

Pearls

Fungi and the Rise of Mammals

Arturo Casadevall*

Departments of Microbiology and Immunology and Medicine, Division of Infectious Diseases, Albert Einstein College of Medicine, Bronx, New York, United States of America

The Mammalian Lifestyle Is Energetically Costly

Here are two indisputable facts: we are living in the age of mammals [1], and immunologically intact mammals are highly resistant to fungal diseases, such that most human systemic fungal are considered “opportunistic” [2]. Could these two facts be connected? The mammalian lifestyle is characterized by endothermy, homeothermy, and care for the young, including nourishment via lactation, all of which are energetically costly activities. In contrast, reptiles, which are ectotherms, require about one-tenth of the daily mammalian energy needs [3], and reptilian development is faster and requires less parental involvement. Given this energy handicap, how did mammals replace reptiles as the dominant land animals? This essay further develops the hypothesis originally proposed seven years ago that fungi contributed to the emergence of mammals by creating a fungal filter at the end of the Cretaceous that selected for the mammalian lifestyle and against reptiles [4].

Mammals Are Naturally Resistant to Fungal Diseases

Mammals are highly resistant to systemic fungal diseases. Whereas dermatophyte-associated diseases are common, these are seldom life threatening. For humans most fungal diseases were described in the 20th century and are associated with changes in the host such as iatrogenic immunosuppression, antibiotic-mediated disruption of the microflora, or other immune-impairing conditions as HIV infection, hematologic malignancies, and rheumatologic conditions. Unlike viral and bacterial diseases human mycoses are seldom contagious.

Endothermy and homeothermy are thought to contribute to mammalian resistance to mycosis by creating a thermal exclusionary zone that inhibits most fungal species [5]. The remarkable resistance of mammals to mycotic diseases is probably a combination of a vertebrate immune system, with both innate and adaptive arms, and elevated body temperatures. Experimental evidence for the synergy of temperature and immunity is apparent from studies of cryptococcal infection in rabbits, which have core temperatures of 40–41°C [6]. Rabbits are naturally resistant to systemic *C. neoformans* infection. However, rabbits can be infected with *C. neoformans* when inoculated in the skin or cornea, which are cooler, but the fungus does not disseminate. However, when rabbits are immunosuppressed with corticosteroids, *C. neoformans* infection is rapidly fatal [7].

Primitive mammals like the platypus, with core temperatures near 32°C, are susceptible to *Mucor amphibiorum*, a fungus with a maximal thermal tolerance of 36°C that would make it avirulent for higher mammals [8]. The resistance of mammals to fungal diseases is in sharp contrast to the vulnerability of other vertebrates, such as amphibians, a group that is currently under severe pressure from a chytrid [9]. Like mammals, amphibians have adaptive immunity, but unlike mammals, they are ectotherms and lack a thermal environment that is exclusionary to fungi. Hence, their vulnerability to fungal diseases echoes the

experimental findings in rabbits whereby high resistance is conferred by a combination of high temperature and vertebrate-level immunity [6]. Amphibians can be cured of chytridomycosis if placed at 37°C [10]. Another example of the protection provided by the combination of vertebrate-level immunity and endothermy comes from bats. In the summer bats manifest high activity and mammalian temperatures, but during winter hibernation their core temperatures drop as they hibernate and become vulnerable to infection with *Geomyces destructans*, a fungus that is decimating several North American bat species [11]. Infected bats woken from hibernation made full recovery when provided with supportive care, as higher body temperature inhibited fungal growth [12]. It is noteworthy that birds, which are also endotherms, are susceptible to *Aspergillus fumigatus* [13], a thermotolerant fungus that can survive up to 55°C [14].

A computation of the optimal temperature that would provide maximal protection against fungi given the caloric needs needed to maintain elevated temperatures yielded a value of 36.7°C, which is very close to mammalian temperatures [13]. This raises the possibility that mammalian resistance to fungi through the combination of vertebrate-level immunity and endothermy could have been the result of selection by pathogenic fungi.

A Fungal Filter at the Cretaceous-Tertiary (K-T) Boundary

Mammals replaced reptiles as the dominant land forms after the catastrophe that marked the end of the Cretaceous and the beginning of the Tertiary, an event known as the K-T boundary. The currently favored hypothesis for the demise of dinosaurs and end of the age of reptiles is a bolide impact approximately 65 million year ago with the possibility that other events, such as increased volcanism, contributed to disrupting the cretaceous ecosystem [15]. That ecological calamity was accompanied by massive deforestation [16], an event followed by a fungal bloom [17], as the earth became a massive compost. Although one cannot know which spores were present at the time, the likelihood that pathogenic fungi existed at the K-T boundary is enhanced by

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* E-mail: arturo.casadevall@einstein.yu.edu

the finding that the potential for pathogenicity probably arose independently several times in evolution [18].

There is now increasing evidence that large dinosaurs were warm bodied [19], as a result of their size, which would have entailed considerable heat generation dependent on food metabolism and their metabolic activities. Large animals at the top of the food chain, such as dinosaurs, are highly vulnerable to ecosystem disruption. The altered ecosystem would have implied disruption of food sources and changed climate, which is thought to have included a significant cooling of the earth [20], by dust clouds and fires. Such stresses would be expected to weaken any survivors of the bolide blast with consequent immunological impairment and could have made survivors, and their eggs, susceptible to fungal diseases, especially if they could not maintain body warmth in the setting of starvation.

Since there are reptiles today, it is clear that some reptiles survived the K-T boundary cataclysm. This raises the question, if reptiles were previously so successful, why did they not reclaim the earth to launch a second reptilian age? It is difficult to imagine how mammals could have replaced reptiles as the dominant land forms without some selection mechanism for this energetically costly lifestyle. This led me to propose the hypothesis that fungal proliferation after the devastation of the KT event preferentially selected for the fungal-resistant endothermic and hindered the re-emergence of a second reptilian age [4]. Although we do not know the timeline for the recovery of the planet climate, it is estimated that photosynthesis was shut down for 6 months and climate cooling persisted for at least 9 years [20], and the occurrence of a fungal bloom sufficient to have left fossil evidence implies that surviving animals were exposed to massive numbers of fungal spores. The darkened skies and cooler temperatures that accompanied the K-T cataclysm [20] would have shielded the sun and reduced the ability of ectothermic creatures such as reptiles to induce fevers by insolation, a necessary activity for protection against fungal diseases. Hence, it is reasonable to posit that ectothermic creatures unable to induce behavioral fevers and in weakened states from environmental stress would have been at a severe disadvantage relative to small mammals with their innate thermal exclusionary zones for fungal growth. Further complicating the situation for reptiles is that eggs can be vulnerable to fungal attack [21], whereas mammalian progeny would be protected in placenta.

Climate Change and Future Fungal Threats

Climate warming implies that the temperate gradient from mammals and average environmental temperatures will be reduced. Higher global temperatures could select for more thermally tolerant fungi, and it is possible that many fungi with

current pathogenic potential for mammals that are unable to cause disease in mammals due to thermal intolerance will acquire the capacity to survive at mammalian temperatures and thus become pathogenic for mammals [22]. This concern is heightened by the fact that some fungi can be easily adapted to higher temperatures by thermal selection, as exemplified by the generation of a thermally resistant entopathogenic fungus as an attempt to create a pest control strain that would be less susceptible to insect-induced behavioral fevers [23].

The Fungal-Mammalian Emergence Hypothesis in Context

The fungal-mammalian emergence hypothesis posits that fungi selected for the emergence of mammals. The hypothesis suggests an explanation for how the highly energy-intensive mammalian lifestyle was selected and for the relative resistance of immunologically intact mammals to fungal diseases. The hypothesis is a plausible synthesis assembled from very disparate lines of evidence. At this time it is unlikely that experimental evidence will be available in the near future to validate or refute this hypothesis simply by the very nature of what it tries to explain, and the remoteness of past events. For example, given that the animals that died as a result of the KT-related events represent an extremely small part of the fossil record, it is unrealistic to imagine finding fossils that could unequivocally be dated to the time in question with evidence of fungal disease. Fungal diseases can leave traces in the fossil record, as manifested by the finding of *Coccidioides*-like spherules in a fossil bison from the Holocene [24], but those fossils are very recent relative to the KT event and fungal effects on bone tissue usually reflect chronic infections. In contrast, fungal diseases caused by microscopic organisms that killed hosts by destroying soft tissues would leave no fossil record. On the other hand, recent developments with amphibian chytridiomycosis and the white nose syndrome in bats provide strong circumstantial evidence for the notion that fungal diseases could have provided strong selection pressures and driven some species to extinction. Although these are examples of individual fungal-host interaction in specialized ecological settings, they do provide precedents for the notion that fungi can be powerful selective forces for vertebrate species. In addition, there is now considerable evidence that fungi are potential threats to entire ecosystems [25]. The fungal-mammalian emergence hypothesis will likely continue to evolve as new information is available and is best considered as a cognitive tool for stimulating thinking and discussion on global issues related to evolutionary selection, infectious diseases, and ecological change.

References

1. Bakker RT (1970) Dinosaur physiology and the origins of mammals. *Evolution* 25: 636–658.
2. Dixon DM, McNeil MM, Cohen ML, Gellin BG, La Montagne JR (1996) Fungal infections: a growing threat. *Public Health Rep* 111: 226–235.
3. Hulbert AJ, Else PL (1981) Comparison of the “mammal machine” and the “reptile machine”: energy use and thyroid activity. *Am J Physiol* 241: R350–R356.
4. Casadevall A (2005) Fungal virulence, vertebrate endothermy, and dinosaur extinction: is there a connection? *Fungal Genet Biol* 42: 98–106.
5. Robert VA, Casadevall A (2009) Vertebrate endothermy restricts most fungi as potential pathogens. *J Infect Dis* 200: 1623–1626.
6. Perfect JR (2006) *Cryptococcus neoformans*: the yeast that likes it hot. *FEMS Yeast Res* 6: 463–468.
7. Perfect JR, Lang SDR, Durack DT (1980) Chronic cryptococcal meningitis. *Am J Pathol* 101: 177–193.
8. Obendorf DL, Peel BF, Munday BL (1993) *Mucor amphibionum* infection in platypus (*Ornithorhynchus anatinus*) from Tasmania. *J Wildl Dis* 29: 485–487.
9. Berger L, Speare R, Daszak P, Green DE, Cunningham AA et al. (1998) Chytridiomycosis causes amphibian mortality associated with population declines in the rain forests of Australia and Central America. *Proc Natl Acad Sci U S A* 95: 9031–9036.
10. Woodhams DC, Alford RA, Marantelli G (2003) Emerging disease of amphibians cured by elevated body temperature. *Dis Aquat Organ* 55: 65–67.
11. Blehert DS, Hicks AC, Behr M, Meteyer CU, Berlowski-Zier BM, et al. (2009) Bat white-nose syndrome: an emerging fungal pathogen? *Science* 323: 227.
12. Meteyer CU, Valent M, Kashmer J, Buckles EL, Lorch JM et al. (2011) Recovery of little brown bats (*Myotis lucifugus*) from natural infection with *Geomyces destructans*, white-nose syndrome. *J Wildl Dis* 47: 618–626.
13. Beernaert LA, Pasmans F, Van WL, Haesebrouck F, Martel A (2010) *Aspergillus* infections in birds: a review. *Avian Pathol* 39: 325–331.
14. Chang YC, Tsai HF, Karos M, Kwon-Chung KJ (2004) THTA, a thermotolerance gene of *Aspergillus fumigatus*. *Fungal Genet Biol* 41: 888–896.
15. Keller G, Sahni A, Bajpai S (2009) Deccan volcanism, the KT mass extinction and dinosaurs. *J Biosci* 34: 709–728.

16. Vajda V, Raine JL, Hollis CJ (2001) Indication of global deforestation at the Cretaceous-Tertiary boundary by New Zealand fern spike. *Science* 294: 1700–1702.
17. Vajda V, McLoughlin S (2004) Fungal proliferation at the Cretaceous-Tertiary boundary. *Science* 303: 1489.
18. Bowman BH, Taylor JW, White TJ (1992) Molecular evolution of the fungi: human pathogens. *Mol Biol Evol* 9: 893–904.
19. Clarke A, Portner HO (2010) Temperature, metabolic power and the evolution of endothermy. *Biol Rev Camb Philos Soc* 85: 703–727.
20. Pope KO, Baines KH, Ocampo AC, Ivanov BA (1997) Energy, volatile production, and climatic effects of the Chicxulub Cretaceous/Tertiary impact. *J Geophys Res* 102: 21645–21664.
21. Moreira PL, Barata M (2005) Egg mortality and early embryo hatching caused by fungal infection of Iberian rock lizard (*Lacerta monticola*) clutches. *Herpetological J* 15: 265–272.
22. Garcia-Solache MA, Casadevall A (2010) Global warming will bring new fungal diseases for mammals. *MBio* 1: e00061-10.
23. de CE, Jaronski S, Lyons B, Lyons TJ, Keyhani NO (2009) Directed evolution of a filamentous fungus for thermotolerance. *BMC Biotechnol* 9: 74.
24. Morrow W (2006) Holocene coccidioidomycosis: Valley Fever in early Holocene bison (*Bison antiquus*). *Mycologia* 98: 669–677.
25. Fisher MC, Henk DA, Briggs CJ, Brownstein JS, Madoff LC et al. (2012) Emerging fungal threats to animal, plant and ecosystem health. *Nature* 484: 186–194.