Using ancient DNA from Sardinia to assess population stability from the Neolithic to present

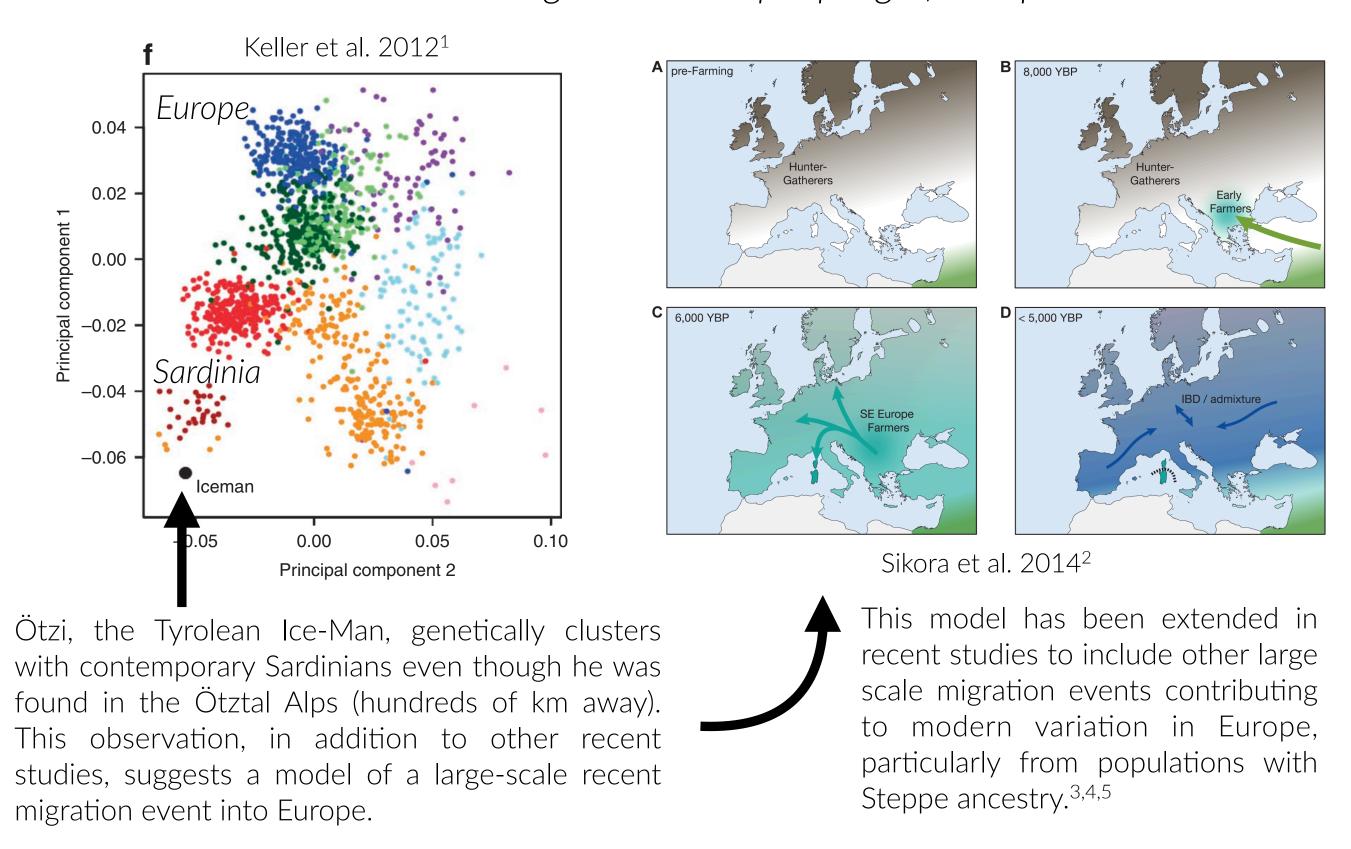


Joseph Marcus¹, Cosimo Posth^{7,8}, Luca Lai¹¹, Carlo Sidore^{4,5,6}, Jessica Beckett¹⁴, Hussein Al-asadi^{2,3}, Anja Furtwängler⁸, $Magdalena\ Zoledziewska^4$, Charleston Chiang 12,13 , Kushal Dey 3 , Maria Giuseppina Gradoli 15 , Goncalo Abecasis 5 , David Schlessinger⁹, Robin Skeates¹⁰, Johannes Krause^{7,8}, Francesco Cucca^{4,6}, John Novembre^{1,2}

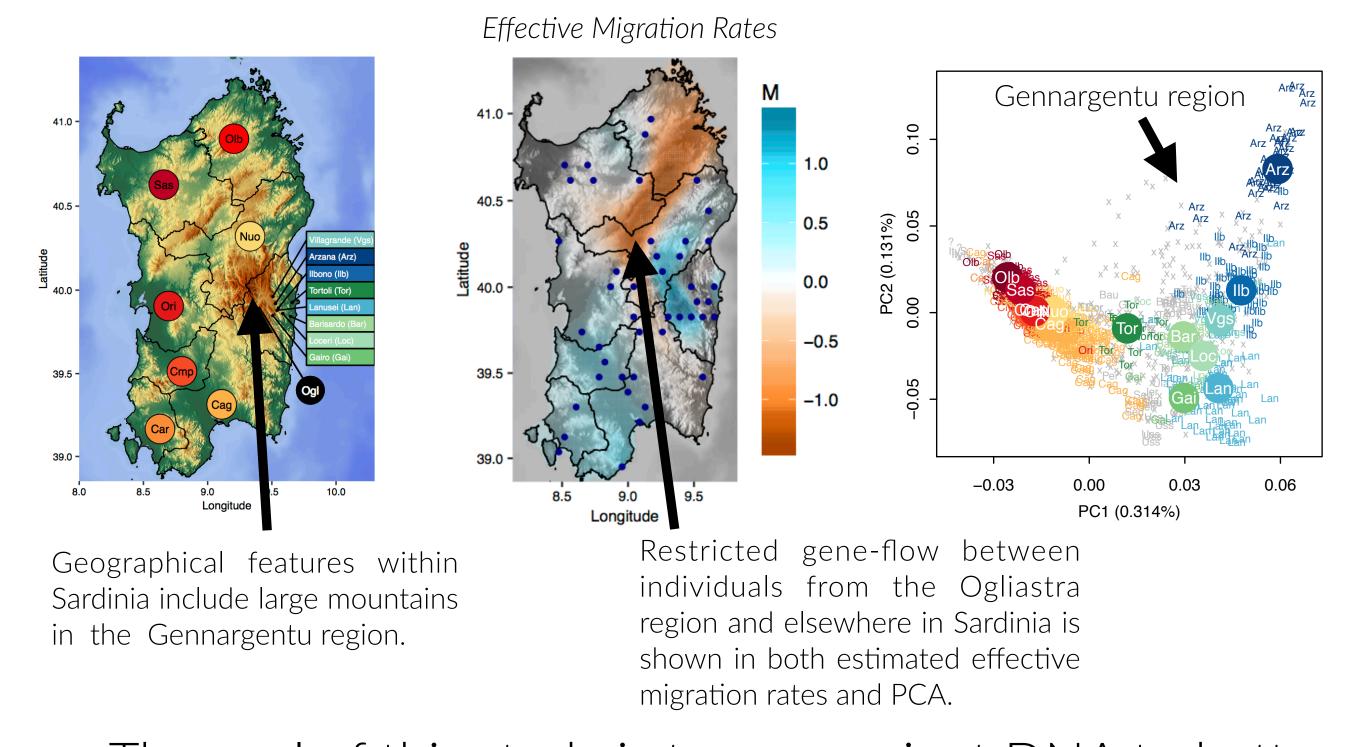
1. Department of Human Genetics, University of Chicago, Chicago, Chicago, Chicago, IL. 2. Department of Ecology and Evolution, University of Chicago, Chicago, IL. 3. Department of Statistics, University of Chicago, IL. 4. Istituto di Ricerca Genetica e Biomedica, CNR, Cagliari, Italy. 5. Center for Statistical Genetics, Department of Statistics, University of Chicago, IL. 3. Department of Statistics, University of Chic Biostatistics, University of Michigan, Ann Arbor, Ml. 6. Dipartimento di Scienze Biomediche, Università di Sassari, Italy. 7. Max Planck Institute for Archaeological Sciences, Archaeo- and Palaeogenetics, University of Tübingen, 9. Tübingen, Germany. 10. Laboratory of Genetics, NIA, NIH, Baltimore, MD. 11. Department of Archaeology, University of South Florida, Tampa, FL. 13. Center for Neurobehavioral Genetics, Semel Institute for Neuroscience and Behavior, University of California, Los Angeles, Los Angeles, California. 14. Department of Ecology and Evolutionary Biology, University of California, Los Angeles, California. 15. Independent Contractor, Cagliari, Italy. 16. School of Archaeology and Ancient History, University of Leicester, Leicester, United Kingdom.

Background

Genetic studies of the Sardinian population have revealed insights into the peopling of Europe

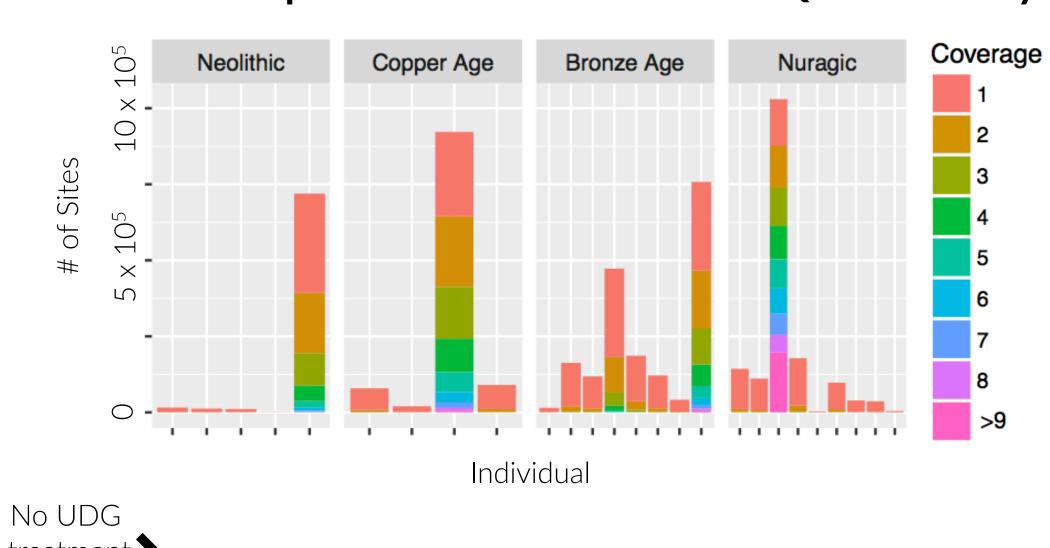


Fine-scale population structure within Sardinia (Chiang et al. 20176)

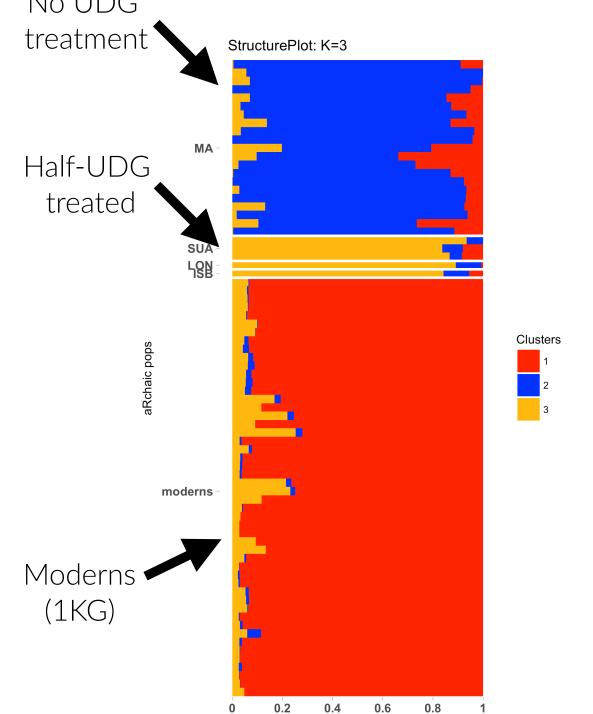


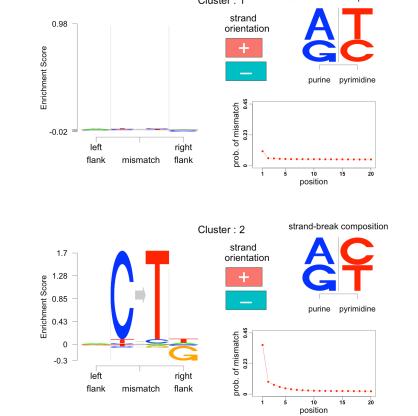
The goal of this study is to use ancient DNA to better understand the population history of the Sardinian people.

Sample Info / Quality Control



We generated low-coverage capture data at a targeted set of ~1240k SNPs in 26 individuals from different time periods in Sardinia's pre-history.





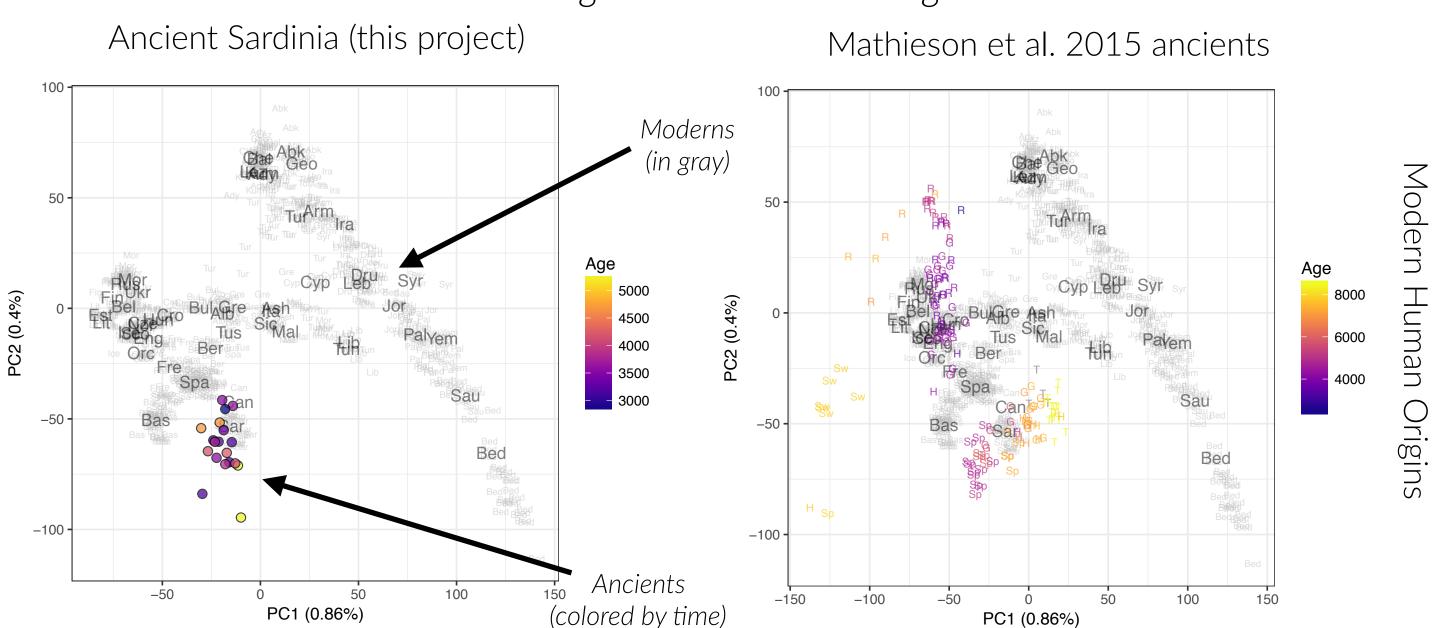
Visualizing ancient DNA damage patterns with a mixed-membership model using aRchaic⁷ shows ancients have a distinct damage profile when compared to moderns.



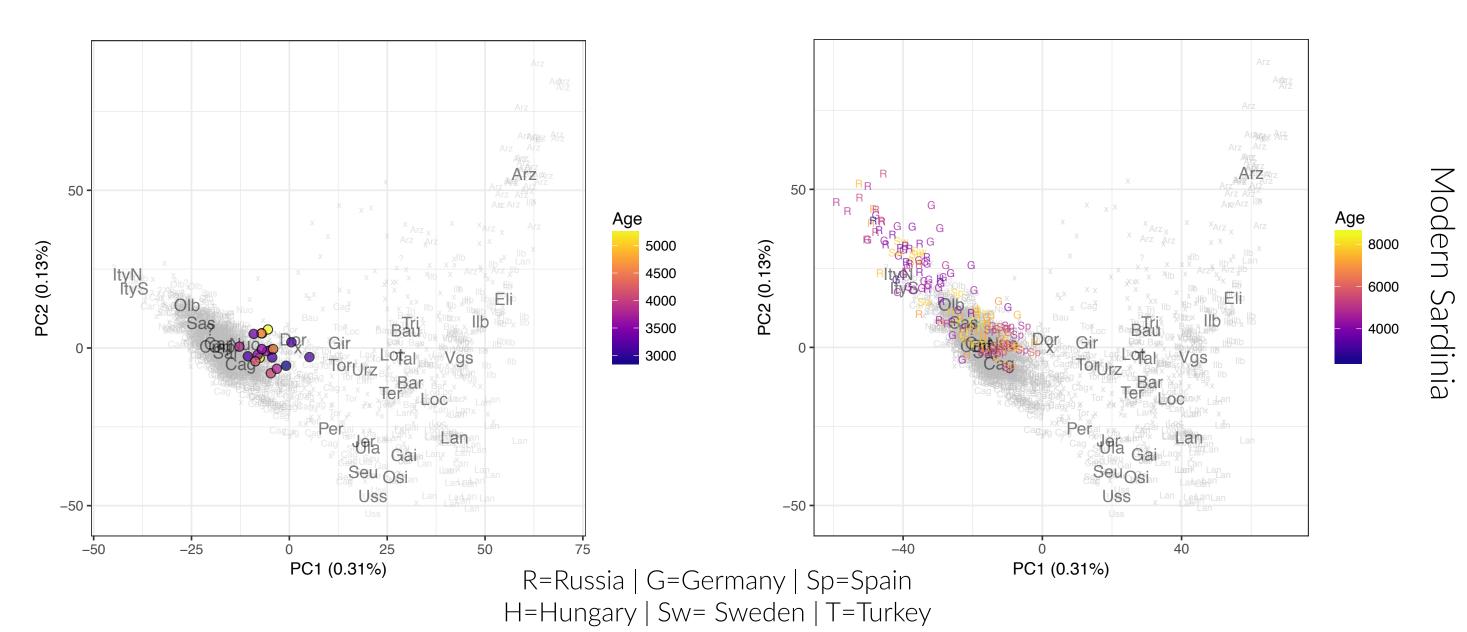
We additionally performed a number of QC steps to filter out individuals who have high contamination rates or too few covered SNPs.

Population Structure

Projecting ancient individuals onto axes defined by modern genetic variation using PCA

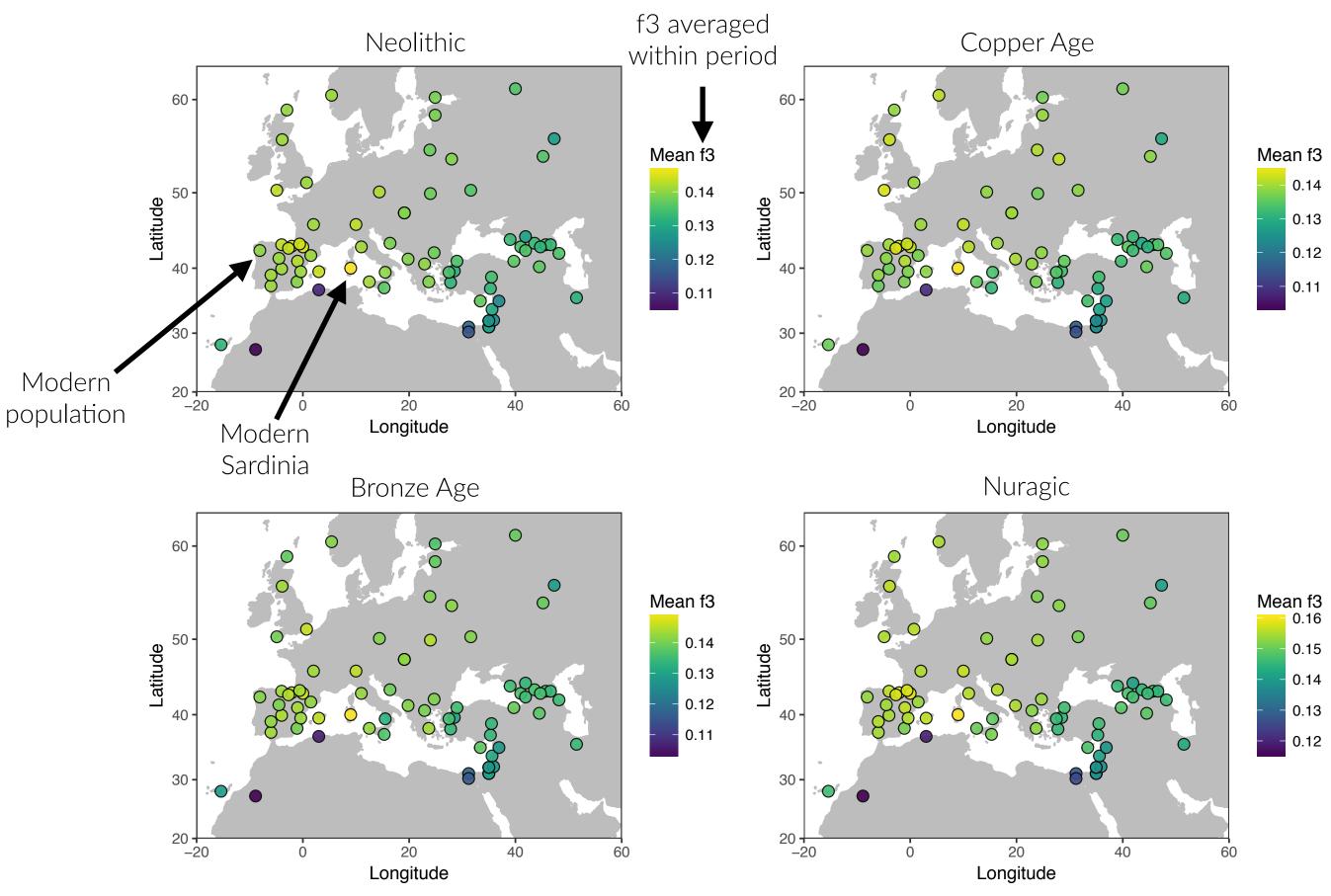


Ancient Sardinians from different time periods cluster with modern Sardinians. This type of temporal stability is not as persistent in mainland Europe.



Ancient individuals project close to Sardinian sub-populations outside the Gennargentu region.

Visualizing shared genetic drift between ancient Sardinia and modern populations through time

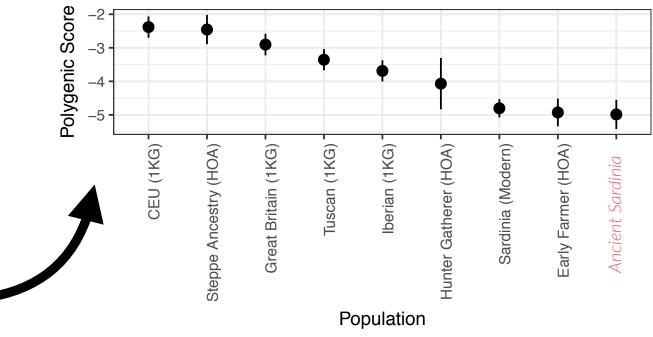


Modern Sardinia and Basque populations shows the strongest shared genetic drift with ancient Sardinia through multiple time periods.

Conclusions & Future Directions

We observe temporal stability within Sardinia when contrasted to other ancient individuals from mainland Europe.

We are conducting an ongoing study of the evolutionary forces acting on quantitative traits through time in Sardinia (particularly height and auto-immune diseases)



⁷Al-Asadi, H.*, Dey, K.*, Novembre, J., Stephens, M. (2017). In Prep.