Older age becomes common late in human evolution

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Increased longevity, expressed as number of individuals surviving to older adulthood, represents one of the ways the human life history pattern differs from other primates. We believe it is a critical demographic factor in the development of human culture. Here, we examine when changes in longevity occurred by assessing the ratio of older to younger adults in four hominid dental samples from successive time periods, and by determining the significance of differences in these ratios. Younger and older adult status is assessed by wear seriation of each sample. Whereas there is significant increased longevity between all groups, indicating a trend of increased adult survivorship over the course of human evolution, there is a dramatic increase in longevity in the modern humans of the Early Upper Paleolithic. We believe that this great increase contributed to population expansions and cultural innovations associated with modernity.

The human life history pattern differs from that observed in the great apes in its delayed maturation, slower growth, higher fertility, and increased longevity, which is associated with menopause in women (refs. 1–4, and see figure 1 in ref. 2 for mortality differences between recent hunter-gatherers and chimpanzees). These are evolutionary changes that have implications for the development of human culture. Longevity, in particular, may be necessary for the transgenerational accumulation and transfer of information that allows for complex kinship systems and other social networks that are uniquely human. It is also a focal point of the grandmother hypothesis, which posits that increased longevity is important in enhancing the inclusive fitness of grandmothers who, perhaps as early as the first Homo populations, invested in their reproductive-age daughters and their offspring (4–6).

Therefore, the details of how longevity increases over the course of human evolution are of great interest (7). Other aspects of the hominid life history pattern, in particular, rates of maturation and growth and development, have been the focus of intensive study in fossil hominins (see, for example, refs. 8–11 and references therein), but changes in longevity itself have not been empirically assessed. Longevity is usually discussed through its correlation with other variables, such as body size, encephalization, and growth and development patterns (12, 13). On the basis of these correlations, some have argued that increased longevity appears quite early in the evolution of Homo (5, 6). Observation of larger brain and body size implied prolonged maturation in Homo ergaster (11, 14, 15), leading some workers to suggest longer lifespan in this taxon. On the other hand, it has also been suggested that early hominins may not have lived long enough to become grandmothers, based on published ages from a few key sites (16). However, a large sample, amenable to meaningful statistical treatment of this issue, has not been analyzed. The actual pattern of change in adult survivorship critical to testing the reality of correlations between brain size and longevity in the human lineage, the predictions of the grandmother hypothesis, and any of the other questions surrounding the evolution of human longevity, has yet to be empirically established. Here, we address the basic issue of whether different hominid groups from successive time periods have different patterns of longevity, regardless of differences in brain size and maturation rates, and whether the fossil record supports the prediction of increased longevity in early Homo.

By longevity, we are not concerned with the maximum life span attainable by members of a species, but with the number of adults who live to be old. In our view, it is the number of individuals living to older adulthood that provides evidence of selection favoring the survivorship of older adults and is important for many of the evolutionary issues surrounding changes in longevity. To avoid the taphonomic and demographic problems involved with the inclusion of juveniles, we limited the present study to adults, examining changes in the ratio of older to younger adults (OY ratio) in the death distribution over time. Although this is not the OY ratio that would be expected in the living populations (17), it does reflect it, and its changes provide insight into the evolution of age structure in the human fossil record.

We tested the null hypothesis of no difference in the composition of older and younger adults among four hominid groups represented by 768 dental individuals, estimated conservatively so that no specimen could be counted more than once: later australopithecines (including specimens attributed to Australopithecus and Paranthropus), Early and Middle Pleistocene Homo, Neandertals from Europe and Western Asia, and post-Neandertal Early Upper Paleolithic Europeans (Table 1). The habilines were excluded from both the Homo and australopithecine samples because their taxonomic placement is uncertain. They may be more closely associated with australopithecines than with early Homo (18, 19), with whom they are penecontemporary, and the sample is too small to stand alone as a separate group. As discussed below, their inclusion in either category would not change results significantly. The four groups were chosen because they provide large samples amenable to dental wear seriation and categorical adult age assessment. Each broad group represents several different sites with different taphonomic histories, reducing the chance that the circumstances of one particular site will significantly bias the results. We chose specimens for whom we had reasonable confidence in both their categorical age assessment and the likelihood that they represented a unique individual. Significance of the differences in OY ratios between the groups was tested by using distributions generated by random resampling with replacement.

Dental age estimates, based on wear seriation, were used to place specimens into categories of older or younger adults. These estimates should not be thought of as absolute numbers, but as ranges within which the actual age probably lies, with resolution depending on condition and age of the specimen, knowledge of the population from which it comes, a number of assumptions about eruption schedule, and, to some extent, the constancy of wear rates between populations and tooth classes (20–24). Wear-based seriation was pioneered by Miles (25) and described and successfully used by many workers (26–30). By using this method, rates of wear are estimated by
in this illustration erupted at 3, 7, and 11 years of age for M1, M2, and M3, respectively, to use one chimpanzee model, the wear would represent less time (2.08 OY ratio with minimal M2 wear; SK 34 (11 plus 3), and STS 36 would remain in the older adult category (15 plus 5), indicating an age of tooth wear. Therefore, this indigenous Australian schedule may be appropriate for primitive cultures with heavy masticatory stresses, as one would expect in Paleolithic groups (20, 35, 36). However, we slightly modified the schedule reported by Brown et al. (34) with respect to M3 eruption, as discussed above (19, 30).

Although, in constituting our categories, we used the same eruption schedule for all fossil groups, we recognize that the maturation rate for australopithecines and early Homo is controversial (8–11, 26, 37). However, our use of OY ratios circumvents this debate because the categories are independent of actual ages as long as hominid dental development is tied to physiological development, as it is in other primates (31). This eruption indicates adulthood, and the relative wear indicators of older adulthood apply regardless of actual age (Fig. 1). Therefore, for consistency and comparability we determined age estimates according to the eruption schedule discussed above, but we emphasize that the categories reflect the degree of molar wear at the time of occlusal eruption of subsequent molars on immature specimens and these rates are then extrapolated to older individuals (Fig. 1). Because our sample is categorical (older or younger adult) and not composed of the age estimates themselves, we do not require that the ages have a high level of resolution. Although determination of the categories is not always unambiguous, the procedures we used minimize error.

Individuals with M3s in occlusal eruption were considered adult, and individuals double that age were considered older adult. As discussed below, although we estimated ages based on a single eruption schedule, these ages were used to allocate specimens to life history categories that are themselves independent of actual ages. Regardless of chronological age, M3 eruption marks the end of the juvenile period and is correlated with physiological maturation, including reproductive maturation, in primates (31, 32). However, it was convenient for purposes of comparison and necessary for objective and repeatable assessment of when individuals were double the age of M3 eruption, to estimate an age at death. In making these assessments, we accept the rationale for M3 eruption at age 15 (in contrast to a later age in recent humans), based on observations of minimal M2 wear at the time of M3 occlusal eruption in Neandertal and some recent humans. Older adults were defined as twice the age of reproductive maturation, the age at which one could theoretically first become a grandmother; i.e., 30 years if women had their first child at the age of 15.

These age estimates use one of the fastest eruption schedules observed in human populations (34), transformed from ginvival to occlusal eruption after Wolpoff (30). Our rationale for this estimate derives from evidence that eruption schedules are, in part, influenced by the rate of exfoliation of deciduous teeth (34), which in turn is related to deciduous tooth wear. Therefore, this indigenous Australian schedule may be appropriate for hunting/gathering groups with heavy masticatory stresses, as one would expect in Paleolithic groups (20, 35, 36).

Table 1. Sample summary: Sample sizes and OY ratio for each group

<table>
<thead>
<tr>
<th>Group</th>
<th>Old</th>
<th>Young</th>
<th>Total</th>
<th>OY</th>
</tr>
</thead>
<tbody>
<tr>
<td>Australopithecines</td>
<td>37</td>
<td>316</td>
<td>353</td>
<td>0.12</td>
</tr>
<tr>
<td>Early Homo</td>
<td>42</td>
<td>166</td>
<td>208</td>
<td>0.25</td>
</tr>
<tr>
<td>Neandertals</td>
<td>37</td>
<td>96</td>
<td>113</td>
<td>0.39</td>
</tr>
<tr>
<td>Early Upper Paleolithic</td>
<td>50</td>
<td>24</td>
<td>74</td>
<td>2.08</td>
</tr>
<tr>
<td>All groups combined</td>
<td>166</td>
<td>602</td>
<td>768</td>
<td>0.28</td>
</tr>
</tbody>
</table>

Our sample derives from our own age estimates and age estimates provided by M. H. Wolpoff and D. W. Frayer.

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relative wear determined through seriation and not the numerical ages attributed to them. They represent physiological age groups, and are therefore minimally affected by variation in maturation rates. This reason is why we chose a categorical approach.

Wear-based seriation was performed independently for different fossil groups, so the assumption of reasonably constant wear rates across populations was made only for samples with no subadults to seriate. Although there is evidence that molar wear rates across fossil populations are reasonably comparable (22, 30), this conclusion has not been established across earlier fossil species, especially those with specialized diets like the robust australopithecines. However, the dietary specializations and associated craniofacial architecture among australopithecines that could contribute to greater tooth wear are also associated with larger teeth and thicker enamel that reduce rates of wear, and there is no direct evidence that wear rates are faster in the more specialized australopithecines. In addition, although different molars (tooth classes) may have slightly different wear rates, Miles (21) suggests that this variation ultimately makes little difference to estimates of age.

The categorical treatment of ages circumvents much of the inaccuracy associated with dental age assessment. Our categories are unaffected by the most common inaccuracies associated with wear-based aging, methods that tend to overage modern humans over the age of 35 and underage very old individuals over the age of 50 (21). Our “older” category, age 30 and above, is an age at which age estimates are still quite accurate. Treating the data categorically also allowed us to maximize our data set by including categorical assessments (for example, as in the case of very old individuals, whose teeth are too worn for age assessment) as well as numerical ages as described above, depending on the condition of specimens and the source of the assessment.

We tested the null hypothesis of no difference in longevity between the fossil groups by comparing the number of older adults relative to that of younger adults in the samples of hominid dental remains. These OY ratios and sample sizes for each of the four groups are given in Table 1. OY ratios increase over time. The significance of the differences among OY ratios was assessed by establishing the probability of finding the observed OY ratio of one fossil group within distributions of OY ratios sampled from the earlier group. These distributions were generated through random resampling with replacement. The approach of data resampling provides a way to solve problems that lie outside the analytical boundaries of classical statistics (38), for example, by using ratios as the statistic of interest as in this study. Resampling addresses the problem of interdependence of data within the death distribution and the potential further dependence of the successive samples on earlier ones because of evolution.

Our null hypothesis, that there is no difference in OY ratios among the different hominid groups, can be stated as a question of probability: how likely is it to observe an OY ratio of a particular hominid group in an earlier group? We rejected the null hypothesis if a ratio the same or greater than the observed OY ratio was found in 2.26% of the generated ratio distribution, rejecting the null hypothesis of no difference.

The large increase in ratios associated with the Upper Paleolithic is most significant (Fig. 2C). The observed Upper Paleolithic ratio (2.08) lies well outside maximum possible OY ratio generated from the Neandertal sample. Because of the unexpected magnitude of this increase, we questioned whether the formulation of our age categories could have introduced bias that influenced the results. For this reason, we used an adulthood criterion of 15 years for the Early Upper Paleolithic as we did for the earlier samples. It is probable that Early Upper Paleolithic M3 eruption was closer to 16–18 years, as it is for some living hunter–gatherers, but the average age at first birth in these populations may have been earlier, as reported for Inuit and some Australian groups (3). Hence, although we may have erred by including some subadults in the Early Upper Paleolithic young adult sample, this approach is the most conservative one because it increases the size of the younger adult portion of the sample. Because the significant change is in the increased OY ratio, this approach would underestimate, rather than overestimate, this increase.

Our data do not directly address the causes of life history differences, the high representation of young adults in earlier samples, or the increased representation of older adults in the Upper Paleolithic. We expect that the data reflect variation in the causes of young adult mortality, among other factors; the systematic nature of the OY ratio changes and the many different site histories sampled make a purely taphonomic explanation unlikely. To further examine the possibility of taphonomic bias, we questioned whether mortuary practices, specifically an increased number of burials, are responsible for the magnitude of the relative increase of the number of older adults in the Upper Paleolithic. Table 2 shows the OY ratios based on burials and nonburials within the Upper Paleolithic sample. The Upper Paleolithic OY ratio based on nonburials (2.06) is slightly lower than the Upper Paleolithic OY ratio based on burials (2.13), but the difference is not significant. OY ratio distributions were generated from the Neandertal sample with sample size of 49 for the nonburials and 25 for the burials, and the probability of observing ratios of 2.06 and 2.13 (respectively) was assessed. Both cases rejected the null hypothesis (Fig. 3). Burials could contribute to the large increase in older individuals.
Two important conclusions emerge from this study: first, there is significant increased longevity between all groups, indicating a trend of increased survivorship of older adults through human evolution. Second, the increase is by far the greatest in the early modern humans of the Upper Paleolithic, when for the first time there are a larger number of older adults than younger adults in the death distribution. Whereas high levels of young adult mortality have been noted for Neandertals (39), the magnitude of the increase in OY ratios in the Upper Paleolithic is nevertheless surprising.

The larger OY ratio in Homo relative to australopithecines weakly supports the predictions of the grandmother hypothesis as applied to Homo erectus (6). The OY ratio doubles in the Early/Middle Pleistocene Homo sample, but is still very low. In contrast, the increased longevity in the Upper Paleolithic is dramatically larger, with OY ratios five times greater than that seen in Middle Paleolithic hominids. Because the OY ratio increases so significantly at this time, we suggest that theories involving the evolutionary value of senescence may be most applicable to the Middle/Upper Paleolithic transition.

Although longevity may be under direct selection as suggested in some kin selection models (40), or may be linked to other traits under direct selection, it is not clear that the increase in adult survivorship reported here necessarily has a genetic basis. Long individual lifespans occur in nonhuman primates (41), not just in humans, and therefore increase in the number of individuals living to older age may not reflect a major genetic change affecting maximum lifespan. However, whether the result of cultural factors, other forms of relaxed selection affecting the mortality of young adults, and/or biological change, the increase in adult survivorship would have considerable evolutionary impact. If related to senescence, it reflects an adaptation that must help compensate for the disabilities and diseases of older age, when gene expressions uncommon in younger adults became more frequent. We believe that this adaptation involves the increased importance of transgenerational relationships that may be critical to the development and survival of social groups with large amounts of complex information to transmit. Increased adult survivorship strengthens those relationships and information transmission by extending the time over which people can learn from older individuals and by the increase in the number of older people, which promotes the acquisition and transmission of specialized knowledge such as that reflected in the Upper Paleolithic.

Table 2. OY ratios for burials and nonburials in the Early Upper Paleolithic

<table>
<thead>
<tr>
<th></th>
<th>Old</th>
<th>Young</th>
<th>Total</th>
<th>OY</th>
</tr>
</thead>
<tbody>
<tr>
<td>Nonburials</td>
<td>33</td>
<td>16</td>
<td>49</td>
<td>2.06</td>
</tr>
<tr>
<td>Burials</td>
<td>17</td>
<td>8</td>
<td>25</td>
<td>2.13</td>
</tr>
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</table>

Fig. 2. Probability of finding observed OY ratios for three hominid groups in distributions of OY ratios generated from hominids from an earlier time period by resampling. Each OY ratio is created by randomly drawing an adult sample of the same size as the test (later) group from the earlier one and calculating the OY ratio in it (sample sizes are shown in Table 1). This procedure was performed 10,000 times in each case, and the generated samples describe the probability of observing the OY ratio from the test (later) sample in the earlier one. For example, each one of the 10,000 OY ratios in A is a random sample of 208 australopithecine adults, because the actual Homo sample size is 208. (A) The observed early Homo OY ratio (vertical bar at 0.25) compared with ratios generated from the australopithecine sample. The observed Homo ratio was never reproduced in 10,000 OY ratios in draws of 208 generated from the australopithecine sample. (B) Observed Neandertal OY ratio (0.39) compared with ratios generated from the Early and Middle Pleistocene Homo sample in 10,000 draws of 133. OY ratios of 0.39 and larger occurred in 2.26% of the generated distribution. (C) The observed Early Upper Paleolithic OY ratio (2.08) compared with ratios generated from the Neandertal sample in 10,000 draws of 74. The observed ratio is far outside this distribution, representing a major increase in the number of adults over the age of 30 in the death distribution.
The potential demographic consequences of increased adult survivorship are significant. Our results are compatible with archeological and genetic evidence suggesting population expansions in the Upper Paleolithic that may be a consequence of population growth (42, 43). There is reason to think that increased longevity would have a direct effect on population growth. Not only does increased survivorship imply greater lifetime fertility for individuals, the investment of older individuals in their children’s families may provide a selective advantage promoting further population increase. The large OY ratio we observe may therefore be a significant factor in the evolution of modernity not only through its importance for transgenerational information transfer but also because of its demographic impact. Recent models (43) suggest demographic factors are responsible for the cultural innovations associated with modernity. Population expansion may have provided social pressures that led to extensive trade networks, increased mobility, and more complex systems of cooperation and competition between groups, resulting in increased personal ornamentation and other material expressions of individual and group identity.

In summary, while the fossil record by its very nature reflects sampling biases and there is a great need to develop and refine techniques for aging older adults to address the evolution of senescence and to understand the changes presented here in more detail, these results suggest a major increase in adult survivorship in the Upper Paleolithic. We suggest that this increase in longevity addresses the meaning of modernity itself. Modernity is a complex concept, incorporating both biological and cultural variables, that has proven difficult, if not impossible, to define (44). However, if there is a single fundamental factor related to biology that underlies the behavioral innovations of modernity, this increase in adult survivorship may be it. We therefore think significant longevity came late in human evolution and was a fundamental demographic component tied to the population expansions and related behavioral innovations associated with modern humans.

We thank David W. Frayer (University of Kansas, Lawrence) and Milford H. Wolpoff (University of Michigan, Ann Arbor) for additional age estimates; Mark Flinn, Thomas Roček, and three anonymous reviewers for advice and comments that significantly improved this manuscript; and the Anthropos Institute, Moravské Muzeum (Brno, Czech Republic), Anthropologické Oddělení, Národní Museum (Prague), Hrvatski Prirodoslovni Muzej (Zagreb, Croatia), Institute for Paleontology and Quaternary Geology, Croatian Academy of Sciences and Arts (Zagreb), Natural History Museum (London), Paleoanthropology Laboratory, Université de Bordeaux (France), Institute for Vertebrate Paleontology and Paleoanthropology (Beijing), Transvaal Museum (Pretoria, South Africa), and the Department of Anatomy, University of the Witwatersrand (Johannesburg) for permission to study the fossils in their care.

News and Views

Taxonomy and longevity: a reply to Minichillo (2005)

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We recently used the fossil record to examine theoretical predictions that had been made about demographic age-structure changes in human evolution (Caspari and Lee, 2004). Although it had been suggested that longevity increased in early Homo populations based on brain size increases and other correlates of longevity, this question, and others involving changes in age structure, had not previously been addressed using the fossil record.

Our paper on adult survivorship was empirical; we were careful to make our assumptions explicit, and we were cautious in the interpretation of our results. These interpretations could be easily biased by preconceived ideas about the relationships between the groups studied, and for that reason we avoided taxonomic or phylogenetic assumptions, focusing instead on variation over time. In his commentary, Minichillo (2005) makes suggestions that require phylogenetic assumptions that our study neither demanded nor addressed.

We tested a null hypothesis of no difference in adult survivorship by examining the ratio of older to younger adults (OY ratio) in four hominid groups, using the largest samples yet to be analyzed: later australopiths, early/middle Pleistocene Homo, Neandertals, and Upper Paleolithic Europeans (Caspari and Lee, 2004). Age categories were based on tooth wear. There were differences, and we assessed their significance using random resampling with replacement. We presented this analysis as a baseline study, one that was intended to provide general characteristics of age structure for several hominid groups. One of our interests was to test whether the fossil record supported a prediction arising from the work of O’Connell et al. (1999) regarding the “grandmother hypothesis,” as applied to early Homo. These authors suggested that a grandmother effect may have been related to the success of early Homo erectus populations, whose adaptation to environmental changes in the early
Pleistocene involved expanded dietary breadth, including a significant number of tubers—a way of life in which grandmothers could have played an important economic and child-rearing role. We expected a consequence of the grandmother hypothesis to be a concomitant increase in adult survivorship in early Homo populations (compared to australopiths), reflecting selection for larger numbers of older adults that promoted intergenerational transfer effects (Lee, 2003). Our question, then, was simple: Was there a difference in OY ratios among the four samples, and in particular, was there an increase in early Homo compared to australopiths? We concluded that there was a significant difference among all four groups (Fig. 1).

We discussed two implications of our results (Caspari and Lee, 2004). First, we found that each group had a significantly higher OY ratio than temporally earlier groups. Therefore, our results could support (i.e., did not refute) the grandmother hypothesis as applied to early Homo because the early Homo OY ratio was significantly higher than that of the australopiths. We do not agree with Minichillo’s (2005) contention that our results provide no support for O’Connell et al.’s (1999) position. However, because these OY ratios are still very low (OY = 0.25), we considered it weak support. The “weak support” Minichillo (2005: 643) refers to in his introduction—“[Caspari and Lee] further stated that these findings weakly support the Grandmother Effect, as articulated by Hawkes and colleagues (Hawkes et al., 1998; Hawkes, 2003), as a primary difference between modern and non-modern peoples” (emphasis added)—actually applied to O’Connell et al.’s position. We wrote: “these findings weakly support the predictions of the grandmother hypothesis as applied to Homo erectus” (Caspari and Lee, 2004: 10898, emphasis added). We think our results could provide strong support for the grandmother effect (or an “older person effect”) as a primary difference between modern and non-modern people.

The second and more important implication of our paper involved the most recent sample, Upper Paleolithic Europeans, whose OY ratio was five times higher than that of Neandertals. We proposed that the behavioral modernity associated with the Upper Paleolithic may be linked to the increase in the number of adults who survived to older ages, and suggested ways that the Upper Paleolithic may have been a consequence of changes in demographic age structure. While we did not rule out the possibility that the change in age structure was an attribute of a modern human lineage, we also didn’t assume it. In fact, we did not test hypotheses about the cause of changes in age structure (i.e., whether they were a consequence of lineage differences or not), and we made no taxonomic inferences.

Minichillo’s (2005: 643) critique is:

Unfortunately, the data Caspari and Lee had available failed to address the questions they would have liked them to. If the question is “Were age structures different between Upper Paleolithic Homo sapiens and Middle Paleolithic Homo neanderthalensis?” then the answer is decidedly “yes,” and stunningly so. But this is neither the question that Caspari and Lee asked nor the result that they emphasized.

To the contrary, this is precisely the question we asked of all the Homo groups, and the results we emphasized, but without the taxonomic emphasis that Minichillo gives.

Minichillo’s comment implies that our paper focused on the modern human origins issue. If so,
it perhaps reflects a misunderstanding of our use of “late” in the title: “Older age becomes common late in human evolution.” “Late,” in the context of our paper, referred to the difference between early and late Pleistocene; we reported that age structure changes were more dramatic in the modern humans of the Upper Paleolithic than in our earlier samples, but these are not the earliest modern humans. We did not address the question of whether age structure changed with the emergence of modern humans or occurred later than that, i.e., in the Upper Paleolithic. Minichillo is interested in whether the increase in longevity is an attribute of modern humans as a taxon, but this is not the question we addressed.

Minichillo assumes a unique African ancestry for modern Europeans. Given that, it is possible to address the question he thinks we should have asked: Is the increase in OY ratios found in Upper Paleolithic Europeans an attribute of a modern human taxon, or did it arise later, perhaps associated with cultural change within modern humans? We recently tested this idea by examining whether the high OY ratio of Upper Paleolithic Europeans is present in Middle Paleolithic moderns (Caspari and Lee, 2005, in press). We conducted two tests. First, since the two groups are both associated with the Middle Paleolithic, yet are often considered to be different species (lineages), a comparison of their OY ratios is informative. Second, since the western Asian moderns are usually considered to be closely related to earlier and contemporary modern humans in Africa, and have been viewed as ancestral to Upper Paleolithic Europeans by workers on all sides of the modern human origins controversy (e.g., Bar-Yosef and Vandermeersch, 1993; Klein, 1999; Wolpoff et al., 2001), we compared western Asian moderns to the later Upper Paleolithic modern humans from Europe.

Our Middle Paleolithic western Asian sample is small (n = 32) compared to the samples in our original study. Using resampling methods to assess the impact of small sample size, we found there was no significant difference in adult survivorship between the roughly contemporary Middle Paleolithic Neandertals (n = 18; OY = 1.0) and modern humans (n = 14; OY = 0.75) in western Asia (Caspari and Lee, 2005, in press). The lack of age structure variation implies that phylogeny is not a factor in the large OY ratio differences between the Neandertals and modern humans in Europe. We also compared the modern humans of the Middle and Upper Paleolithic. The Upper Paleolithic Europeans had an OY ratio (2.10) that was significantly larger than that of the Middle Paleolithic moderns (0.75). This result indicates that the large increase in the European moderns is not directly related to phylogeny (Caspari and Lee, 2005, in press).

In conclusion, the causes of adult mortality and survivorship across the span of human evolution are complex—caused by a variety of ecological, behavioral, and possibly phylogenetic factors. However, the consequences of a reduction in young-adult mortality may be linked to demographic changes that could have influenced the behavioral expressions of the Upper Paleolithic, and the intergenerational transfers that make older individuals in social species evolutionarily important, despite their decreasing fertility (Lee, 2003; Caspari and Lee, 2004; Rosenberg, 2004). Our original work made no assumptions about potential causes, but used the fossil record to test limited questions about changes in adult survivorship that may provide insight into life history evolution. This kind of empirical approach uses the prehistoric record to test hypotheses generated from evolutionary theory, and when keeping assumptions to a minimum, provides a basis for the generation of further hypotheses about the causes and effects of life history changes over the course of human evolution.

\[1\] We refer to the non-Neandertal Middle Paleolithic hominids as “moderns,” although we recognize that their categorization as “modern humans” or “anatomically modern humans” is controversial. Definitions of modernity are complex, and “modern humans” may not represent a monophyletic group. Nevertheless, it is conventional to use “modern” to refer to the non-Neandertal Middle Paleolithic specimens from western Asia, and we employ it here.
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References