anthropogenic fragmentation, loss of natural habitats and climate change in a highly adaptable primate system. For example, baboons may rapidly change how long they allocate time and energy to different behaviors or where they range, in response to human-related activities and habitat changes (Fehlmann et al., 2017; Ferreira da Silva et al., 2018). Studies of individual baboon behavior can use sophisticated GPS tracking and non-invasive genetic tools to make broad-scale inferences about movements and processes at the population level (Kopp et al., 2014; Strandburg-Peshkin et al., 2015). These inferences can then be applied to questions of how other large populations of mammals will respond to changes in their environment.

Baboons are not considered a global priority in conservation, with the exception of Guinea baboons which are categorized as Near Threatened by the IUCN (Oates et al., 2008). However, populations geographically overlapping with human communities often damage crops and infrastructure and are persecuted as pests. In some locations, people consume substantial number of baboons and sell their meat in bushmeat markets (e.g., Minhós et al., 2013). Humans and baboons often compete for space and hunting of specific individuals or even entire groups is increasingly frequent, leading to fragmented populations and local extinctions (Ferreira da Silva et al., 2018). Non-monitored populations living outside protected areas may be at a higher risk of silently disappearing. The challenge is to assess the risk of different

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**Box 2. Outstanding questions about the natural history of baboons.**

- How have changes in population density and environmental conditions (e.g., opening and closing of forest and other barriers) affected dispersal and mating patterns, and ultimately given rise to different social systems?
- What is the genetic architecture of baboon social behavior (including social style, patterns of dispersal, and degree of reproductive skew according to social status)? How does that architecture promote or restrict evolutionary flexibility in social systems?
- Does the link between sociality and reproductive success vary among species or even local populations?
- Do the different species vary in the way they represent the social relationships around them and how they attend to social information?
- How responsive are baboons to changes in temperature patterns due to global warming, as well as to associated changes in aridity or habitat type?

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populations and develop appropriate conservation plans.

The long-term nature of many baboon field studies has provided great insight into how populations may rise and fall rapidly in response to changes in the environment. The Amboseli Baboon Project, for example, has been running continuously for 50 years, and has documented numerous periods of relative drought or rainfall abundance (Alberts et al., 2005; Alberts and Altmann, 2011). This variation in precipitation has been linked to variation in fecundity and survival (Beehner et al., 2006; Lea et al., 2015) and to subsequent changes in population structure (Altmann et al., 1985). Periods of environmental change, and consequent boom and bust cycles in populations, are driven by both natural phenomena, such as natural aging of woodland, and anthropogenic influences, such as overgrazing by pastoralists (Alberts and Altmann, 2011). Many long-term baboon field sites also carefully collect detailed data on temperature and rainfall, as well as food availability and diseases. They also monitor the habitats in addition to the baboon populations. The breadth and scope of such data ensure that the baboon represents an outstanding model of both individual-level and population-level responses to environmental change.

**Conclusion**

Baboons constitute a fascinating and informative analog model for hominin evolution in savanna habitats, with their ongoing patterns of range expansions and contractions, and regular occurrences of hybridization where two species meet. Given the availability of long-term data and the variation in the types of societies baboons live in, they constitute an excellent test case to study the link between sociality, health, longevity and reproductive success, as well as the emergence and spread of diseases. Such studies are extremely important to put biomedical data from captive baboon studies into natural context. For future research, we propose an approach that integrates field observations and carefully designed field experiments with cutting-edge measures of genomic variation, gene expression, non-invasive endocrinology and immunology. The fact that baboons have been studied in a wide range of habitats at sites across Africa for several decades also make them an informative example to investigate how populations of large mammals respond to environmental diversity and change (see Box 2 for suggested future research questions).

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References

Abegglen JJ. 1984. On Socialization in Hamadryas Baboons: A Field Study. Bucknell University Press.

Ackermann RR, Arnold ML, Baiz MD, Cahill JA, Cortés-Ortiz L, Evans BJ, Grant BR, Grant PR, Hallgrimson B, Humphreys RA, Jolly CJ, Malukiewicz J, Percival CJ, Ritzman TB, Roos C, Roseman CC, Schroeder L, Smith FH, Warren KA, Wayne RK, et al. 2019. Hybridization in human evolution: insights from other organisms. Evolutionary Anthropology: Issues,
News, and Reviews 28:189–209. DOI: https://doi.org/10.1002/evan.21787

Alberts SC, Watts HE, Altmann J. 2003. Queueing and queue-jumping: long-term patterns of reproductive skew in male savannah baboons, Papio cynocephalus. Animal Behaviour 65:821–840. DOI: https://doi.org/10.1016/S0003-3472(03)00216-0

Alberts SC, Hollister-Smith JA, Mututua RS, Sayiuel SN, Muruthi PM, Kinyua Warutere J, Altmann J. 2005. Seasonality and long-term change in a savanna environment. In: Brockmann D, K, van Schaik C P (Eds). Seasonality in Primates. Cambridge University Press. p. 157–195. DOI: https://doi.org/10.1017/CBO9780511542343.007

Alberts SC, Altmann J. 1995. Balancing costs and opportunities: dispersal in male baboons. The American Naturalist 145:279–306. DOI: https://doi.org/10.1086/285740

Alberts SC, Altmann J. 2011. The Amboseli Baboon Research Project: 40 Years of Continuity and Change. In: Kappeler P, Watts D (Eds). Long-Term Field Studies of Primates. Springer Science & Business Media, p. 261–287. DOI: https://doi.org/10.1007/978-3-642-22514-7_12

Alfred J, Baldwin IT. 2015. New opportunities at the wild frontier. eLife 4:e06956. DOI: https://doi.org/10.7554/eLife.06956

Altmann J, Altmann SA, Hausfater G, McCuskey SA. 1977. Life history of yellow baboons: physical development, reproductive parameters, and infant mortality. Primates 18:315–330. DOI: https://doi.org/10.1007/BF02383111

Altmann J, Hausfater G, Altmann SA. 1985. Demography of amboseli baboons, 1963-1983. American Journal of Primatology 8:113–125. DOI: https://doi.org/10.1080/03071758509448734

Altmann J, Alberts SC, Haines SA, Dubach J, Muruthi P, Coote T, Geffen E, Cheesman DJ, Mututua RS, Sayiuel SN, Wayne RK, Lacy RC, Bruford MW. 1996. Behavior predicts genes structure in a wild primate group. PNAS 93:5797–5801. DOI: https://doi.org/10.1073/pnas.93.12.5797

Amici F, Aureli F, Call J. 2008. Fission-fusion dynamics, behavioral flexibility, and inhibitory control in primates. Current Biology 18:1415–1419. DOI: https://doi.org/10.1016/j.cub.2008.08.020

Anandam MV, Bennett EL, Davenport TRB, Davies NJ, Detwiler KM, Engelhardt A, Eudey AA, Gadsby EL, Groves CP, Healy A, Karanth KP, Molor S, Nadler T, Richardson MC, Riley EP, Roos C, Rylands AB, Sheeran LK, Ting N, Wallis J, et al. 2013. Species accounts of Cercopithecidae. In: Mittermeier R, Rylands A, Wilson DS (Eds). Handbook of the Mammals of the World. 3 Barcelona, Primates: Lynx. 628–753.

Archie EA, Altmann J, Alberts SC. 2012. Social status predicts wound healing in wild baboons. PNAS 109:9017–9022. DOI: https://doi.org/10.1073/pnas.1206391109

Archie EA, Tung J, Clark M, Altmann J, Alberts SC. 2014. Social affiliation matters: both same-sex and opposite-sex relationships predict survival in wild female baboons. Proceedings of the Royal Society B: Biological Sciences 281:20141261. DOI: https://doi.org/10.1098/rspb.2014.1261

Aureli F, Schaffner CM, Boesch C, Bearder SK, Call J, Chapman CA, Connor R, Fiore AD, Dunbar RIM, Henzi SP, Holekamp K, Korstjens AH, Layton R, Lee P, Lehmann J, Manson JH, Ramos-Fernandez G, Stier KB, Schaik CPvan. 2008. Fission-fusion dynamics. Current Anthropology 49:627–654. DOI: https://doi.org/10.1086/588708

Bedford NL, Hoekstra HE. 2015. Peromyscus mice as a model for studying natural variation. eLife 4:e06813. DOI: https://doi.org/10.7554/eLife.06813

Beehner JC, Bergman TJ, Cheney DL, Seyfarth RM, Whitten PL. 2005. The effect of new alpha males on female stress in free-ranging baboons. Animal Behaviour 69:1211–1221. DOI: https://doi.org/10.1016/j.anbehav.2004.08.014

Beehner JC, Onderdonk DA, Alberts SC, Altmann J. 2006. The ecology of conception and pregnancy failure in wild baboons. Behavioral Ecology 17:741–750. DOI: https://doi.org/10.1093/beheco/arl006

Beck J, Esquiere L, Seyfarth RM, Cheney DL, Alberts SC, Altmann J. 2009. Testosterone related to age and life-history stages in male baboons and geladas. Hormones and Behavior 56:472–480. DOI: https://doi.org/10.1016/j.yhbeh.2009.08.005

Bergen CM, Phillips-Conroy JE, Disotell TR, Jolly CJ. 2016. Dopamine pathway is highly diverged in primate species that differ markedly in social behavior. PNAS 113:6178–6181. DOI: https://doi.org/10.1073/pnas.1525530113

Bergman TJ. 2003. Hierarchical classification by rank and kinship in baboons. Science 302:1234–1236. DOI: https://doi.org/10.1126/science.1087513

Bergman TJ, Beehner JC, Cheney DL, Seyfarth RM, Whitten PL. 2005. Correlates of stress in free-ranging male chacma baboons, Papio hamadryas ursinus. Animal Behaviour 70:703–713. DOI: https://doi.org/10.1016/j.anbehav.2004.12.017

Buchan JC, Alberts SC, Silk JB, Altmann J. 2003. True paternal care in a multi-male primate society. Nature 425:179–181. DOI: https://doi.org/10.1038/nature01866

Bulger JB. 1993. Adult Male Savanna Baboon Mating Activity and Interactions with Infants. Springer.

Chada D, Roos C, Svenning J-C, Zinner D. 2019. Species-specific effects of climate change on the distribution of suitable baboon habitats – Ecological niche modeling of current and Last Glacial Maximum conditions. Journal of Human Evolution 132:215–226. DOI: https://doi.org/10.1016/j.jhevol.2019.05.003

Cheney DL, Moscovice LR, Heesen M, Mundy R, Seyfarth RM. 2010. Contingent cooperation between wild female baboons. PNAS 107:9562–9566. DOI: https://doi.org/10.1073/pnas.1001862107

Cheney DL, Seyfarth RM. 2008. Baboon Metaphysics: The Evolution of a Social Mind. University of Chicago Press. DOI: https://doi.org/10.7208/chicago/9780226102429.001.0001

Crockford C. 1993. Species concepts and speciation analysis. Current Ornithology:159–187. DOI: https://doi.org/10.1007/978-1-4615-6781-3_6

Crocket CF, Wittig RM, Seyfarth RM, Cheney DL. 2007. Baboons eavesdrop to deduce mating opportunities. Animal Behaviour 73:885–890. DOI: https://doi.org/10.1016/j.anbehav.2006.10.016

Dal Pesco F, Fischer J. 2018. Greetings in male guinea baboons and the function of rituals in complex social groups. Journal of Human Evolution 125:87–98. DOI: https://doi.org/10.1016/j.jhevol.2018.10.007
et al. eLife 2019;8:e50989. DOI: https://doi.org/10.7554/eLife.50989

Fischer et al. eLife 2019;8:e50989. DOI: https://doi.org/10.7554/eLife.50989

Fischer, J., Kopp, G.H., Dals Pesco, F., Goffe, A., Hammerschmidt, K., Kalbitzer, U., Klapproth, M., Maciej, P., Ndao, I., Patzelt, A., Zinner, D. 2017. Charting the neglected west: the social system of Guinea baboons. American Journal of Physical Anthropology 162:15–31. DOI: https://doi.org/10.1002/ajpa.23144

Fleagle, J.G. 2013. Old World monkeys. In: Primates Adaptation and Evolution. 3rd Edition. London: Academic Press. p. 119–150. DOI: https://doi.org/10.1016/C2009-0-01979-5

Frost, S.R., Marcus, L.F., Bookstein, F.L., Reddy, D.P., Delson, E. 2003. Cranial allometry, phyleogeography, and systematics of large-bodied papionins (primates: cercopithecinae) inferred from geometric morphometric analysis of landmark data. The Anatomical Record 275A:1048–1072. DOI: https://doi.org/10.1002/ar.a.10112

Fuchs, A.J., Gilbert, C.C., Kamil, J.M. 2018. Ecological niche modeling of the genus Papio. American Journal of Physical Anthropology 166:812–823. DOI: https://doi.org/10.1002/ajpa.23470. PMID: 29607482

Gesquiere, L.R., Lear, N.H., Simao, M.C.M., Onyango, P.O., Alberts, S.C., Altmann, J. 2011. Life at the top: rank and stress in wild male baboons. Science 333:357–360. DOI: https://doi.org/10.1126/science.1207120

Gesquiere LR, Altmann J, Archie EA, Alberts SC. 2018. Interbirth intervals in wild baboons: environmental predictors and hormonal correlates. American Journal of Physical Anthropology 166:107–126. DOI: https://doi.org/10.1002/ajpa.23407

Goffe AS, Zinner D, Fischer J. 2016. Sex and friendship in a multilevel society: behavioral patterns and associations between female and male Guinea baboons. Behavioral Ecology and Sociobiology 70:323–336. DOI: https://doi.org/10.1007/s00265-015-2050-6

Goffe AS, Fischer J. 2016. Meat sharing between male and female Guinea baboons (Papio papio). Primate Biology 3:1–8. DOI: https://doi.org/10.5194/pb-3-1-2016

Green RE, Krause J, Briggs AW, Maricic T, Stenzel U, Kircher M, Patterson N, Li H, Zhai W, Fritz MH, Hansen NF, Durand EY, Malaspinas AS, Jensen JD, Marques-Bonet T, Alkan C, Pruefer K, Meyer M, Burbano HA, Good JM, et al. 2010. A draft sequence of the neanderthal genome. Science 328:710–722. DOI: https://doi.org/10.1126/science.1188021

Hammerschmidt K, Fischer J. 2019. Baboon vocal repertoires and the evolution of primate vocal diversity. Journal of Human Evolution 126:1–13. DOI: https://doi.org/10.1016/j.jhevol.2018.10.010

Henzi SP, Barrett L. 1999. The historical socioecology of savanna baboons (Papio hamadryas). Journal of Zoology 265:215–226. DOI: https://doi.org/10.1017/S0952836904006399

Higham JP, MacLarnon AM, Ross C, Heistermann M, Semple S. 2008. Baboon sexual swellings: information content of size and color. Hormones and Behavior 53:452–462. DOI: https://doi.org/10.1016/j.yhbeh.2007.11.019

Higham JP, Warren Y, Adanu J, Umaru BN, MacLarnon AM, Sommer V, Ross C. 2009. Living on the edge: life-history of olive baboons at Gashaka-Gumti National Park, Nigeria. American Journal of Primatology 71:293–304. DOI: https://doi.org/10.1002/ajp.20651

Huchard E, Alvergne A, Féjan D, Knapp LA, Cowlishaw G, Raymond M. 2010. More than friends? Behavioural and genetic aspects of heterosexual associations in wild chacma baboons. Behavioral Ecology and Sociobiology 64:769–781. DOI: https://doi.org/10.1007/s00265-009-0894-3

Huchard E, Charpentier MJ, Marshall H, King AJ, Knapp LA, Cowlishaw G. 2013. Paternal effects on access to resources in a promiscuous primate society. Behavioral Ecology 24:229–236. DOI: https://doi.org/10.1093/beheco/ars158

Johnson SE. 2003. Life history and the competitive environment: trajectories of growth, maturation, and reproductive output among chacma baboons. American Journal of Physical Anthropology 120:83–98. DOI: https://doi.org/10.1002/ajpa.10139

Jolly CJ. 1993. Species, subspecies, and baboon systematics. In: Kimbel WH (Ed). Species Concepts and Primate Evolution. Boston, MA: Springer. p. 67–107. DOI: https://doi.org/10.1007/978-1-4899-3745-2_4

Jolly CJ, Phillips-Conroy JE, Kaplan JR, Mann JJ. 2008. Cerebrospinal fluid monoaminergic metabolites in wild Papio anubis and P. hamadryas are concordant with Taxon-specific behavioral ontogeny. International
Journal of Primatology 29:1549–1566. DOI: https://doi.org/10.1076/0176-4060(94)90144-3

Jolly CJ. 2019. Philopatry at the frontier: a demographically-driven scenario for the evolution of multi-level societies in baboons (Papio). Journal of Human Evolution. In press.

Kallbitzer U, Heistermann M, Cheney D, Seyfarth R, Fischer J. 2015. Social behavior and patterns of testosterone and glucocorticoid levels differ between male chacma and Guinea baboons. Hormones and Behavior 75:100–110. DOI: https://doi.org/10.1016/j.yhbeh.2015.08.013

Kallbitzer U, Roos C, Kopp GH, Butynski TM, Knauf S, Zinner D, Fischer J. 2016. Insights into the genetic foundation of aggression in Papio and the evolution of two length-polymorphisms in the promoter regions of serotonin-related genes (5-HTTLPR and MAOALPR) in Papionini. BMC Evolutionary Biology 16:121.

DOI: https://doi.org/10.1186/s12862-016-0693-1

Kappeker PM. 2019. A framework for studying social coalitions. Behavioral Ecology and Sociobiology 73:12. DOI: https://doi.org/10.1007/s00265-018-2601-8

Kitchen DM, Seyfarth RM, Cheney DL. 1997. The structure and function of a multilevel primate society. Cambridge, MA: Harvard University Press.

Kummer H. 1968. Social Organization of Hamadryas Baboons. University of Chicago Press.

Lea AJ, Altmann J, Alberts SC, Tung J. 2015. Developmental constraints in a wild primate. The American Naturalist 185:809–821. DOI: https://doi.org/10.1088/681016

Lemasson A, Palombit RA, Jubin R. 2008. Friendships between males and lactating females in a free-ranging group of olive baboons (Papio hamadryas anubis): evidence from playback experiments. Behavioral Ecology and Sociobiology 62:1027–1035. DOI: https://doi.org/10.1007/s00265-007-0530-z

Maciej P, Patzelt A, Ndou I, Hammerschmidt K, Fischer J. 2013. Social monitoring in a multilevel society: a playback study with male Guinea baboons. Behavioral Ecology and Sociobiology 67:61–68.

DOI: https://doi.org/10.1007/s00265-012-1425-1

Marais EN. 1939. My Friends the Baboons.London: Methuen.

Markham AC, Gesquiere LR, Alberts SC, Altmann J. 2015. Optimal group size in a highly social mammal. PNAS 112:14882–14887. DOI: https://doi.org/10.1073/pnas.1511779112

Mayr E. 1963. Animal Species and Evolution. Cambridge, MA: Harvard University Press.

Minhós T, Wallace E, Ferreira da Silva MJ, Sá RM, Carmo M, Barata A, Brufورد MW. 2013. DNA identification of primate bushmeat from urban markets in Guinea-Bissau and its implications for conservation. Biological Conservation 167:43–49. DOI: https://doi.org/10.1016/j.biocon.2013.07.018

Moscoveci LR, Di Fiore A, Crockford C, Kitchen DM, Wittig R, Seyfarth RM, Cheney DL. 2010. Hedging their bets? male and female chacma baboons form friendships based on likelihood of paternity. Animal Behaviour 79:1007–1015. DOI: https://doi.org/10.1016/j.anbehav.2010.01.013

Newman TK, Jolly CJ, Rogers J. 2004. Mitochondrial phylogeny and systematics of baboons (Papio). American Journal of Physical Anthropology 124:17–27.

DOI: https://doi.org/10.1002/ajpa.10340

Nguyen N, Van Horn RC, Alberts SC, Altmann J. 2009. “Friendships” between new mothers and adult males: adaptive benefits and determinants in wild baboons (Papio cynocephalus). Behavioral Ecology and Sociobiology 63:1331–1344. DOI: https://doi.org/10.1007/s00265-009-0786-6

Noëi R, Slijjer AA. 1990. Reproductive tactics of male savanna baboons. Behaviour 113:117–169.

DOI: https://doi.org/10.1163/156853999X00458

Noëi R, Slijjer AA. 1995. Which adult male savanna baboons form coalitions? International Journal of Primatology 16:77–105. DOI: https://doi.org/10.1007/BF02700154

O’Connell SM, Cowlishaw G. 1994. Infanticide avoidance, sperm competition and mate choice: the function of copulation calls in female baboons. Animal Behaviour 48:687–694. DOI: https://doi.org/10.1006/anbe.1994.1288

Oates JF, Gippoliti S, Groves CP. 2008. Papio Papio, Guinea Baboon: IUCN Red List of Threatened Species. DOI: https://doi.org/10.2305/IUCN.UK.2008.RLTS.T16018A5354225.en

Palombit RA, Patzelt A, Kopp GH, Ndao I, Kalbitzer U, Zinner D, Fischer J, Brito JC, Van Horn RC, Alberts SC, Altmann J, Kalbitzer U, Zinner D, Fischer J, Brito JC, Van Horn RC, Alberts SC, Altmann J. 2009. “Friendships” between new mothers and adult males: adaptive benefits and determinants in wild baboons (Papio cynocephalus). Behavioral Ecology and Sociobiology 63:1331–1344. DOI: https://doi.org/10.1007/s00265-009-0786-6

Noëi R, Slijjer AA. 1995. Which adult male savanna baboons form coalitions? International Journal of Primatology 16:77–105. DOI: https://doi.org/10.1007/BF02700154

O’Connell SM, Cowlishaw G. 1994. Infanticide avoidance, sperm competition and mate choice: the function of copulation calls in female baboons. Animal Behaviour 48:687–694. DOI: https://doi.org/10.1006/anbe.1994.1288

Oates JF, Gippoliti S, Groves CP. 2008. Papio Papio, Guinea Baboon: IUCN Red List of Threatened Species. DOI: https://doi.org/10.2305/IUCN.UK.2008.RLTS.T16018A5354225.en

Palombit RA, Seyfarth RM, Cheney DL. 1997. The adaptive value of ‘friendships’ to female baboons: experimental and observational evidence. Animal Behaviour 54:599–614. DOI: https://doi.org/10.1006/anbe.1996.0457

Palombit RA, Cheney DL, Seyfarth RM. 2001. Female–females compete for male ‘friends’ in wild chacma baboons (Papio cynocephalus ursinus). Animal Behaviour 61:1159–1171. DOI: https://doi.org/10.1006/anbe.2000.1690

Patzelt A, Zinner D, Fickenscher G, Diedhiou S, Camara B, Stahl D, Fischer J. 2011. Group composition of guinea baboons (Papio papio) at a water place suggests a fluid social organization. International Journal of Primatology 32:652–668. DOI: https://doi.org/10.1007/s10764-011-9493-z

Patzelt A, Kopp GH, Ndao I, Kallbitzer U, Zinner D, Fischer J. 2014. Male tolerance and male–male bonds in a multilevel primate society. PNAS 111:14740–14745. DOI: https://doi.org/10.1073/pnas.1405811111

Perry GH, Marioni JC, Melsted P, Gilad Y. 2010. Genomic-scale capture and sequencing of endogenous DNA from feces. Molecular Ecology 19:5332–5344.
enhance infant survival. eLife 2019;8:e50989. DOI:https://doi.org/10.7554/eLife.50989

Petersdorf M, Weyher AH, Kamilar JM, Dubuc C, Higham JP. 2019. Sexual selection in the Kinda baboon. Journal of Human Evolution 135:102635. DOI:https://doi.org/10.1016/j.jhevol.2019.06.006

Phifer-Rixey M, Nachman MW. 2015. Insights into mammalian biology from the wild house mouse Mus musculus. eLife 4:e05959. DOI:https://doi.org/10.1013/nature12886

Prüfer K, de Filippo C, Grote S, Mafessoni F, Korlević P, Hajdinjak M, Vernot B, Skov L, Hsieh P, Peyrégné S, Reher D, Hoppe C, Nagel S, Maricic T, Fu Q, Theunert C, Rogers R, Skoglund P, Chintalapati M, Dannemann M, et al. 2017. A high-coverage Neandertal genome from Vindija Cave in Croatia. Science 358:655–658. DOI:https://doi.org/10.1126/science.aao1887

Reich D, Green RE, Kircher M, Krause J, Patterson N, Durand EY, Viola B, Briggs AW, Stenzel U, Johnson PLF, Maricic T, Good JM, Marques-Bonet T, Alkan C, Fu Q, Mallick S, Li H, Meyer M, Eichler EE, Green RE, Kircher M, Krause J, Patterson N, Durand EY, Viola B, Briggs AW, Stenzel U, Johnson PLF, Maricic T, Good JM, Marques-Bonet T, Alkan C, Fu Q, Mallick S, Li H, Meyer M, Eichler EE, Stoneking M, et al. 2010. Genetic history of an archaic hominin group from Denisova cave in Siberia. Nature 468: 1053–1060. DOI:https://doi.org/10.1038/nature09710

Robinson JA, Belsare S, Birnbaum S, Newman DE, Chan J, Glenn JP, Ferguson B, Cox LA, Wall JD. 2019. Analysis of 100 high-coverage genomes from a pedigreed captive baboon colony. Genome Research 29:848–856. DOI:https://doi.org/10.1101/gr.247122.118

Rogers J. 2018. The behavioral genetics of nonhuman primates: status and prospects. American Journal of Physical Anthropology 165:23–36. DOI:https://doi.org/10.1002/ajpa.23384

Rogers J, Raveendran M, Harris RA, Mailund T, Leppäla K, Athanasiadis G, Schierup MH, Cheng J, Munch K, Walker JA, Konkel MK, Jordan V, Steely BJ, Beckstrom TO, Bergey C, Burrell A, Schrempf D, Noll A, Kothe M, Kuhlwilm M, et al. 2014. The complete genome sequence of a Neanderthal from the Altai Mountains. Current Biology 24:R102635. DOI:https://doi.org/10.1016/j.cub.2014.05.067

Silk JB, Alberts SC, Wray GA. 2010. Strong and consistent social bonds enhance the longevity of female baboons. Current Biology 20:1359–1361. DOI:https://doi.org/10.1016/j.cub.2010.05.067

Silk JB, Roberts ER, Barrett BJ, Patterson SK, Strum SC. 2017. Female–male relationships influence the form of female–female relationships in olive baboons, Papio anubis. Animal Behaviour 131:89–98. DOI:https://doi.org/10.1010/j.anbehav.2017.07.015

Smuts BB. 1985. Sex and Friendship in Baboons. New York: Aldine Publishing Company.

Smuts B, Nicolson N. 1989. Reproduction in wild female baboons. American Journal of Primatology 19:229–246. DOI:https://doi.org/10.1002/ajp.1000190040

Snyder-Mackler N, Majoros WH, Yuan ML, Shaver AO, Gordon JB, Kopp GH, Schelbusch SA, Wall JD, Alberts SC, Mukherjee S, Zhou H, Tung J. 2016. Efficient Genome-Wide sequencing and Low-Coverage pedigree analysis from noninvasively collected samples. Genetics 203:699–714. DOI:https://doi.org/10.1034/genes.11817492

Städele V, Van Doren V, Pines M, Sweddell L, Vigilant L. 2015. Fine-scale genetic assessment of sex-specific dispersal patterns in a multilevel primate society. Journal of Human Evolution 78:103–113. DOI:https://doi.org/10.1016/j.jhevol.2014.10.019

Städele V, Pines M, Sweddell L, Vigilant L. 2016. The ties that bind: maternal kin bias in a multilevel primate society despite natal dispersal by both sexes. American Journal of Primatology 78:731–744. DOI:https://doi.org/10.1002/ajp.22537

Städele V, Roberts ER, Barrett BJ, Strum SC, Vigilant L, Silk JB. 2019. Male–female relationships in olive baboons (Papio anubis): Parenting or mating effort? Journal of Human Evolution 127:81–92. DOI:https://doi.org/10.1010/j.jhevol.2018.09.003

Strandburg-Peshkin A, Farine DR, Cousin ID, Crofoot MC. 2015. Shared decision-making drives collective movement in wild baboons. Science 348:1358–1361. DOI:https://doi.org/10.1126/science.aaa5099

Stringer C. 2016. The origin and evolution of Homo sapiens. Philosophical Transactions of the Royal Society B: Biological Sciences 371:20150237. DOI:https://doi.org/10.1098/rstb.2015.0237

Sweddell L. 2011. African Papionins: Diversity of Social Organization and Ecological Flexibility. In: Campbell C, Fuentes A, MacKinnon K, Bearder S, Stumpf R (Eds). Primates in Perspective. New York: Oxford University Press. 241–277.

Sweddell L 2013. Hamadryas Baboon (Papio hamadryas). In: Butynski T, Kingdon J (Eds). The Mammals of Africa. Primates. London: Bloomsbury Publishing. p. 221–224. DOI:https://doi.org/10.5040/9781472926920.0038

Tung J, Alberts SC, Wray GA. 2010. Evolutionary genetics in wild primates: combining genetic approaches with field studies of natural populations. Trends in Genetics 26:353–362. DOI:https://doi.org/10.1016/j.tig.2010.05.005

VandeBerg JL. 2003. Social bonds of female baboons: status and prospects. In: Campbell C, Fuentes A, MacKinnon K, Bearder S, Stumpf R (Eds). The Baboon in Primatology. London: Bloomsbury Media. p. i–xxiii. DOI:https://doi.org/10.1007/978-0-387-75991-3

Fischer et al. eLife 2019;8:e50989. DOI:https://doi.org/10.7554/eLife.50989

15 of 16
Vilgalys TP, Rogers J, Jolly CJ, Mukherjee S, Tung J. 2019. Evolution of DNA methylation in Papio baboons. *Molecular Biology and Evolution* **36**:527–540. DOI: https://doi.org/10.1093/molbev/msy227, PMID: 30521003

Vore DI, Washburn SL. 1961. The social life of baboons. *Scientific American* **204**:62–72. DOI: https://doi.org/10.1038/scientificamerican0661-62

Walker JA, Jordan VE, Steely CJ, Beckstrom TO, McDaniel CL, St. Romain CP, Bennett EC, Robichaux A, Clement BN, Konkel MK, Batzer MA. 2017. Papio baboon species indicative alu elements. *Genome Biology and Evolution* **9**:1788–1796. DOI: https://doi.org/10.1093/gbe/evx130

Wall JD, Schlebusch SA, Alberts SC, Cox LA, Snyder-Mackler N, Nevonen KA, Carbone L, Tung J. 2016. Genomewide ancestry and divergence patterns from low-coverage sequencing data reveal a complex history of admixture in wild baboons. *Molecular Ecology* **25**:3469–3483. DOI: https://doi.org/10.1111/mec.13864

Wasser SK, Norton GW, Kleindorfer S, Rhine RJ. 2004. Population trend alters the effects of maternal dominance rank on lifetime reproductive success in yellow baboons (Papio cynocephalus). *Behavioral Ecology and Sociobiology* **56**:338–345. DOI: https://doi.org/10.1007/s00265-004-0797-2

Zinner D, Groeneveld LF, Keller C, Roos C. 2009. Mitochondrial phylogeography of baboons (Papio spp.) – Indication for introgressive hybridization? *BMC Evolutionary Biology* **9**:83. DOI: https://doi.org/10.1186/1471-2148-9-83

Zinner D, Arnold ML, Roos C. 2011. The strange blood: natural hybridization in primates. *Evolutionary Anthropology: Issues, News, and Reviews* **20**:96–103. DOI: https://doi.org/10.1002/evan.20301

Zinner D, Wertheimer J, Liedigk R, Groeneveld LF, Roos C. 2013. Baboon phylogeny as inferred from complete mitochondrial genomes. *American Journal of Physical Anthropology* **150**:133–140. DOI: https://doi.org/10.1002/ajpa.22185, PMID: 23180628