



said Professor Willerslev, Lundbeck Foundation Professor at the University of Copenhagen and Prince Philip Professor at the University of Cambridge. "Our diet changed as we developed from hunter-gatherers into farmers, our settlement patterns changed, and there have been changes in pressure of infection from the pathogenic micro-organisms to which we were exposed due to altered living conditions. We also know that chronic viral, bacterial and fungal infections might be causative factors in neuropsychiatric diseases, so there is every reason to believe that the analyses of DNA from this period will show significant trends, giving us the ability to create new, publicly available reference sets, to enhance both the scientific and healthcare communities' understanding of disease evolution."

In order to decode the genetic origins and evolution of human diseases, Professor Willerslev and his team will utilize Illumina's most powerful sequencer, the NovaSeq™6000 System, which was designed to open new horizons for larger, more robust experiments, providing the throughput to properly power large-cohort studies. Projects of this scale benefit from speed, throughput and data quality, and as such this project will leverage the S4 flow cell to sequence up to 20 billion ancient DNA fragments every two days.

"The NovaSeq 6000 was the obvious choice for this project with its unrivaled data quality and high-throughput capabilities," remarks Professor Willerslev. "While we conceived this project to explore the evolutionary origins of genetic disorders years ago, it was simply impossible to realize before Illumina's NovaSeq System came on the market. We are delighted that Lundbeck Foundation had the foresight to see the importance of our project and that Illumina's technology will make the research possible."

"Illumina is thrilled to be working with Professor Willerslev and his team on this extraordinary project," said Paula Dowdy, Senior Vice President and General Manager for Europe, Middle East and Africa at Illumina. "It will be extremely valuable, if by going back 10,000 years, we can acquire new information about when, and under which environmental conditions, a brain disorder may have been introduced into human DNA. This project has the potential to influence future product developments in genetics and precision medicine by providing invaluable insights to those affected by mental health issues."