

Historical genomes reveal recent changes in genetic health of eastern gorillas

December 27 2018



Historical collections of eastern gorilla specimens spanning the last 100 years were used to provide a glimpse into the recent past of this critically endangered species. Credit: Katerina Guschanski

The critically endangered Grauer's gorilla has recently lost genetic diversity and has experienced an increase in harmful mutations. These conclusions were reached by an international team of researchers who sequenced eleven genomes from eastern gorilla specimens collected up to 100 years ago, and compared these with genomes from present-day individuals. The results are now published in *Current Biology*.

Many wild animals have declined in numbers over the past century, and scientists have long worried that these declines have resulted in losses of genetic diversity, increased inbreeding and an accumulation of harmful mutations. Although this could lead to an even higher risk of extinction in threatened species, investigating recent changes in genetic viability has been difficult. In a new study, a team led by scientists from Uppsala University and the Swedish Museum of Natural History has used specimens stored in [museum collections](#) to analyse changes in eastern gorilla genomes over the past 100 years.

"We found that the [genetic diversity](#) in Grauer's gorilla has declined significantly in just a few generations," says Tom van der Valk, a Ph.D. student at Uppsala University in Sweden.

Grauer's gorillas are found in the Democratic Republic of the Congo and have declined by 80 per cent in recent decades due to poaching and habitat destruction. The results from the comparison of historical and modern genomes show that this [decline](#) has led to increased inbreeding and a loss of genetic variation. This in turn means that Grauer's gorillas have likely become less able to adapt to future disease outbreaks and changes in their environment. In addition, the scientists identified several mutations that are probably harmful and that have increased in frequency over the past four to five generations as a consequence of the decline in [population size](#). In the closely related mountain gorilla, however, the scientists did not discover any significant genetic changes, suggesting that its genetic viability has remained stable over the past 100

years.

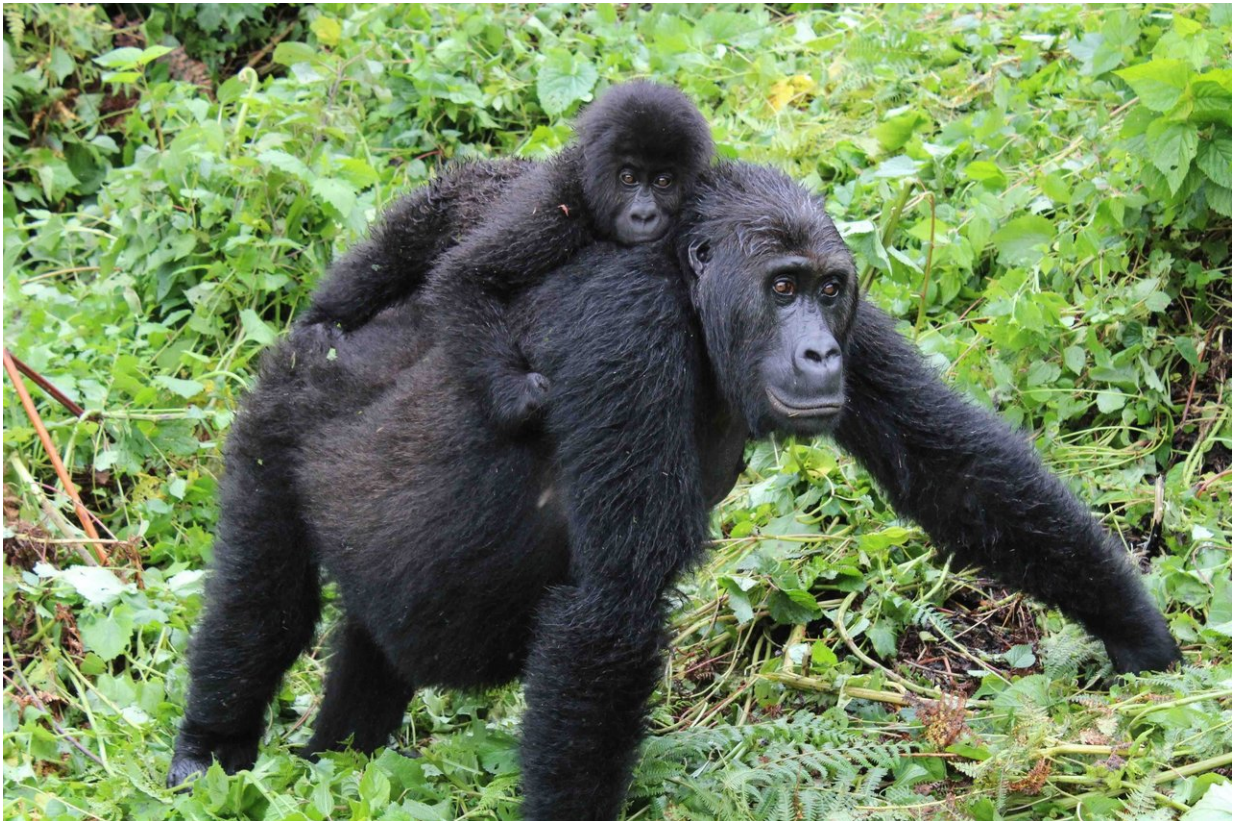


The magnificent Grauer's gorillas have experienced severe population declines in the last two decades. This population loss has left a deep mark in the genomes of this critically endangered great ape. Credit: Damien Caillaud, Dian Fossey Gorilla Fund International.

"This recent increase in harmful mutations really emphasises the need to reverse the ongoing population decline in Grauer's gorillas," says Love Dalén at the Swedish Museum of Natural History.

Some of the potentially harmful mutations that have increased in frequency were found in genes that affect disease resistance and male fertility. In addition, the researchers identified [mutations](#) leading to loss of function in genes associated with finger and toe development, which likely explains why present-day gorillas sometimes have fused digits.

"Our study highlights that historical museum specimens constitute a unique resource for monitoring recent changes in the genetic status of endangered species," says Katerina Guschanski at Uppsala University.



The magnificent Grauer's gorillas have experienced severe population declines in the last two decades. This population loss has left a deep mark in the genomes of this critically endangered great ape. Credit: Amy Porter, Dian Fossey Gorilla Fund International

Interestingly, the reason why Grauer's gorillas have been more severely affected than mountain gorillas may lie in their deeper history. While Grauer's gorillas went through a major increase in numbers between 5,000 and 10,000 years ago, mountain [gorillas](#) have been rare for several thousands of years. This long-term small population size may have enabled natural selection to remove [harmful mutations](#) before mountain gorilla numbers started to decline in the 20th century.

More information: van der Valk T, Díez-del-Molino D, Marques-Bonet T, Guschanski K, Dalén L. Historical genomes reveal the genomic consequences of recent population decline in eastern gorillas. *Current Biology* (2018). [DOI: 10.1016/j.cub.2018.11.055](https://doi.org/10.1016/j.cub.2018.11.055)

Provided by Uppsala University

Citation: Historical genomes reveal recent changes in genetic health of eastern gorillas (2018, December 27) retrieved 18 April 2024 from <https://phys.org/news/2018-12-historical-genomes-reveal-genetic-health.html>

This document is subject to copyright. Apart from any fair dealing for the purpose of private study or research, no part may be reproduced without the written permission. The content is provided for information purposes only.