







Professor Milford H. Wolpoff from the University of Michigan (USA), one of the first advocates of the hypothesis about the polycentric origin of man. *Altai Mountains, 2018*



Excursion in Denisova Cave. Center: Viviane Slon, a paleogeneticist from Prof. S. Pääbo's laboratory. Her presentation was devoted to the sensational results of Denisova-11 DNA sequencing. *Altai Mountains, 2018*

Although today many people still believe in the divine act of man's creation, over the last hundred years science has collected vast compelling evidence of the long and gradual evolution of our very distant ancestors, who branched about 6–7 million years ago from the common evolutionary trunk they shared with apes. However, human genesis turned out to be not near as simple and straightforward as the first researchers believed it to be.

This picture became much clearer through the development of paleogenetics, a science that emerged in the 1980s at the junction of molecular biology, classical archeology and paleontology. Now scientists have learned to isolate and identify DNA sequences from archeological and paleontological remains and can study, from this perspective, populations of organisms that had vanished from the Earth, including human ancestors.

Until recently, one of the most likely candidates for the role of our ancestors has been the Neanderthal man, whose brain volume would match that of modern humans and who felt comfortable at temperate latitudes. However, based on the results of the first paleogenetic

studies of the mitochondrial DNA from fossil bone remains, scientists inferred that the Neanderthals had been a dead-end branch of mankind. New results rehabilitated them only at the beginning of the new millennium, with the emergence of advanced DNA sequencing technology, able to “read” nuclear DNA and prove that the genome of an average human with non-African roots has 1–2% of Neanderthal genes.

The Max Planck Institute for Evolutionary Anthropology (Germany) analyzed bone anthropological remains of the Neanderthals, including specimens from the Altai caves—Denisova and Okladnikova. These specimens showed exceptional preservation due to a special temperature regime in the caves. Among them was a small bone from Denisova Cave—a phalanx of a child's little finger, which created a real sensation.

Professor Svante Pääbo, Director of the Department of Genetics at the Max Planck Institute, recalls that in the spring of 2009, they received another bone fragment from Anatoly Derevyanko, found the previous year in Denisova Cave. The bone was tiny, and Pääbo thought it unimportant

and only decided to run some mtDNA tests, when time permits. But then came the unexpected: “On December 3, 2009, I was attending a meeting on the rat genome... As I walked from the dining hall to the lecture hall after breakfast, my cell phone rang. It was Johannes Krause calling from Leipzig and he sounded strangely excited. I asked him what the matter was. He asked me if I was sitting down. When I said no, he said I'd better sit down before hearing what he had to tell me...” The stunning news was the discovery of a new human species, unknown to science, which literally turned upside down the established ideas about the origin and evolution of *Homo sapiens*.

Paleogenetic analysis of the fossil bone remains of the Denisovan man showed that these hominins, like the Neanderthals, left a mark in the genome of many modern human populations. Judging by the genetic diversity, it was the Denisovans who dominated in North Asia in the Stone Age.

The fact that archaeologists find bone remains of the Denisovans and Neanderthals in the same cultural layers of Denisova Cave suggests that these hominins lived this area at the same time. But how did they get along with each other? We cannot answer this question decisively, but the last unique find speaks volumes.

Denisova 11 is a small bone whose exact position in the skeleton remains unknown. By comparing it with the known long bones of Neanderthals and modern humans, scientists found that it belonged to a female not younger than 13 years of age. After sequencing the nuclear DNA extracted from the specimen, scientists associated the bone with a young crossbred girl, whose nuclear DNA contains about 42% Denisovan genes and 39% Neanderthal ones. In other words, her ancestors came from the two

