On the apparent rarity of epithelial cancers in captive chimpanzees

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Malignant neoplasms arising from epithelial cells are called carcinomas. Such cancers are diagnosed in about one in three humans in ‘developed’ countries, with the most common sites affected being lung, breast, prostate, colon, ovary and pancreas. By contrast, carcinomas are said to be rare in captive chimpanzees, which share more than 99% protein sequence homology with humans (and possibly in other related ‘great apes’—bonobos, gorillas and orangutans). Simple ascertainment bias is an unlikely explanation, as these nonhuman hominids are recipients of excellent veterinary care in research facilities and zoos, and are typically subjected to necropsies when they die. In keeping with this notion, benign tumours and cancers that are less common in humans are well documented in this population. In this brief overview, we discuss other possible explanations for the reported rarity of carcinomas in our closest evolutionary cousins, including inadequacy of numbers surveyed, differences in life expectancy, diet, genetic susceptibility, immune responses or their microbiomes, and other potential environmental factors. We conclude that while relative carcinoma risk is a likely difference between humans and chimpanzees (and possibly other ‘great apes’), a more systematic survey of available data is required for validation of this claim.

1. What are carcinomas?

During embryogenesis, the three germ layers (endoderm, ectoderm, mesoderm) differentiate into epithelial and nonepithelial cells, which eventually form differentiated tissues and organs [1]. Epithelial cells arise from stem cells and often line body surfaces that interact directly with the environment [2]. The type of epithelium reflects location and function. For example, the squamous epithelium of the skin epidermis, oropharynx and uterine cervix is stratified, impervious and protective against shear forces. By contrast, the epithelial lining of the lumen of the gastrointestinal tract and bronchial airways, and the ducts of pancreas, breast and prostate are columnar or cuboidal (arranged in a single column), with basal nuclei and abundant cytoplasm, and can have secretory or absorptive functions. Epithelial cells are typically attached to underlying connective tissue by a basement membrane, and the underlying stroma includes blood vessels, lymphatic vessels, haematopoietic cells, stromal fibroblasts, extracellular matrix, neuronal structures, smooth muscle and adipose tissue [1].

Neoplasia is the term given to the new growth of cells proliferating without regard to stop signals, with attendant new blood vessels, forming a tumour mass (neoplasm) [3]. If the growing neoplasm within an epithelial layer stays restricted by a basement membrane and does not invade that barrier, it is designated as a benign tumour, which grows locally and may have physical effects, or occasionally secrete bioactive molecules. Benign neoplasms arising from columnar epithelia that have not breached the basement membrane are termed adenomas (or adenomatous polyps, if they protrude into the lumen of a hollow organ).

However, once tumour cells undergo multiple genetic changes [4,5] and breach and infiltrate through the basement membrane, the neoplasm becomes
a markedly increased human lifespan in many countries [9]. Accompanying this demographic transition has been a major increase in the lifetime risk of developing the types of carcinomas mentioned above [7,8]. While rare familial genetic variants contribute to a small fraction of these tumours in humans, the great majority are initiated by sporadic and spontaneous genetic mutations [3,10]. There is lifetime risk of about approximately 40% of males and approximately 30% of females being diagnosed with one of these tumours in so-called ‘developed’ countries, where these cancers now constitute a major proportion of overall causes of death [7,8].

3. ‘Great apes’, our closest evolutionary cousins
Abundant data now indicate that we humans are very closely related to other hominids including chimpanzees, bonobos, gorillas and orangutans (the so-called ‘great apes’, hereafter called nonhuman hominids, or NHHs) [11–13]. Indeed, humans are closer genetically to chimpanzees and bonobos than they are to gorillas and orangutans. And humans are closer genetically to the chimpanzee/bonobo clade than mice and rats are to each other [12,13]. Thus, it was reasonable to expect that chimpanzees would be good models for understanding human disease [14]. Prior to the recognition of their extreme cognitive similarities to humans and the resulting ethical concerns [15–18] large numbers of chimpanzees were thus placed in captivity, to be used for modelling of human diseases. While recent recommendations have markedly curtailed and almost eliminated the use of chimpanzees in biomedical research [17], there was already a considerable body of published and unpublished information regarding their disease profiles, particularly from facilities in the USA. Surprisingly, surveys of this existing information suggest that several common human diseases may be partially or completely unique to our species, and that the captive chimpanzee population may suffer from different profiles of pathology [19–22].

4. Extant literature suggests that carcinomas are uncommon in ‘great apes’
Among these apparent differences in disease incidence, one that has been emphasized in multiple reports is the rarity of occurrence of common human carcinomas in captive chimpanzees. Earlier surveys reporting on disease profiles and causes of death in captive chimpanzees from US facilities noted the rarity of these cancers [23–27]. A more recent thorough analysis [25] listed all neoplasms documented at two long-standing major US facilities from their inception many decades ago, through mid-2008. While the denominator (the total number of chimpanzees and life years at risk) was not made clear, there were only nine spontaneously arising epithelial malignancies reported over these many decades of observations of colonies that measured in the many hundreds. Moreover, even these cancers did not arise in the usual sites observed in humans, and were: one kidney carcinoma, one malignant carcinoma of the uterus, one basal-squamous carcinoma, one thyroid carcinoma, one adenocarcinoma of the parotid salivary gland, one nasopharyngeal carcinoma and three hepatocellular carcinomas (HCCs). There were a few additional cases of HCCs that were associated with experimental chronic hepatitis infection. No carcinomas of colon, breast, lung, prostate, stomach,
pancreas or ovary were reported in this cohort [25]. Another recent review [28] indicates that the incidence of tumours in non-human primates does increase with age but that malignancies of the haematopoietic system are most frequent.

5. Simple ascertainment bias appears an unlikely explanation

In addition to ethical considerations, ‘great apes’ are charismatic animals, and also represent a very expensive and valuable long-term investment. Thus captive NHHs in research facilities and zoos are typically the recipients of excellent veterinary care and are often subjected to careful necropsies when they die. As a consequence, benign tumours and other neoplasms that are relatively uncommon in humans are routinely reported in this population (the above-mentioned review [25] documented 89 benign tumours). Overall, it thus seems unlikely that the apparent rarity of the common human carcinomas in NHHs is simply a matter of ascertainment bias. Notably, we are not just referring to microscopic disease, which is nowadays picked up by early detection methods in humans. Prior to the era of modern medicine, untreated carcinomas in humans used to manifest during an individual’s lifetime with grossly evident or symptomatic primary or metastatic tumours. One would expect the same to happen in NHHs, who are not subjected to surveillance for microscopic disease. On the other hand, examination at post-mortem is of course biased towards internal organs.

6. Inadequacy of numbers surveyed

All told, there are probably only a few thousand NHHs that have been carefully studied in captivity over the past century. Thus, no reliable comments can be made about diseases that have the relative frequency of less than 1 per 100 in the human population. However, many of human carcinomas (other than a few examples such as red meat consumption), detection of which must also be taken into consideration. Primate facilities have typically used commercially prepared blended foods (‘extruded pellets’ or ‘biscuits’, composed of many by-products of human food production), supplemented by fresh produce. However, a systematic survey of the dietary composition experienced by captive NHHs is lacking. Moreover, dietary requirements and gastrointestinal anatomy and physiology for each NHH are markedly different, so comparisons must be done carefully. Also, it is likely that dietary regimes at various facilities have been a moving target over time. Regardless, it seems unlikely that the dietary composition of captive NHHs is markedly different from that experienced by humans. With regard to one particular risk factor for human carcinomas, our recent work suggests that it is specific for human carcinomas. With regard to one particular risk factor for human carcinomas, our recent work suggests that it is human-specific. In this instance, the nonhuman sialic acid Neu5Gc is metabolically incorporated from red meats (lamb, pork and beef) into human epithelia and endothelia, and therein interacts with circulating anti-Neu5Gc antibodies [39]. Our evidence suggests that the resulting chronic inflammation may contribute to cancer progression [39,40]. Even if the diet of NHHs happens to contain lots of Neu5Gc, this risk factor would not apply to the great apes as they all already express Neu5Gc naturally [41]. Regardless, the impact of red meat consumption on human carcinoma risk is modest [37,38], and this factor cannot by itself account for the overall difference.

7. Differences in life expectancy and population demographics

As mentioned earlier, most human carcinomas occur later in life. Thus, given the fact that the NHHs appear to age more rapidly and die earlier [32], one possible explanation is that they simply do not live long enough to experience the same carcinoma risk that humans do. However, with progressive improvements in veterinary care, these animals can now live into the fourth or even early fifth decade of life, and a few survive to the age of 60 [32]. Thus, one might have expected to see a few cases of adenocarcinomas of the colon, breast, ovary or pancreas, which do occur in humans before the age of 50. But any conclusions must remain speculative until there is a systematic survey of the entire population demographics of captive chimpanzees and other ‘great apes’ from all facilities that have routinely performed complete necropsies. Of course, this question is impractical to address for NHHs that live in the wild, because they die earlier, and also because necropsies are hardly ever possible in such cases. Necropsies performed on African and Asian sanctuary NHHs may be more detailed, but their accuracy would depend on the time of recovery of the bodies after death, as post-mortem autolysis would be rapid under tropical conditions.

8. Dietary factors

Dietary factors such as soluble fibre protection from colon cancer [33] and a red meat consumption risk for multiple carcinomas [34–40] must also be taken into consideration. Primate facilities have typically used commercially prepared blended foods (‘extruded pellets’ or ‘biscuits’, composed of many by-products of human food production), supplemented by fresh produce. However, a systematic survey of the dietary composition experienced by captive NHHs is lacking. Moreover, dietary requirements and gastrointestinal anatomy and physiology for each NHH are markedly different, so comparisons must be done carefully. Also, it is likely that dietary regimes at various facilities have been a moving target over time. Regardless, it seems unlikely that the dietary composition of captive NHHs is markedly different from that experienced by humans. With regard to one particular risk factor for human carcinomas, our recent work suggests that it is human-specific. In this instance, the nonhuman sialic acid Neu5Gc is metabolically incorporated from red meats (lamb, pork and beef) into human epithelia and endothelia, and therein interacts with circulating anti-Neu5Gc antibodies [39]. Our evidence suggests that the resulting chronic inflammation may contribute to cancer progression [39,40]. Even if the diet of NHHs happens to contain lots of Neu5Gc, this risk factor would not apply to the great apes as they all already express Neu5Gc naturally [41]. Regardless, the impact of red meat consumption on human carcinoma risk is modest [37,38], and this factor cannot by itself account for the overall difference.

9. Lifestyle and environmental factors

While the common squamous cell carcinomas in humans are typically associated with definitive environmental risk factors (smoking for lung cancer, ultraviolet radiation for skin cancer, papilloma virus infection for cervical cancer, etc.), the adenocarcinomas associated with internal organs do not have such clear-cut associations with environmental factors (other than a few examples such as red meat consumption). Captive chimpanzees likely also experience somewhat similar exposure to potential carcinogens in the air, water and food. With regard to breast cancer risk, it is worth noting that systematic vasectomy of males has long been used to curtail fertility in almost all current chimpanzee research and retirement facilities [17,42]. Thus, there are many living captive adult female chimpanzees that have never experienced pregnancy, live births or lactation. While these lifestyle changes constitute well-known risk factors for breast cancer in humans [43], there have been no reported cases of breast...
cancer in any chimpanzee to date. Again what is needed is a systematic survey of the population at potential risk, taking into account the number of years that have passed since the vasectomy policies were instituted.

10. Potential differences in immunosurveillance and/or immunostimulation

It is now clear that surveillance by cells of the immune system serves to eliminate many early tumours [44]. On the other hand, many of the same immune responses can facilitate the growth of established tumours by virtue of contributing to local chronic inflammation [45–47]. While this ‘double-edged sword’ role of the immune system in cancer is well recognized, it has only recently been appreciated how narrow the quantitative gap between the two mechanisms might be [48,49]. Siglecs are sialic-acid binding immunoglobulin-like lectins expressed on immune cells, playing multiple roles in human disease, including cancer [50]. In this regard, we have noted that the relative lack of Siglecs on human immune cells compared with those from the chimpanzee might contribute to increased immunoreactivity of some cell types [51,52]. While purely speculative, it is possible that this difference in inflammatory potential could also contribute to the increased risk of tumour progression in humans. More extensive surveys of the differences in responses of immune cells in whole blood samples suggest that the situation may be more complex [53].

11. Genetic susceptibility

As mentioned in §7, the rare inherited genetic variants that increase human cancer risk cannot account for the apparent difference in rates of sporadic adenocarcinomas. Attempts have therefore been made to look for genetic features unique to humans as a species that could contribute to increased risk. A comparative analysis of a defined set of 333 orthologous cancer genes in humans and chimpanzees showed a high degree of conservation [54]. However, a detailed analysis detected small differences in certain tumour suppressor genes, which might potentially influence the differences in cancer susceptibility [54]. Given the important role of telomere caps at chromosomal ends in modulating cell survival, ageing and cancer [55,56], it was reasonable to look for a difference in humans. Paradoxical to the finding of extended lifespan in humans [32,57], telomere lengths were found to be greater in nonhuman primates than in humans, and telomere shortening rates were not apparently different between humans and chimpanzees [58,59]. Given the importance of genome damage and DNA repair in cancer, it is also interesting that one study indicated that humans and chimpanzees differ in their cellular response to DNA damage and noncoding sequence elements of DNA repair-associated genes, with evidence for accelerated evolution in some promoter regions and introns [60]. Yet another theory considers the finding that expression of genes involved in programmed cell death of brain neurons is different between humans and chimpanzees, and predicts a reduced level of neuronal apoptosis in humans [61]. This pattern of expression is evidently maintained in other human organs, suggesting that cellular apoptosis may be generally reduced in humans relative to chimpanzees. Assuming that a decreased rate of neuronal apoptosis may have been important for increased cognitive ability in humans, this evolutionary process could have coincidently resulted in an increased risk of cancer and other diseases associated with reduced apoptotic functions [61]. A related mystery arises from the recent report indicating that the lifetime risk of cancers of many different human tissue types is strongly correlated with the total number of divisions of the normal self-renewing cells maintaining that tissue’s homeostasis, i.e. the suggestion that the majority of cancers result from ‘bad luck’—random mutations arising during DNA replication in normal, noncancerous stem cells [10]. If this line of reasoning is correct, it suggests that there may be something very different about rates of stem renewal and/or apoptosis in humans versus other hominids. Considering another possibility, comparative epigenomic studies have observed differing patterns of DNA methylation in the brain between humans and chimpanzees [62,63]. While interpretation was cautioned because DNA methylation changes also vary with age, it is possible that epigenomic changes required for modulating brain gene expression during human evolution could have secondarily affected the impact of the non-neural epigenome in modulating cancer progression. Overall, while these or other potential genetic differences may contribute to the apparent discrepancy in carcinoma risk, there is no single leading candidate mechanism.

12. Infectious agents

It is well known that chronic infections can contribute to the incidence and progression of various cancers [45–47,64]. In this regard, a classic example is the high frequency of HCC in humans who have suffered long-term chronic infection with the hepatitis B or C viruses [65]. While the literature is limited, the impression is that HCCs following long-term hepatitis infections in chimpanzees are rather rare [25,66]. What are needed are systematic and complete data on how many chimpanzees in total have been chronically infected with HBV or HCV, their years at risk following infection, and the number of cases of HCCs that have since occurred in this population. Another striking difference between humans and NHHs is the lack of long-term endemic infectious retroviral infections in human populations [67–69]—setting aside the recent introduction of HIV and HTLV viruses into the human population from NHH sources [67,70,71]. As chronic retroviral infections are associated with increased risk of cancer, in general, this fact also appears counterintuitive.

13. Impact of the microbiome

It is now well established that the complex communities of microbes that normally live in different sites of the body (the microbiome) can have a very strong influence on local inflammation, food metabolism, as well as on the processing of bioactive molecules such as carcinogens. It also seems likely that components of the microbiota can promote chronic inflammation and tumour development via complex interactions with the innate and adaptive immune systems [72]. Given that many epithelial surfaces are in contact with the exterior world, it is not surprising that these locations feature the most extensive microbiomes. Assuming differences between human and NHH microbiomes [73–75], it is reasonable to speculate that this factor contributes to the relative risk of carcinoma development. In this regard, another consideration is
the increasing use of antibiotics in humans, which could cause significant effects on the composition of the microbiome, and secondary effects on various biological processes [76,77]. As part of any survey, it would be interesting to compare the pattern and frequency of antibiotic usage between humans and captive NHHs.

14. Conclusion and perspectives
Considering all available evidence, it is not possible to definitively conclude that humans as a species are at higher risk of cancer biology and therapeutics. To paraphrase Carl Sagan, absence of definitive evidence is not definitive evidence of absence. While the existing data make it likely that there is a major difference, a more systematic survey of all facilities that have cared for chimpanzees and other NHHs and carried out good necropsies is needed. While this had been the initial hope [15,16,18], it now seems unlikely that we will get much in-depth information from chimpanzee retirement facilities, which still await funding, and other resources to perform complete and thorough necropsies for comparative analyses. Collaboration with veterinarians working with in situ NHHs (free ranging and sanctuaries) could also yield some information about this comparison. However, funding and resources to achieve this goal are again limiting. The situation is particularly unfortunate: these naturally ageing populations are valuable and ephemeral sources for important demographic comparisons with humans, adding key insights into the genesis of some of the most common diseases of humans, and those of our closest evolutionary relatives.

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