

Rediscovery of an “extinct” Galápagos tortoise

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Over the past three centuries, humans have demonstrated increasing scientific curiosity about biodiversity. During this time, we developed a classification scheme that accelerated the description and codification of species (1), profound new theories to explain diversity (2), explicit frameworks of taxonomic study (3), and DNA-based methods for characterizing lineages (4). Despite these academic advancements, the progress of biological discovery has struggled to keep pace with an increasingly modified natural environment, because the past 300 years have also witnessed the exponential growth of human population (from 6×10^8 to 6×10^9) and concomitant anthropogenic impacts on biodiversity. Human-mediated extinctions, translocations, and genetic pollution obscure and erode natural patterns of distribution and variation (which is especially true for economically and calorically valuable species, like turtles). A study by Poulakakis *et al.* (5) in this issue of PNAS wrestles with all three of these complicating factors and provides an excellent example of the resources and methodologies required to tease apart patterns resulting from natural and artificial processes. In doing so, this study reaches an astonishing conclusion: A species that we thought was eaten to extinction still survives. . . in part.

The species in question is a Galápagos giant tortoise, one of 15 species (13 formally described) from the famous archipelago (5). The Galápagos Islands are well known for bearing the fauna that helped inspire Darwin to develop his landmark theory of natural selection (2). The giant tortoises should be given partial credit for this inspiration, because the diagnostic phenotypes on each island provided Darwin with a clear example of regional variation and adaptation (6). But even as Darwin observed these tortoise populations, they were declining precipitously. Humans had already altered the tortoises' habitat by introducing numerous feral mammals, but primarily reduced their numbers through direct harvest for consumption [even Darwin ate tortoises (6)]. Darwin and the tortoises exemplify the aforementioned challenge of biologists to uncover and record biological patterns while humans simultaneously obscure them. In fact, Darwin arrived at one of the islands, Floreana (also known as

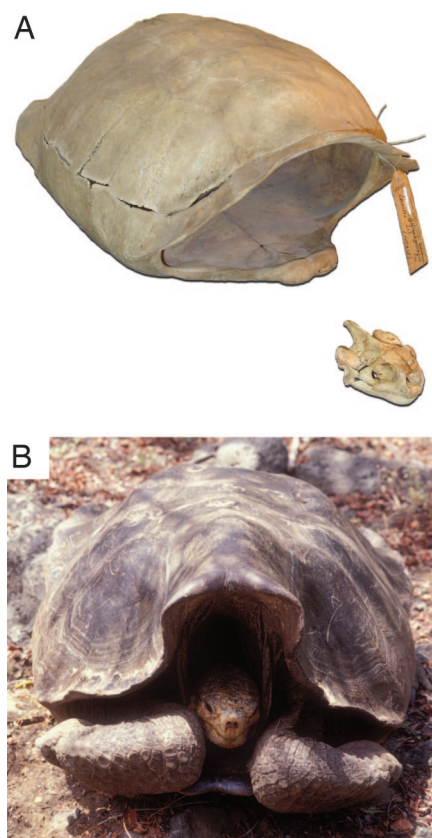


Fig. 1. Galápagos tortoises. (A) A subfossil specimen of the species from Floreana Island (Field Museum of Natural History no. 13525). Until the study by Poulakakis *et al.* (5), the Floreana species was only known from museum specimens such as this one. (B) A living descendant of the “extinct” Floreana species from Isabela Island (photo by C. Ciofi).

Santa María or Charles), in 1835, within years of the putative extinction of its endemic form (7). Both Darwin and the captain of the HMS Beagle remarked on the depredated carcasses and scarcity of the Floreana tortoise, seeing no live tortoises themselves (6, 7). In the case of Floreana, the ultimate result of human activities, extirpation, preceded scientific observation and collection. Consequently, much of what we know about the Floreana species is based on subfossil skeletal remains from caves (8, 9) (Fig. 1A). Because extinction is a permanent phenomenon, the Floreana tortoise seemed destined to remain another sad footnote in biodiversity registers, lost like the dodo.

The report by Poulakakis *et al.* (5) provides new data showing that descen-

dants of Floreana tortoises actually still exist (albeit not on Floreana) (Fig. 1B). That it is still possible to find such an important result within a well studied group of species is at first surprising. But, in fact, it is the detailed genetic studies of Galápagos tortoises by the Yale University-led team (10–12) that provided the necessary background to achieve this insight. Poulakakis *et al.* augment and use their genetic database to address lineage diversity from a population genetics perspective, applying appropriate markers and methods to elucidate the ancestry of extant individuals. But their work is more than a well executed genetic survey of a celebrated clade. One of the salient aspects of their study is the incorporation of data from long-dead Floreana tortoises into their genetic database. By using historic DNA techniques, Poulakakis *et al.* are able to sequence museum specimens from Floreana. Without the incorporation of these specimens, the true ancestry of the Floreana descendants would still be a mystery. Indeed, their study is another shining example of the irreplaceable role of museum specimens for biodiversity science. No matter how fast and powerful genomics becomes, we always need museums to inform, identify, and compare our findings (13). Older, historic collections are especially valuable, irreplaceable even, because they archive biological data from a time when life on Earth was more abundant and diverse.

The discovery of Floreana tortoise descendants is unexpected, but the circumstances that led to the preservation of this lineage are nothing short of ironic. It was, after all, the massive harvest of Floreana tortoises that ultimately led to the persistence of this lineage. Whereas turtle populations may be vulnerable to decimation, individual animals are durable and capable of surviving great physical hardship (such as desiccation and starvation in the hull of a ship). This preservation potential, combined with a wide appreciation for the taste of tortoise flesh, made Galápagos animals a highly sought after source of protein for

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