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and ERIC DELSON². ¹Department of Computer Science, University of California at Davis, ²Department of Anthropolgy, Lehman College of City University of New York, ³Department of Statistics, State University of New York at Stony Brook, ⁴Department of New York at Stony Brook, ⁵Department of Anthropology, University of Oregon, ⁵Stratovan Inc, Davis CA, ⁶Department of Computer Science, Lehman College of City University of New York.

Phylogenies for groups of living organisms are routinelyconstructed based on molecular data. These phylogenies generate hypotheses about the morphology of implied ancestor species, often in the form ancestral morphotypes.

present some computer visualizations of the crania of the hypothetical ancestors of the Old World monkeys, based on the crania of their living descendants. These reconstructions are based on geometric morphometric techniques using a large series of landmarks and semilandmarks on laser surface scans to estimate the morphology of ancestor species. Ancestral morphology is based on an estimated landmark and configuration based on those of extant sepcies weighted by phylogenetic distance using squared change parsimony. Phylogenetic distance is based on current molecular phylogenies from the literature and their estimated divergence dates.

Our reconstructions of the basal ancestors clearly differ from the oldest fossils for the group. We consider several of the technical issues involved in integrating the fossil information into the morphological hypothesis, and we demonstrate how including the fossils improves the estimated morphology and can affect the inferences we draw from the tree. This study was funded by the NSF, grants number 0513660 and 0513894.

Evolutionary morphing.

NINA AMENTA¹, DEBOSHMITA GHOSH¹, ANDREI SHARF¹, F. JAMES ROHLF³, WILL HARCOURT-SMITH², STEVE FROST⁴, DAVID WILEY³, KATHERINE ST. JOHN⁶