# Correspondence

# Evolution of the extinct Sabre-tooths and the American Cheetahlike cat

Ross Barnett<sup>1</sup>, Ian Barnes<sup>2</sup>, Matthew J. Phillips<sup>1</sup>, Larry D. Martin<sup>3</sup>, C. Richard Harington<sup>4</sup>, Jennifer A. Leonard<sup>5,6</sup>, and Alan Cooper<sup>7</sup>

The sabretooths (Smilodon and Homotherium) and the American cheetahlike cat (Miracinonyx) were the top predators in Late Pleistocene America, but became extinct about 13 thousand years ago [1]. As the evolutionary history of these taxa remains poorly understood [1,2], we analysed their phylogenetic relationship to extant felids. In contrast to previous molecular studies [3], our results show that the sabretooths diverge early and are not closely related to any living cats. This supports their morphological placement in a separate subfamily (Machairodontinae) [1]. Despite its remarkable morphological similarity to the African cheetah (Acinonyx jubatus), Miracinonyx appears to have evolved from a puma-like ancestor, presumably in response to similar ecological pressures [4,5].

Even though the evolutionary relationships within the cat family (Felidae) have been analysed using morphological, molecular and palaeontological data [1], many proposed relationships remain contentious, including those of recently extinct species. For example, the sabretooth cats are known from plentiful Late Pleistocene fossil material and have been placed, based on morphological criteria, outside of all extant cats (subfamily Felinae) as a separate subfamily, the Machairodontinae [1]. In contrast, an early ancient DNA study tentatively suggested that Smilodon was a member of the

Felinae and thus part of the modern cat radiation [3].

The relationships of the American cheetahlike cat, Miracinonyx trumani, are similarly confused as it had previously been mistaken for an ancestral or modern puma (e.g. Puma trumani). It was only recognised as a truly distinct group after abundant post-cranial fossil material became available [6]. Morphologically, M. trumani strongly resembles the modern African cheetah (Acinonyx jubatus) [4], which has led to uncertainty about whether it is more closely related to the Old World cheetah or the New World puma [2,4,6,7].

To address these questions, we used ancient DNA techniques, including independent replication, to generate mitochondrial DNA sequences from several specimens of *Smilodon populator* 

from Patagonia and a *M. trumani* specimen from Wyoming. In total, 1302 bp of *cytochrome b (cytb)*, *NADH5*, *ATP8*, as well as 12S and 16S ribosomal RNA genes were recovered (Supplemental Data). We also obtained a short fragment of *cytb* from a Yukon scimitartoothed cat (*Homotherium serum*), another recent member of the Machairodontinae.

Phylogenetic analyses (Figure 1; Supplemental Data) show that *Smilodon* and *Homotherium* fall outside the extant cats, the Felinae. This is in agreement with their traditional morphological placement [1]. The deep genetic split between the two sabretooth taxa is also consistent with morphological analyses [1].

These phylogenetic analyses also reveal that *M. trumani* is the sister taxon to the puma, rather than the African cheetah. Another

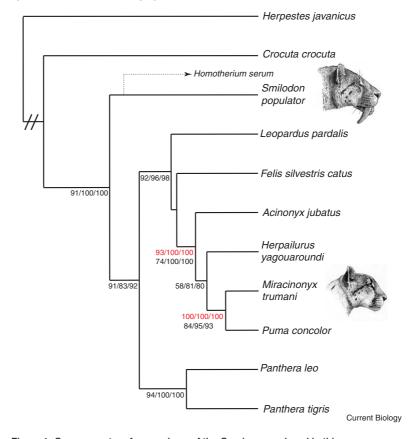


Figure 1. Consensus tree for members of the Carnivora analysed in this paper. Maximum parsimony bootstrap replicates and Bayesian posterior values are given for nodes. The numbers at the nodes show: maximum parsimony bootstrap support (1000 replicates), the Bayesian GTR+I+G posterior probabilities (5 million generations, burnin at 1 million) and the Bayesian GTR+cov+G posterior probabilities (5 million generations, burnin at 2.5 million). The values given in red represent results after omission of the jaguarundi from analysis (Supplemental Data). The maximum likelihood position of Homotherium serum is based on a short dataset (Supplemental Data). Figures of Smilodon and Miracinonyx courtesy of M. Anton.

endemic American cat, the jaguarundi, groups with Miracinonyx and the puma. This result corroborates the first morphological studies using complete specimens of Miracinonyx [6,7], but also raises questions about the anatomical structures used to classify Miracinonyx as a cheetah [4]. Reanalysis of these characters [2,7] has emphasised that many of them are associated with a highly cursorial lifestyle, such as elongated limbs and enlarged nares [2,4], which increase running speed and air-intake efficiency, respectively. In contrast, other limb characters, such as a well-developed anteriorly projecting flange on the head of the fibula, link Miracinonyx, puma and jaguarundi

It has been suggested that the cheetahs originated in the New World [4] and later migrated to the Old World. However, the mitochondrial sequence analysis together with recent fossil data (Supplemental Data) suggests that they originated in the Old World and that a puma-like cat then invaded North America around six million years ago [5,7,8]. Around 3.2 million years ago, this ancestor diverged into Miracinonyx and Puma, which is broadly contemporaneous with increasing prairie in North America [9]. The expansion of this habitat and its effect on ungulate prey, e.g. the pronghorn antelope (Antilocapra americana), may have driven the evolution of cursoriality in Miracinonyx, allowing it to excel in high-speed pursuit [1].

## Acknowledgments

We thank P.J.H. van Bree (Zoological Museum Amsterdam) for access to *Smilodon* samples, Lars Werdelin (Swedish Museum of Natural History) for helpful information on specimens, Andrew Kitchener (National Museum of Scotland) for modern felid samples and Trish McLenachan and David Penny (Massey University) for access to mongoose data. Work at UCLA on *Homotherium* was supported by an NSF grant to Robert Wayne and Blaire Van Valkenburgh (OPP-9617068). This research was supported by NSF (AC, JAL), NERC (RB and AC), Leverhulme

(AC), BBSRC (RB), and Wellcome (IB and AC). We also thank Mauricio Anton for use of his artwork.

### Supplemental data

Supplemental data including Experimental Procedures are available at http://www.currentbiology.com/cgi/content/full/15/15/Rxxx/ DC1/

### References

- Turner, A., and Anton, M. (1997).
  The Big Cats and Their Fossil
  Relatives (New York: Columbia
  University Press).
- Herrington, S.J. (1986).
   Phylogenetic relationships of the wild cats of the world. Ph.D thesis, University of Kansas.
- Janczewski, D.N., Yuhki, N., Gilbert, D.A., Jefferson, G.T., and O'Brien, S.J. (1992). Molecular phylogenetic inference from sabertoothed cat fossils of Rancho La Brea. Proc. Natl. Acad. Sci. USA 89. 9769–9773.
- Adams, D.B. (1979). The Cheetah: Native American. Science 205, 1155–1158.
- Hemmer, H., Kahlke, R.D., and Vekua, A.K. (2004). The Old World puma - Puma pardoides (Owen, 1846) (Carnivora: Felidae) - in the Lower Villafranchian (Upper Pliocene) of Kvabebi (East Georgia, Transcaucasia) and its evolutionary and biogeographical significance. Neues Jahrbuch für Geologie und Paläontologie 233, 197–231.
- Martin, L.D., Gilbert, B.M., and Adams, D.B. (1977). A Cheetah-like cat in the North American Pleistocene. Science 195, 981–982.
- van Valkenburgh, B., Grady, F., and Kurten, B. (1990). The Plio-Pleistocene Cheetah-like Cat Miracinonyx inexpectatus of North America. Journal of Vertebrate Palaeontology 10, 434–454.
- 3. Zhanxiang, Q. (2003). Chapter 2: Dispersals of Neogene Carnivorans between Asia and North America. Bull. Am. Mus. Nat. Hist. 279,
- Janis, C.M., Damuth, J., and Theodor, J.M. (2002). The origins and evolution of the North American grassland biome: the story from the hoofed mammals. Palaeogeogr. Palaeoclimatol. Palaeoecol. 177, 183–198.

<sup>1</sup>Henry Wellcome Ancient Biomolecules Centre, Department of Zoology, University of Oxford, South Parks Road, Oxford OX1 3PS, UK. <sup>2</sup>The Centre for Genetic Anthropology, Department of Biology, Darwin Building, University College London, Gower Street, London WC1E 6BT, UK. <sup>3</sup>Department of Ecology and Evolutionary Biology, University of Kansas, Lawrence, KS66045, Kansas, USA. <sup>4</sup>Canadian Museum of Nature (Paleobiology), Ottawa, Ontario, Canada K1P 6P4. <sup>5</sup>Department of Ecology and

Evolutionary Biology, University of California, Los Angeles, CA 90095-1606, USA. <sup>6</sup>Department of Evolutionary Biology, Uppsala University, Norbyvagen 18D, 752 36 Uppsala, Sweden. <sup>7</sup>Darling Building (DP 418), School of Earth and Environmental Sciences, University of Adelaide, SA 5005, Australia. E-mail: ross.barnett@zoo.ox.ac.uk; alan.cooper@adelaide.edu.au