

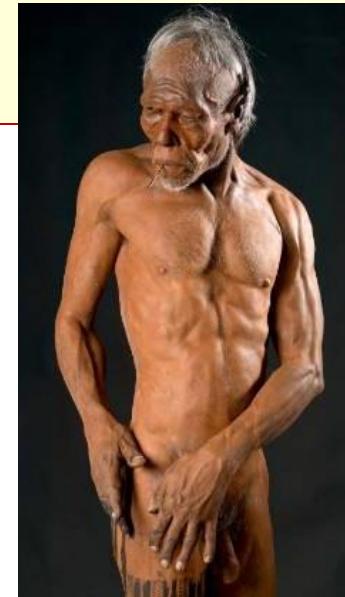
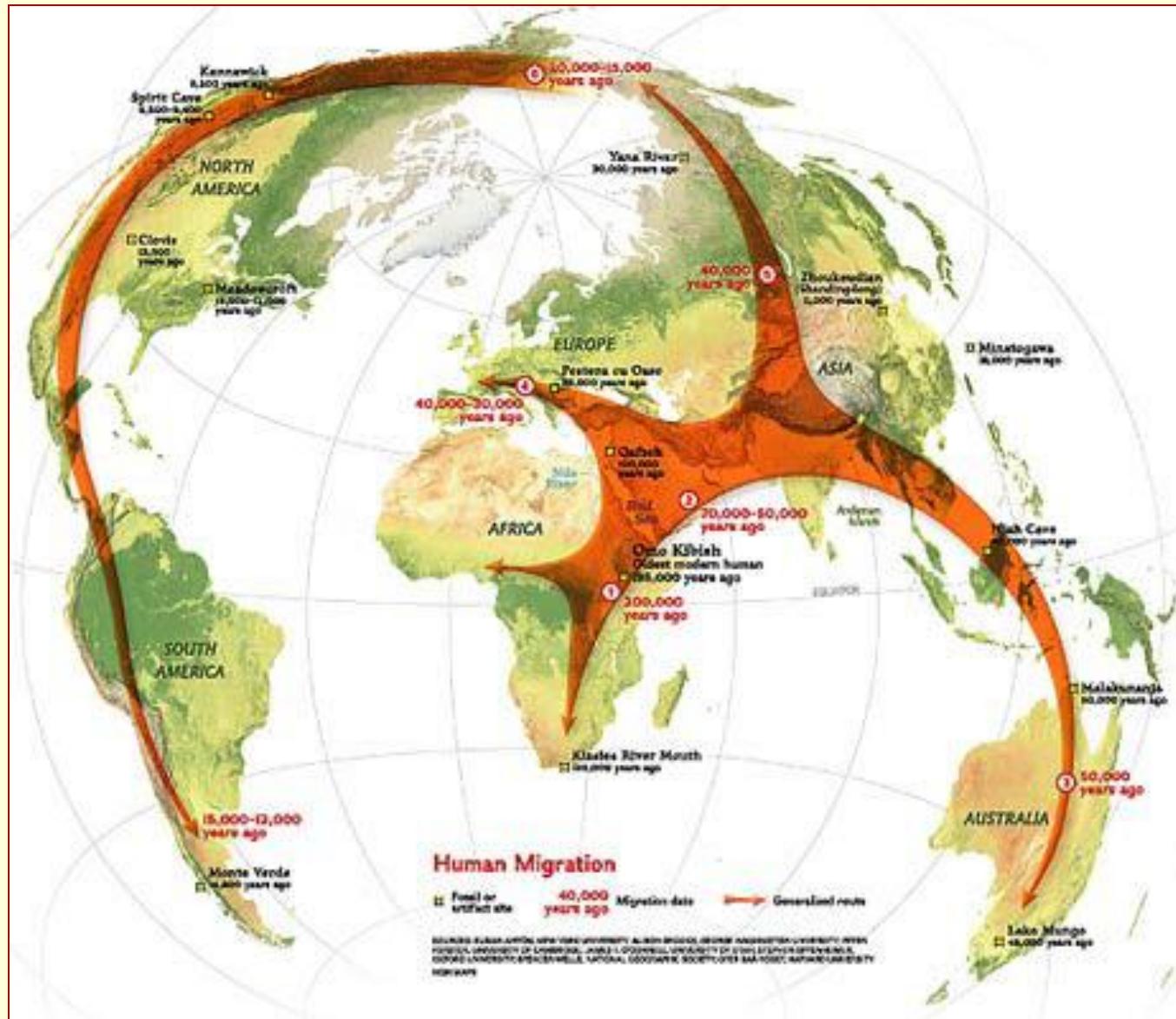
Università
degli Studi
di Ferrara

DIVULGAZIONE E FRUIZIONE DEL PATRIMONIO ANTROPOLOGICO

Lezione 3. Evoluzione umana dall'origine africana alla diffusione di *Homo sapiens*

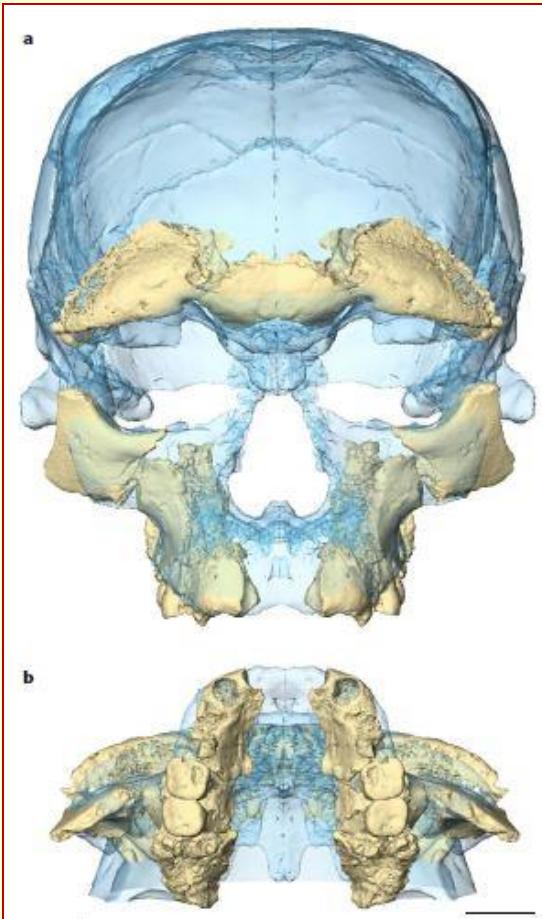
Marco Peresani
Dipartimento di Studi Umanistici
Sezione di Scienze Preistoriche e Antropologiche

Homo sapiens



New fossils from Jebel Irhoud, Morocco and the pan-African origin of *Homo sapiens*

Jean-Jacques Hublin^{1,2}, Abdelouahed Ben-Ncer³, Shara E. Bailey⁴, Sarah E. Freidline¹, Simon Neubauer¹, Matthew M. Skinner⁵, Inga Bergmann¹, Adeline Le Cabec¹, Stefano Benazzi⁶, Katerina Harvati⁷ & Philipp Gunz¹

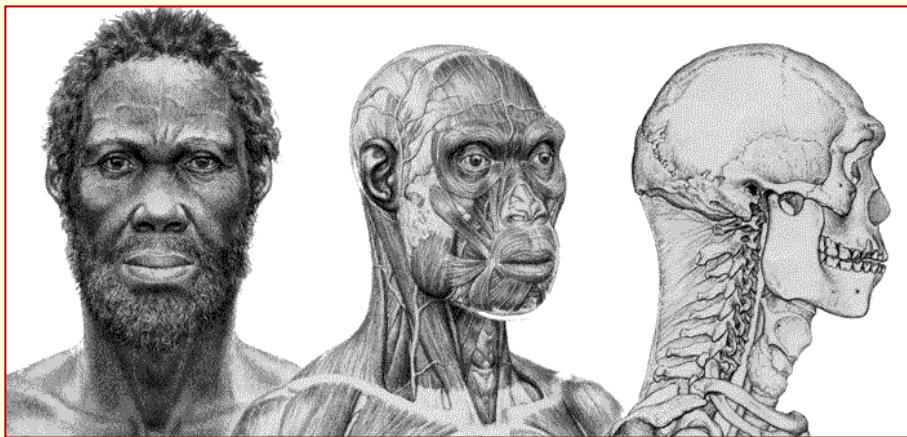


The earliest H.s.

"We identified a **mosaic of features** including facial, mandibular and dental morphology that aligns the Jebel Irhoud material with early or recent anatomically modern humans and more primitive neurocranial and endocranial morphology."

In combination with an age of **315 ± 34 thousand years**, this evidence makes Jebel Irhoud the oldest and richest African Middle Stone Age hominin site that documents early stages of the *H. sapiens* clade in which key features of modern morphology were established.

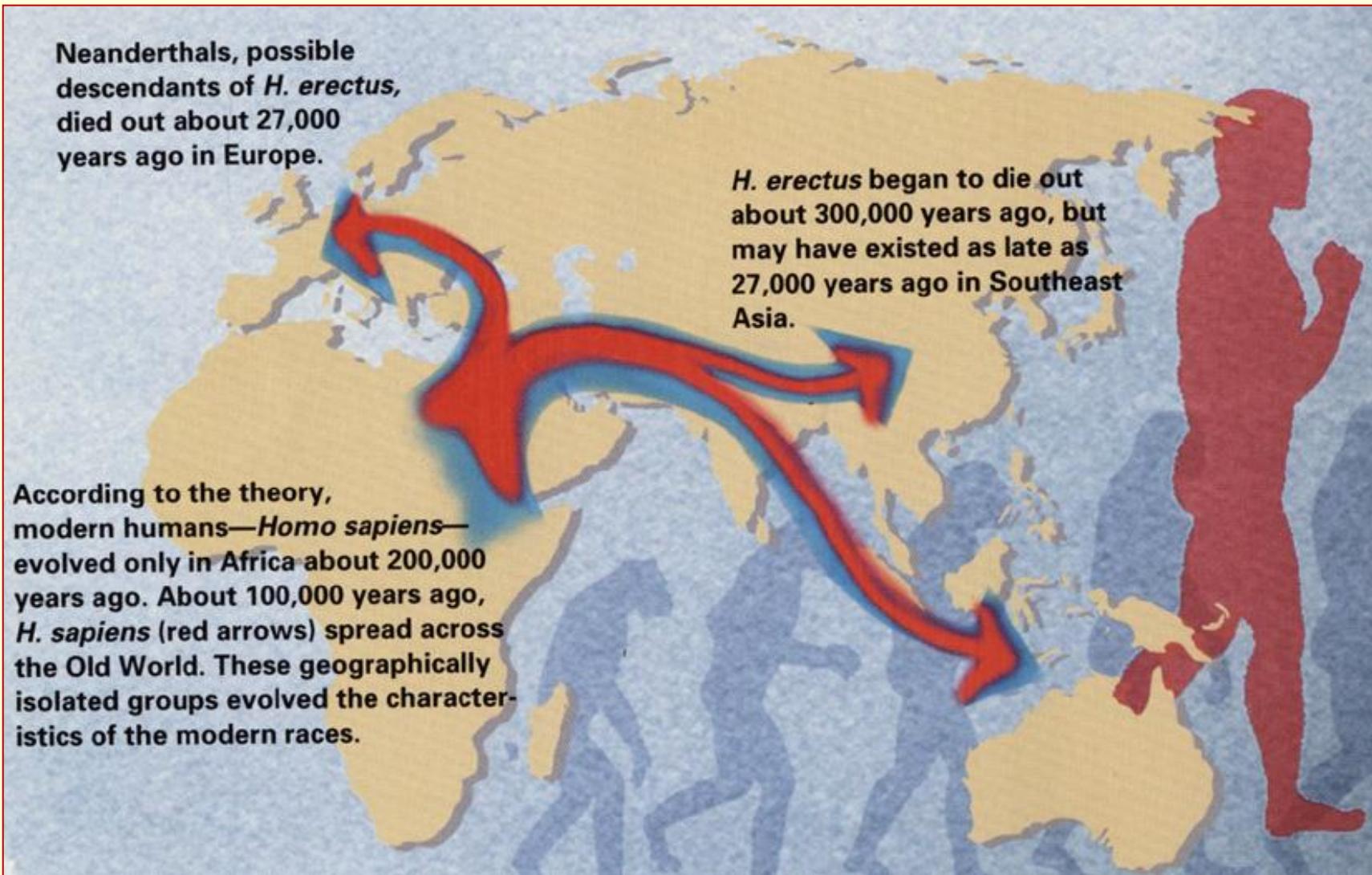
Furthermore, it shows that the evolutionary processes behind the emergence of *H. sapiens* involved the whole African continent."



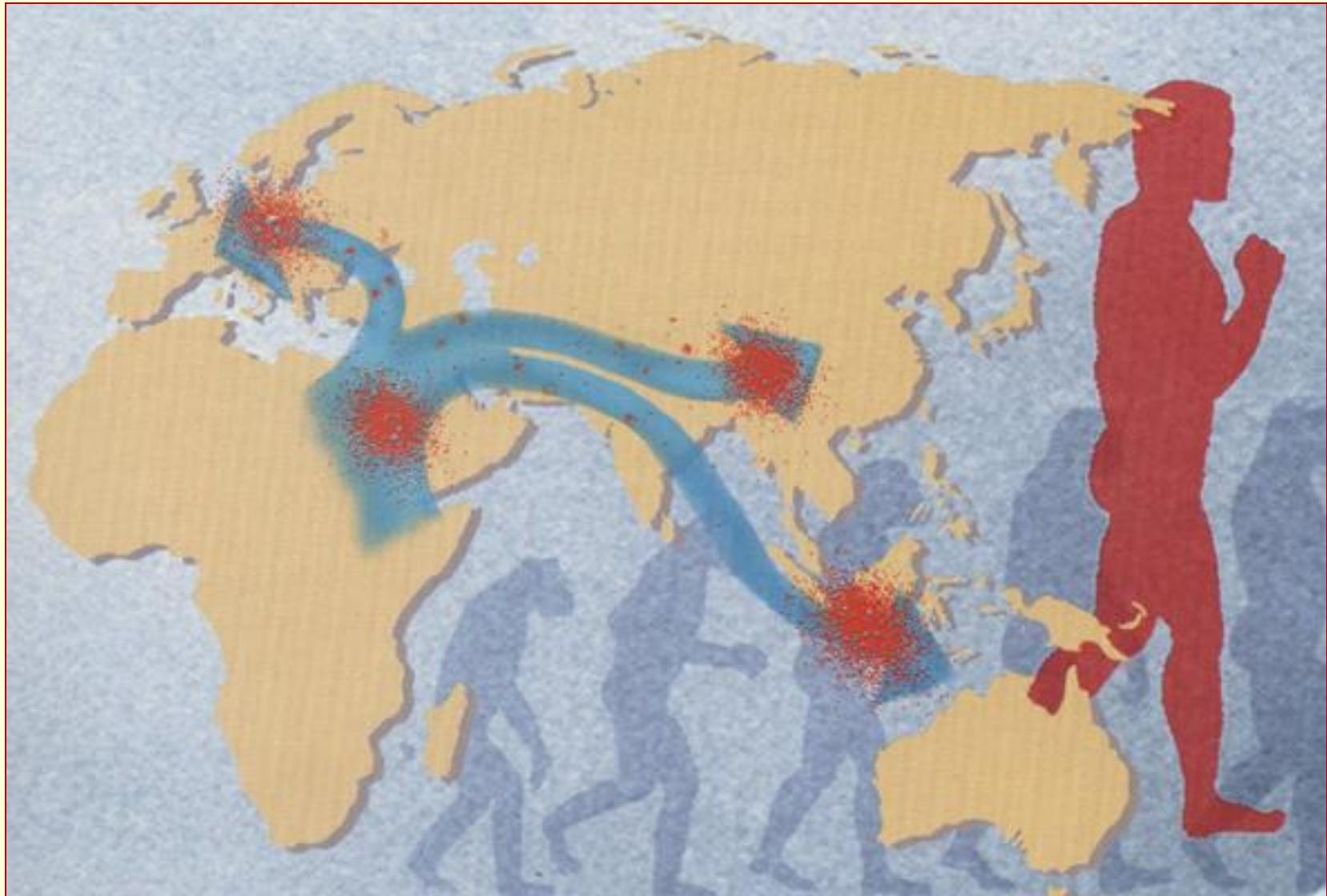
Homo sapiens hidaltu, 160K



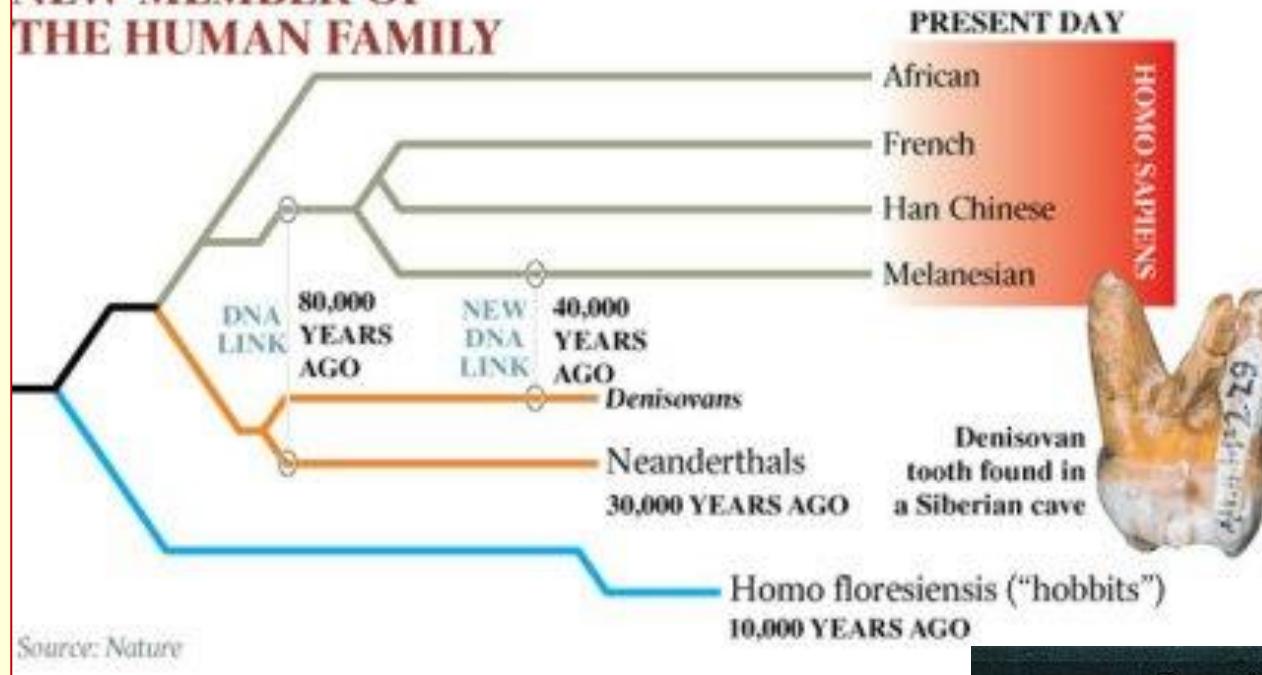
Substitution model about the appearance of Modern Humans



Multiregional model about the appearance of Modern Humans

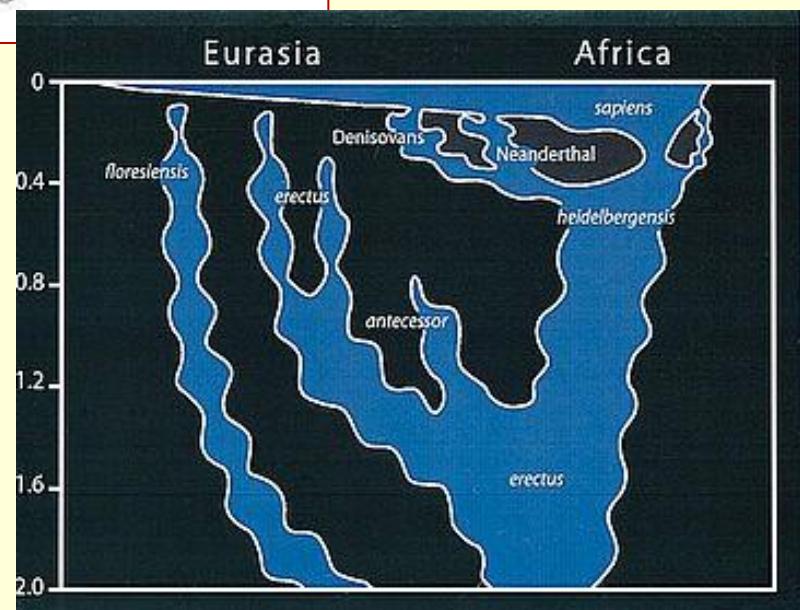


NEW MEMBER OF THE HUMAN FAMILY



Multiple hybridizations?

Hypothesis Out of Africa + partial interbreeding



Ancestors evolve into Neanderthals and first modern humans



Neanderthals die out

Neanderthal

Common ancestor with Neanderthal

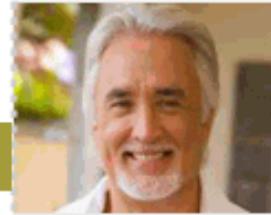
Some Neanderthal and *Homo sapiens* interbreeding

Homo sapiens



Researchers looked at five groups of modern humans

French



Han-Chinese



Papuan



Some modern humans leave Africa

Yoruba



San



Source: Science journal

Note: Time periods not to scale



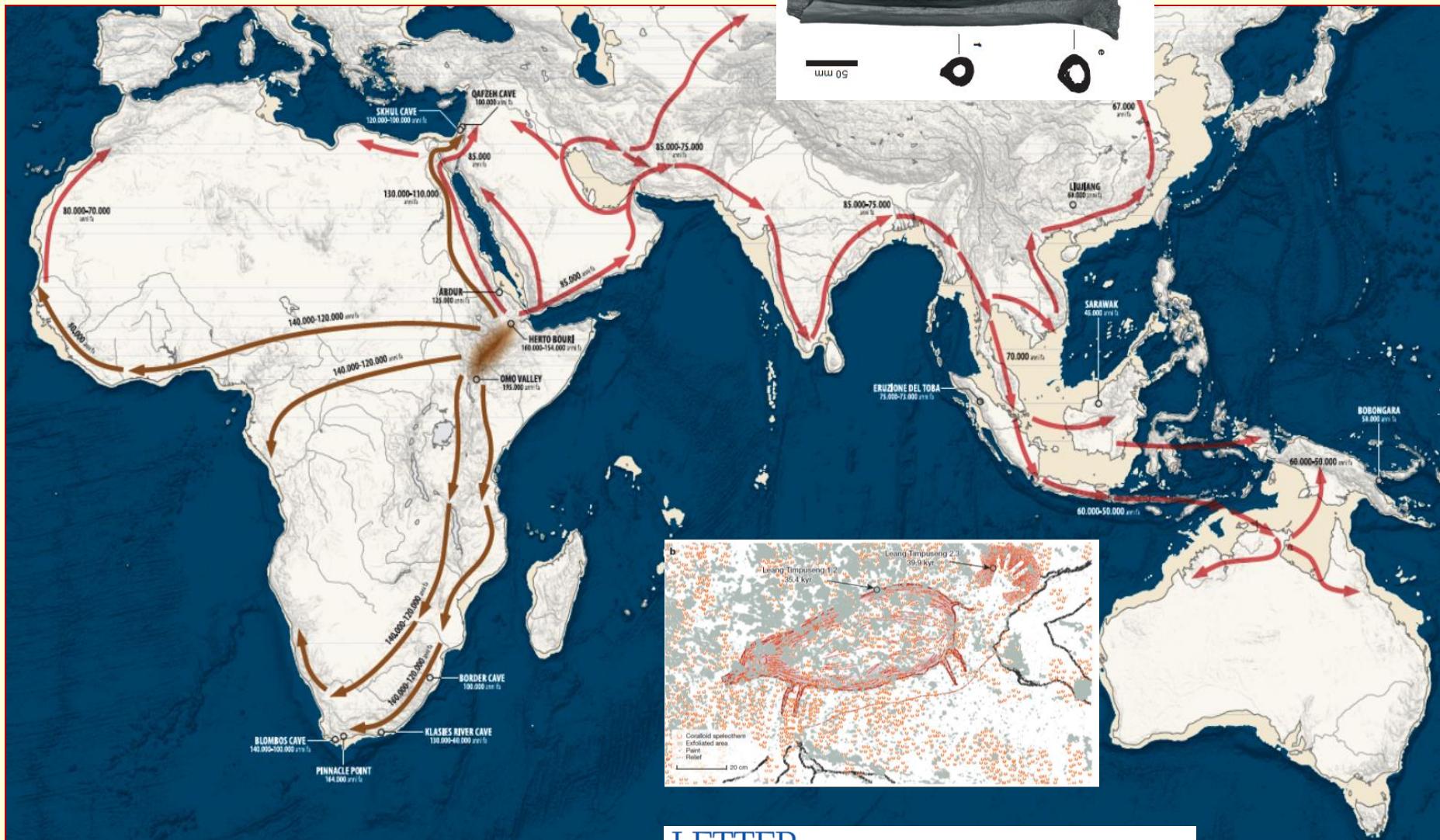
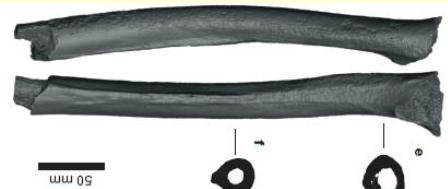
Svante Paabo

"Close Encounters of the Prehistoric Kind"

Science, May 7, 2010

"The long-awaited sequence of the Neanderthal genome suggests that modern humans and Neanderthals interbred tens of thousands of years ago, perhaps in the Middle East"

Ust-ishim femore

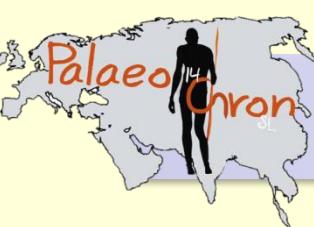


LETTER

doi:10.1038/nature13422

Pleistocene cave art from Sulawesi, Indonesia

M. Aubert^{1,2*}, A. Brumm^{1,3*}, M. Ramil³, T. Sutikna^{1,4}, E. W. Saptomo⁴, B. Hakim⁵, M. J. Morwood^{1,6}, G. D. van den Berg¹, L. Kinsley⁶ & A. Dosseto^{7,8}



The human story 100-30,000 yr ago

Neanderthals

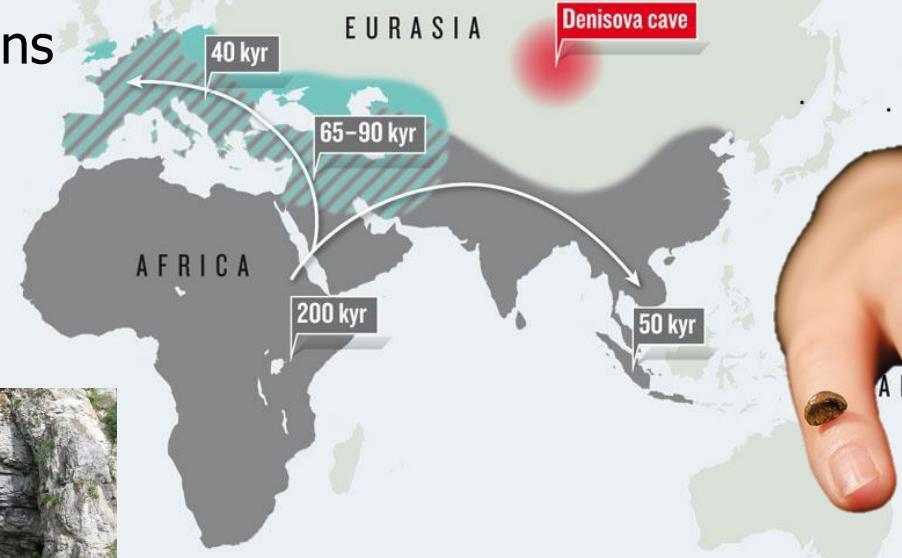
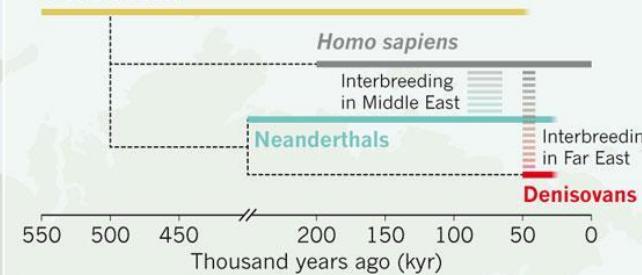
Modern humans

Denisovans

THE HUMAN STRAIN

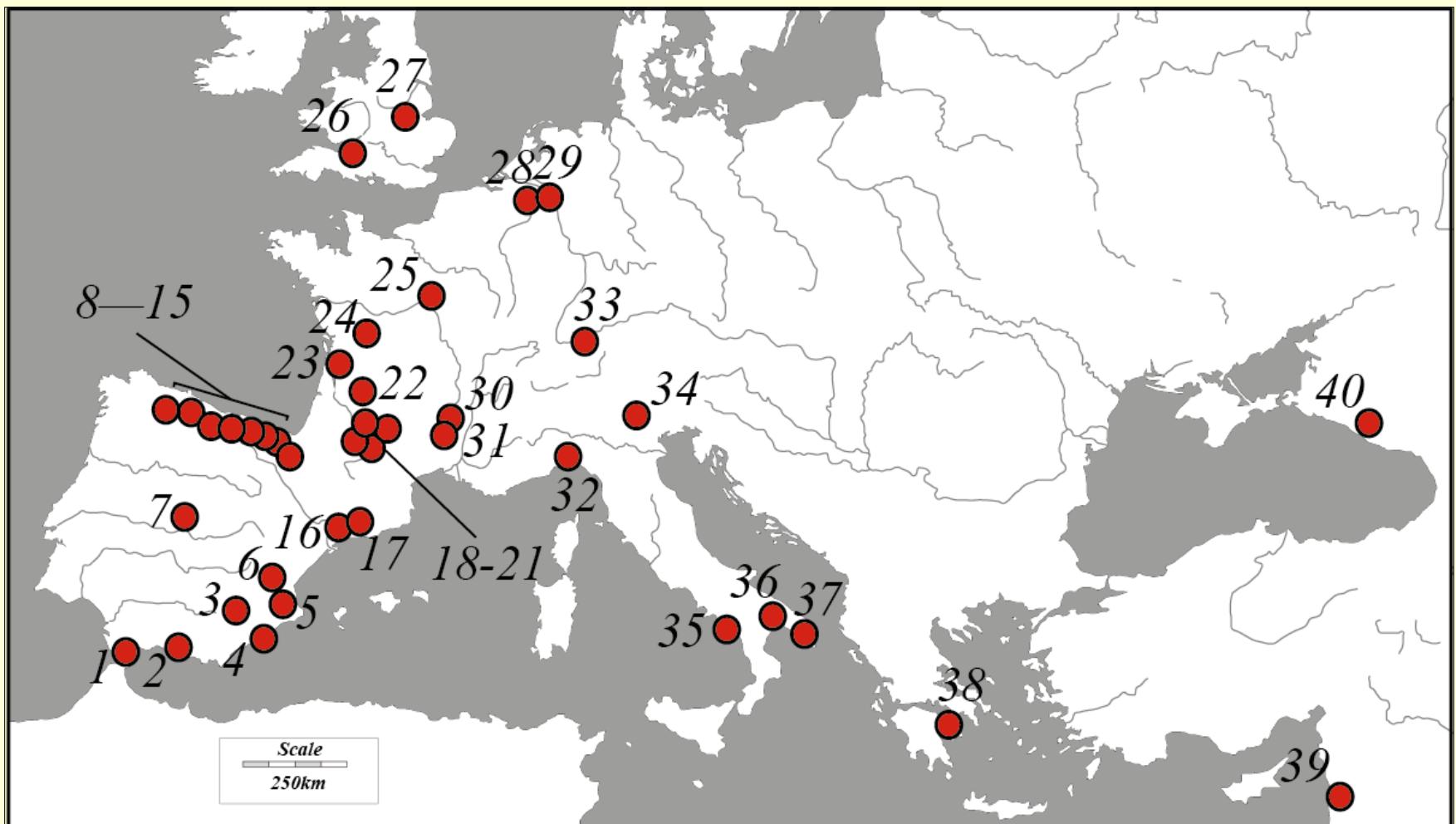
As *Homo sapiens* evolved and migrated across the world, they apparently interbred with archaic humans such as Neanderthals and Denisovans.

Homo erectus



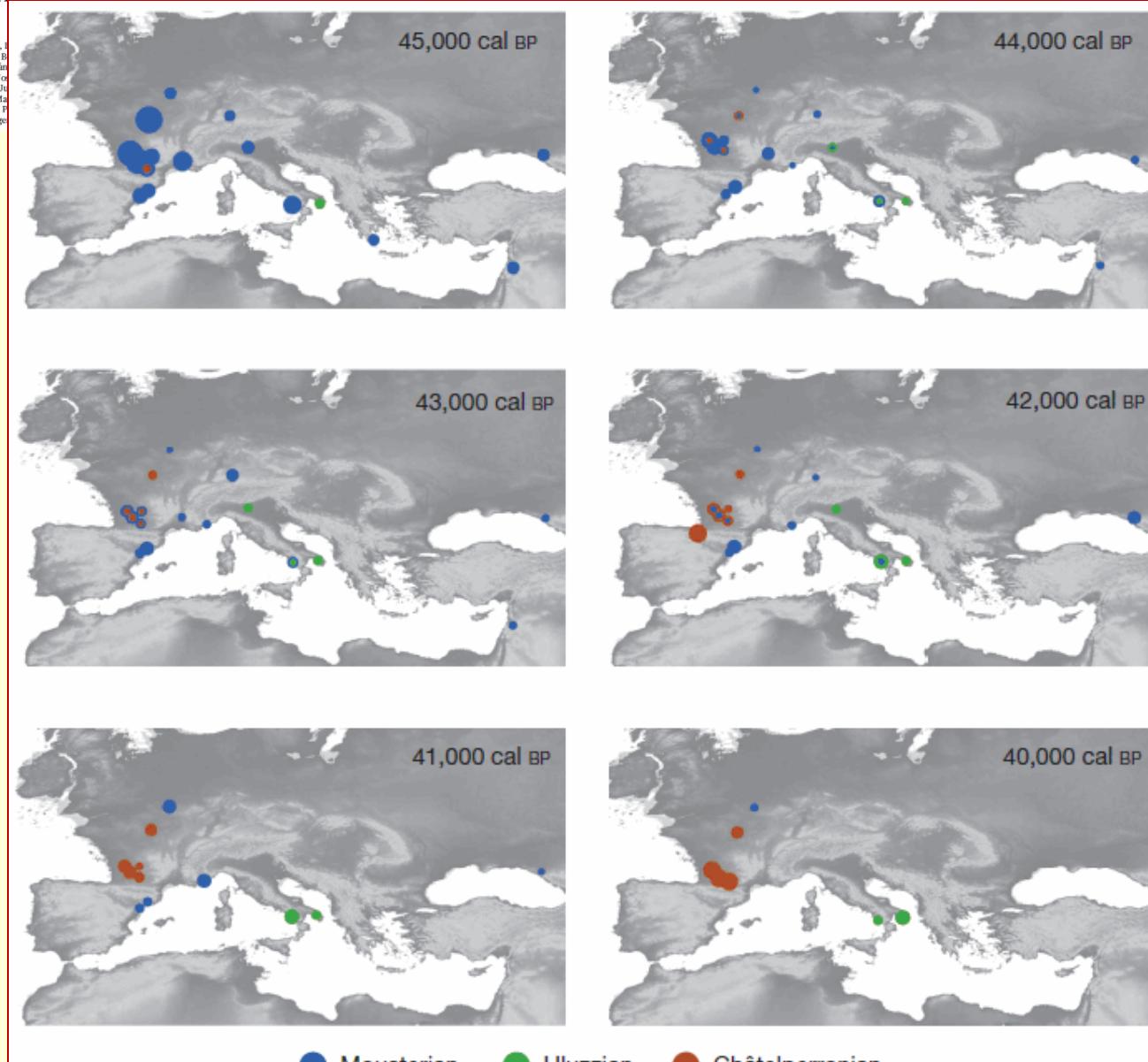
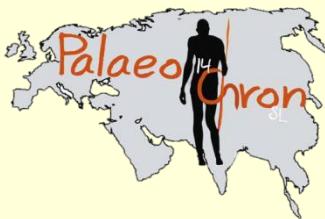
When did Neanderthals disappear?

~200 new AMS dates related to Neanderthal occupation

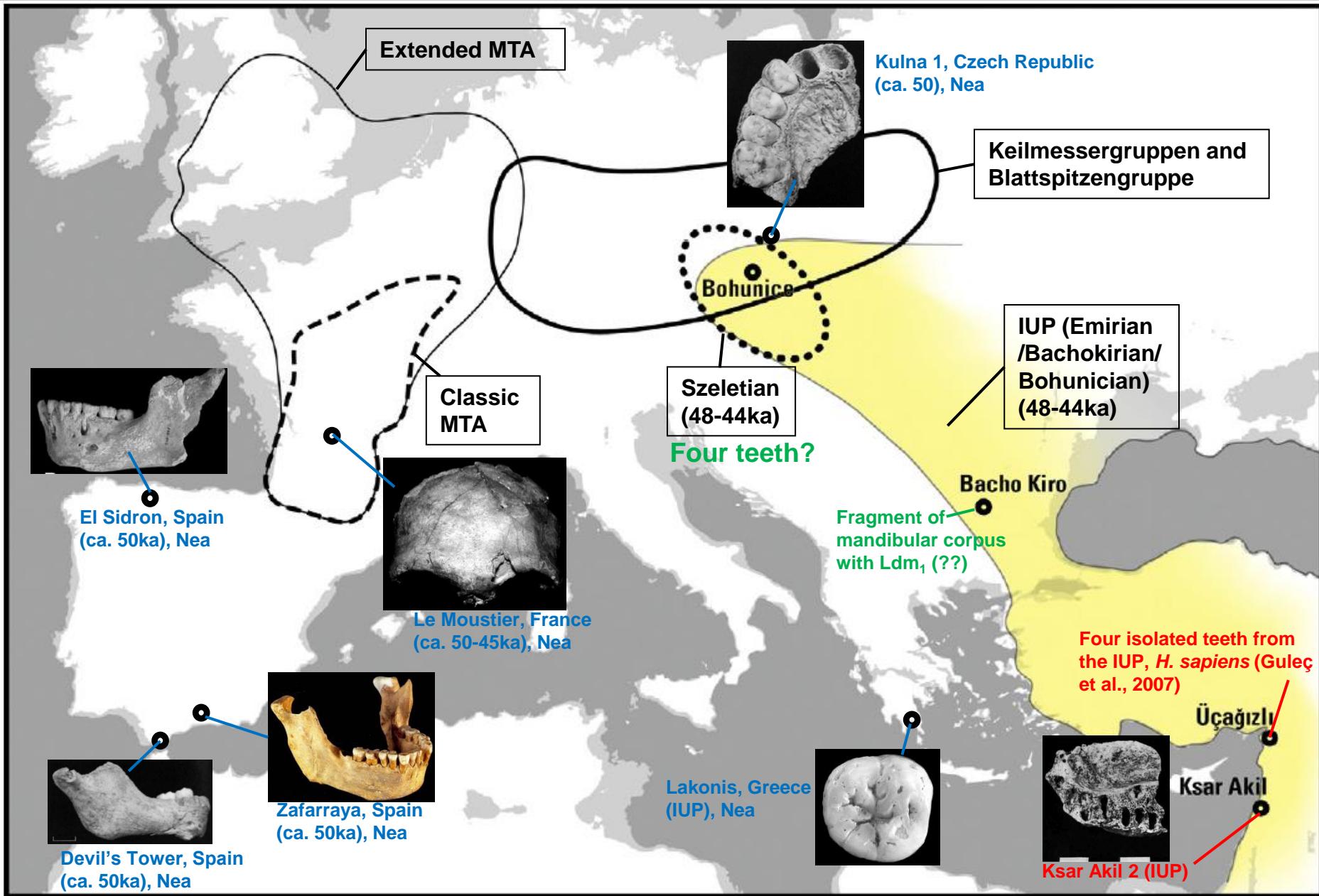


The timing and spatiotemporal patterning of Neanderthal disappearance

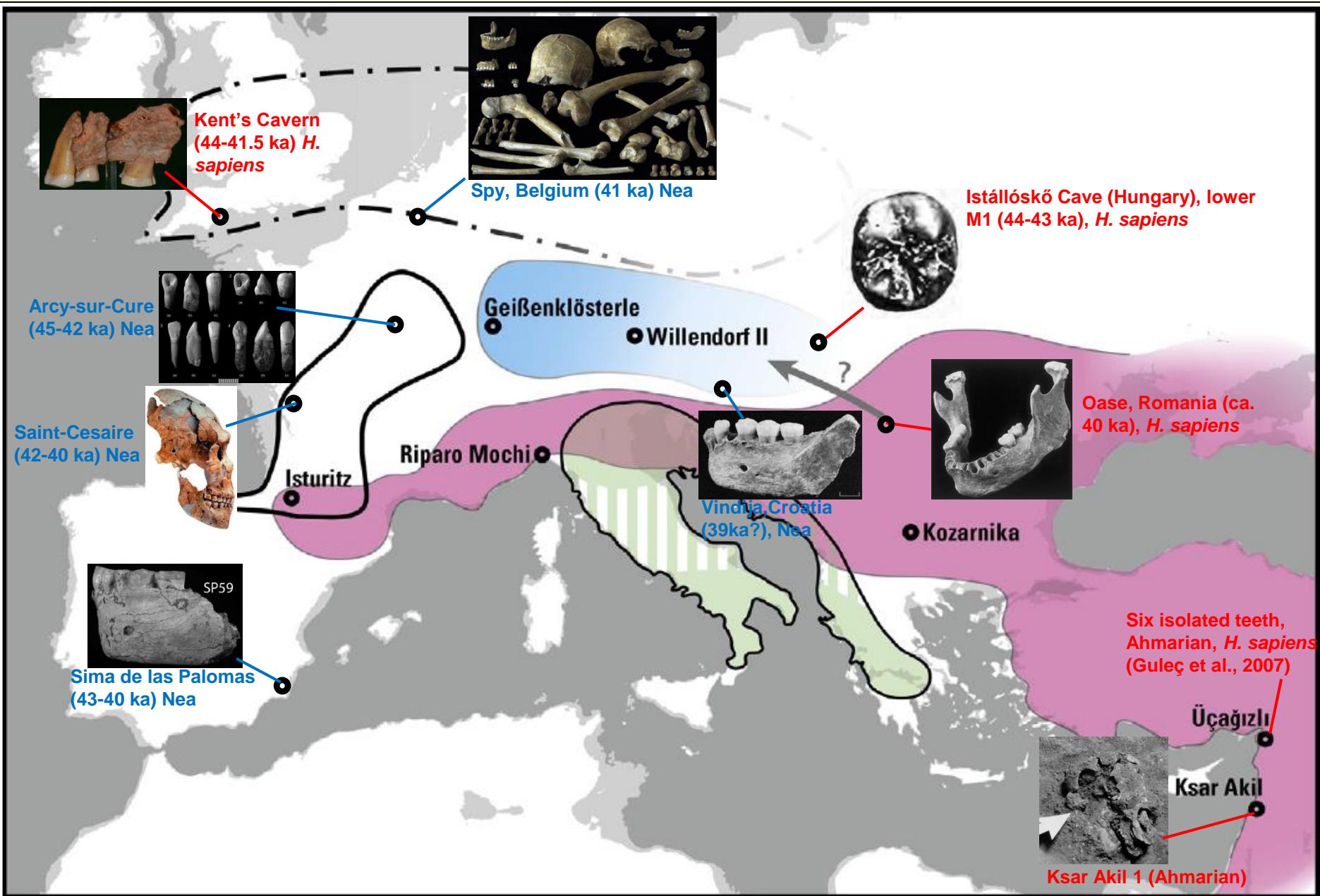
Tom Higham¹, Katherine Douka¹, Rachel Wood^{1,2}, Christopher Brook-Sargeant³, Flora Buck¹, J. Alberto Arsuaga^{4,5}, Javier Baena⁴, Cecilio Barreiro-Bailey⁶, Christopher Bergman⁶, Corinne Blundell-Jones⁷, Miguel Cárdenas¹, Nicholas J. Conard^{1,2,3}, Christelle Drull^{8,9}, Alain Froment¹⁰, Bertrand Galéra¹¹, Alejandro García-Moreno^{12,13}, Stefano Grimaldi¹⁴, Paul Haesaert¹⁵, Brigitte Holt¹⁶, María-José Ibarrola¹⁷, Arthur Jelinek²¹, Jesús F. Jordá Pardo²², José-Manuel Málaga-Fernández²³, Anar Marom^{1,23}, Júlia Martínez-Navarro²⁴, Marco de la Rasilla¹⁸, Julián Riel-Salvador²⁵, Annamaria Ronchitelli¹⁹, David Santamaría¹⁹, P. Ludovic Slimak²⁰, Joaquim Soler²¹, Narcís Soler²¹, Aritzia Valla²¹, Ron Pinhasi¹⁸ & Roger



Da 50 e 45 ka cal BP



Da 45 a 40 ka cal BP





Ancient DNA pinpoints Paleolithic liaison in Europe

Romanian fossil was the great-great-great-grandson of a Neandertal—but an evolutionary dead end



....that the Oase man had far more Neandertal DNA—composing 4.8% to 11.3% of his genome—than either the ancient modern humans from Russia or living Europeans and Asians...

An early modern human from the Peștera cu Oase, Romania

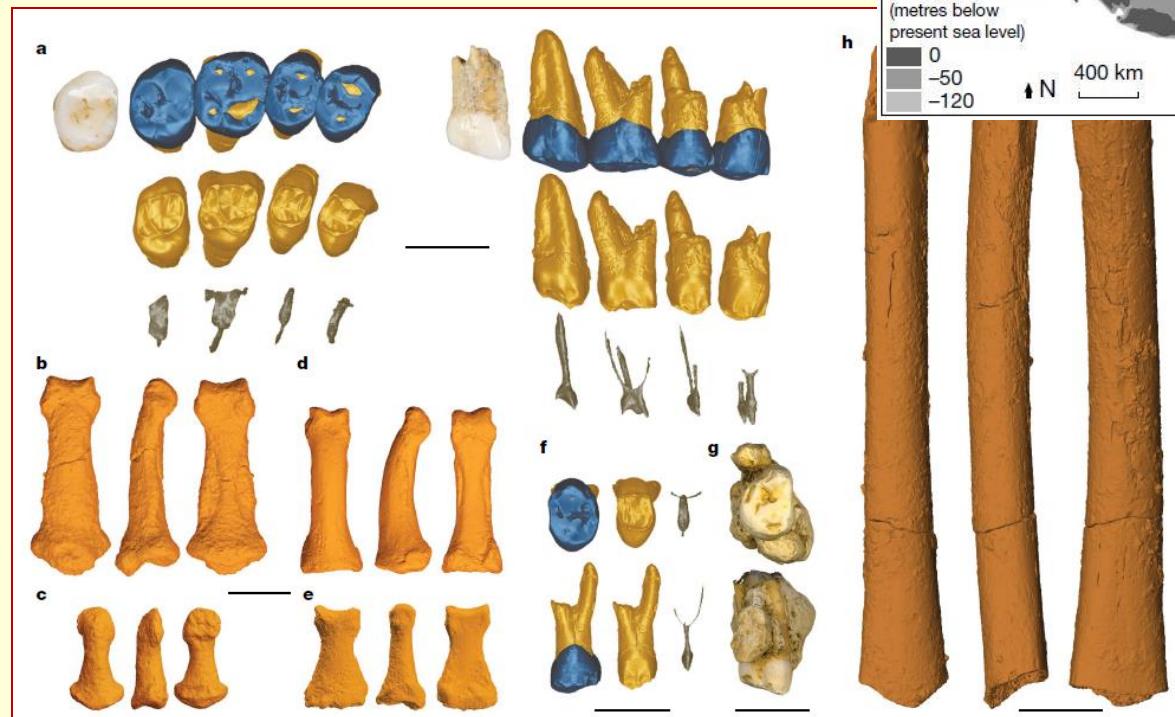
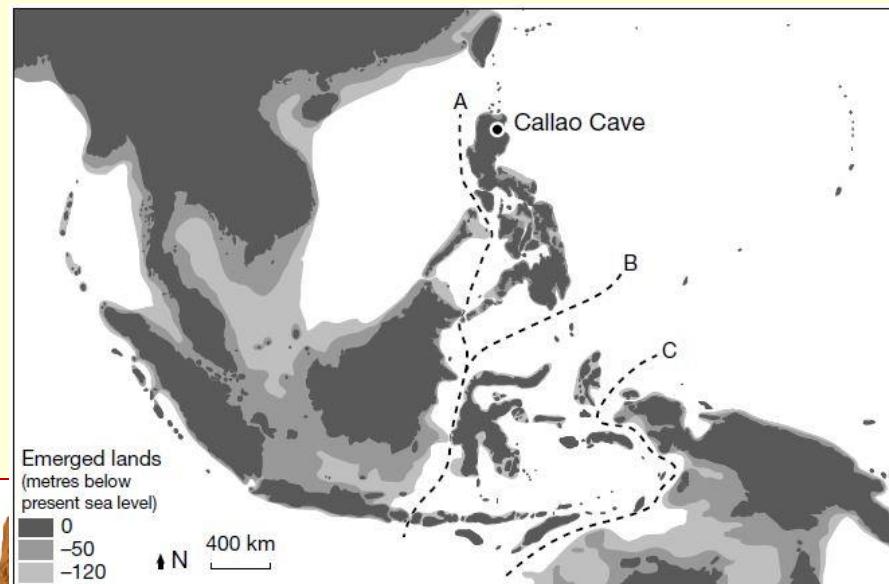
Erik Trinkaus^{*†}, Oana Moldovan[‡], Ștefan Milota[§], Adrian Bilgär[¶], Laurențiu Sarcina[§], Sheela Athreya^{||}, Shara E. Bailey^{**}, Ricardo Rodrigo^{††}, Gherase Mircea[§], Thomas Higham^{††}, Christopher Bronk Ramsey^{††}, and Johannes van der Plicht^{§§}

Popolamento dell'Australia



A new species of *Homo* from the Late Pleistocene of the Philippines

Florent Détroit^{1,8}, Armand Salvador Mijares^{2,3*}, Julien Corny¹, Guillaume Daver⁴, Clément Zanolli^{5,6}, Eusebio Dizon³, Emil Robles², Rainer Grün^{7,8} & Philip J. Piper^{3,9}



Homo floresiensis (Flores)



100,000-60,000 ka

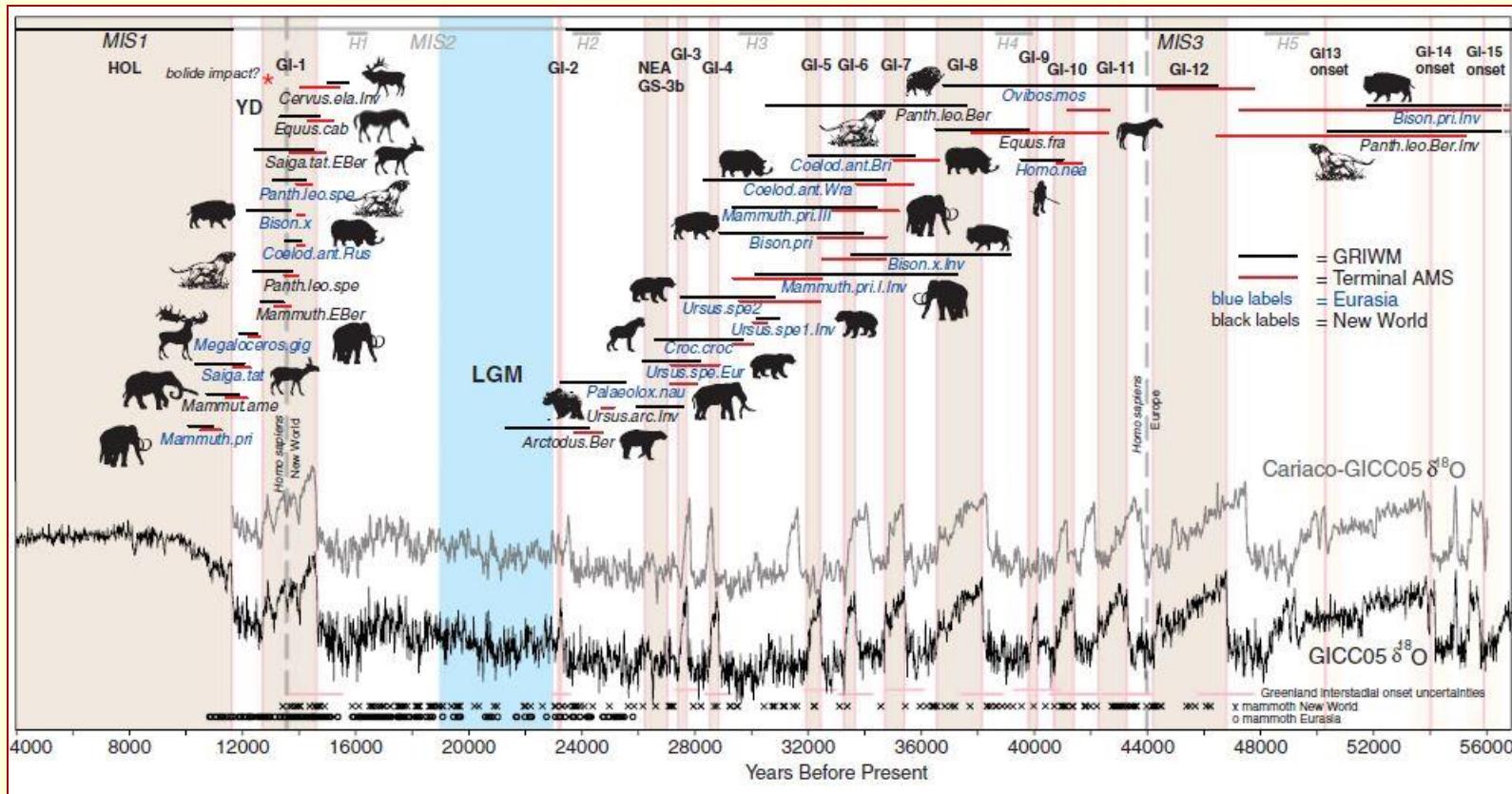
Liang Bua

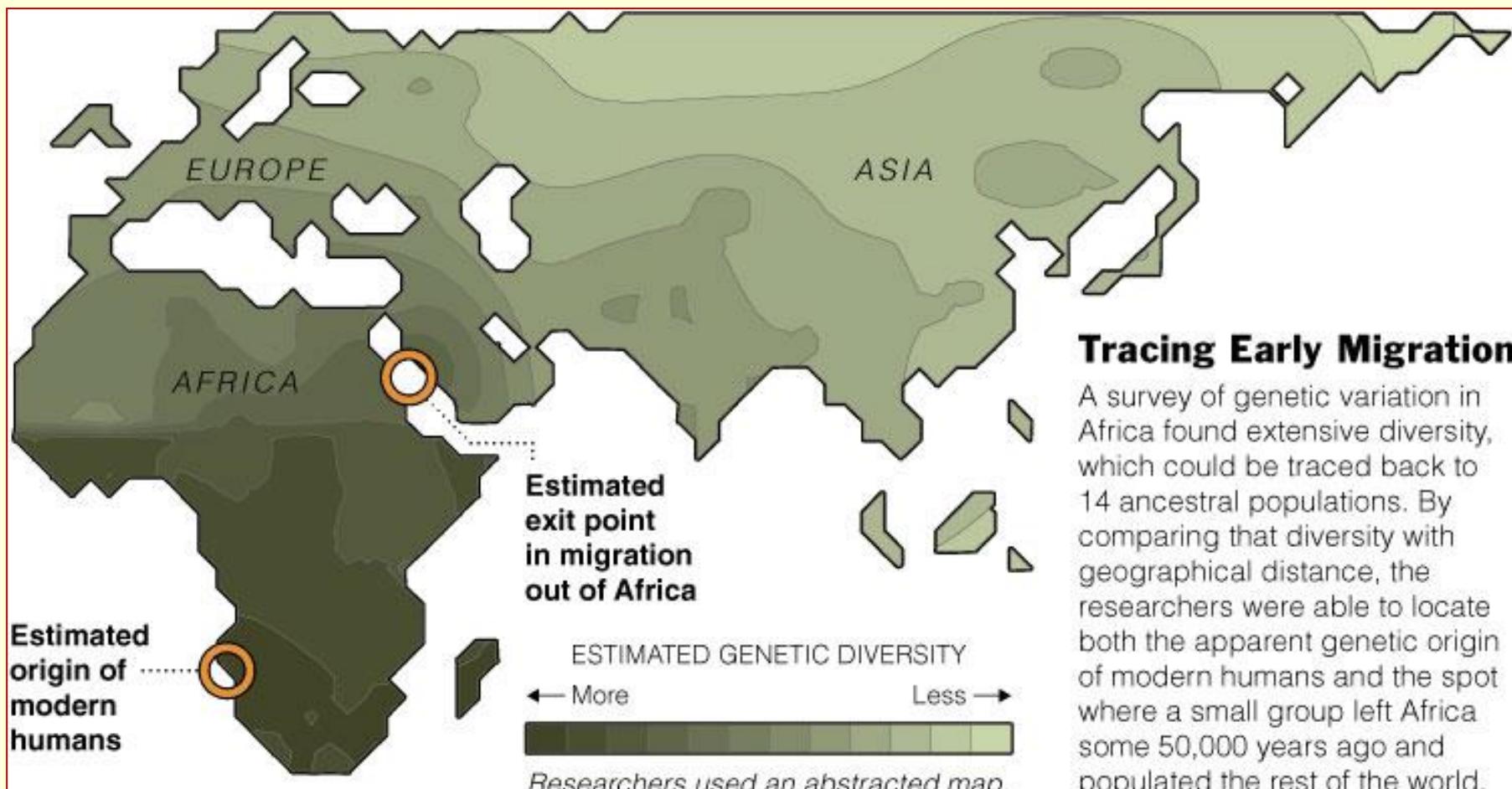


Abrupt warming events drove Late Pleistocene Holarctic megafaunal turnover

Alan Cooper,^{1*} Chris Turney,^{2*} Konrad A. Hughen,³ Barry W. Brook,^{4,5}
H. Gregory McDonald,⁶ Corey J. A. Bradshaw⁴

Eventi di estinzione-trasformazione della megaflora e records paleoclimatici del Pleistocene superiore





Sources: Science; Sarah A. Tishkoff, et al.

THE NEW YORK TIMES

Variabilità genetica popolazione africana

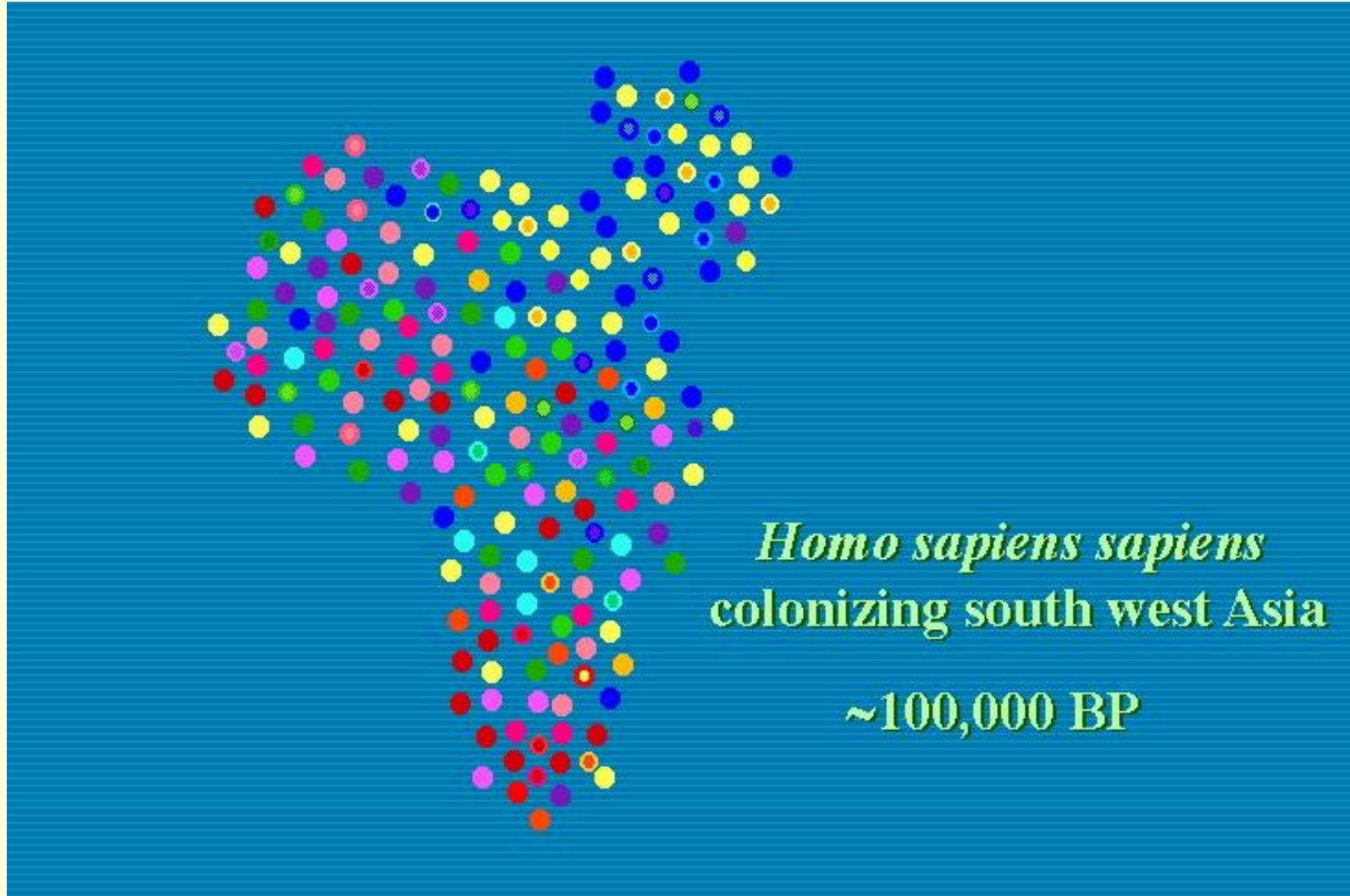
*Early *Homo sapiens sapiens*
in Africa*

150,000 to 100,000 BP



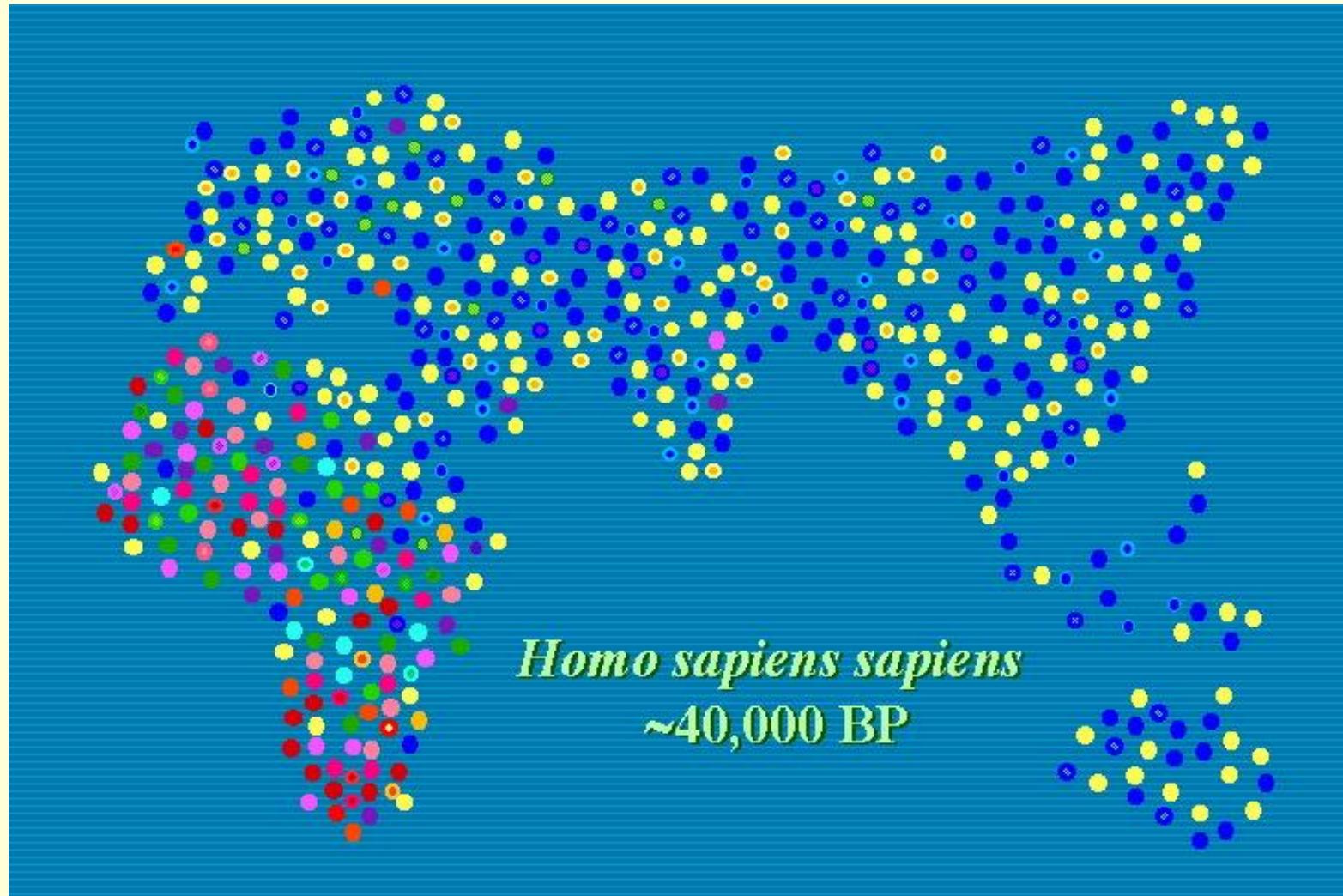
(courtesy, Kenneth Kidd, Yale University)

Deriva genetica delle prime AMH popolazioni out of Africa



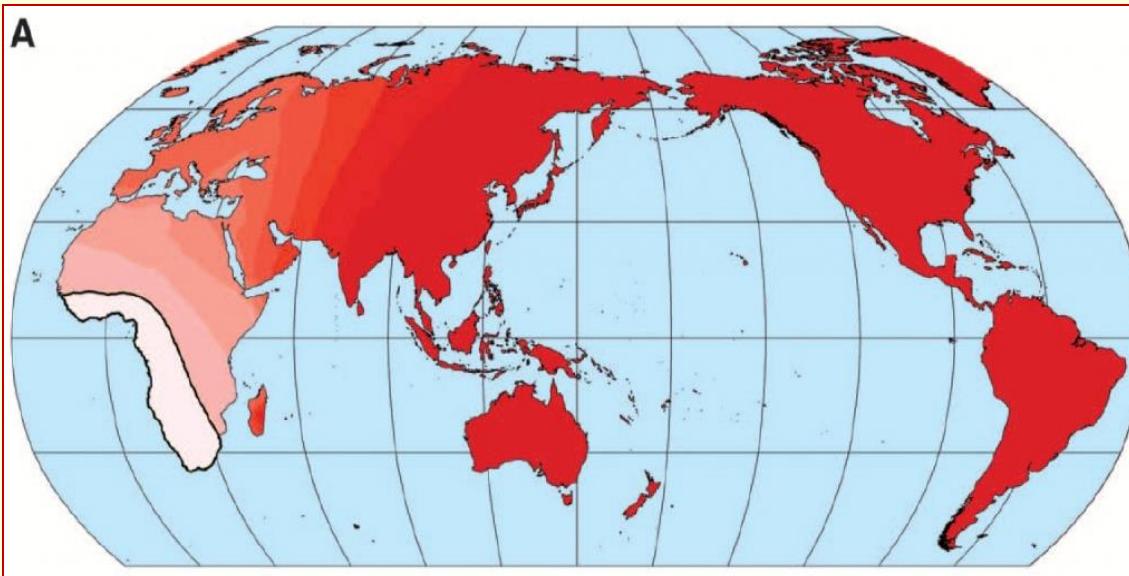
(courtesy, Kenneth Kidd, Yale University)

Riduzione della variabilità genetica della popolazione euroasiatica



(courtesy, Kenneth Kidd, Yale University)

Una relazione tra linguaggio ed espansione?



Atkinson, Science 2011

"Truly modern language, akin to languages spoken today, may thus have been the key cultural innovation that allowed the emergence of these and other hallmarks of behavioral modernity and ultimately led to our colonization of the globe"

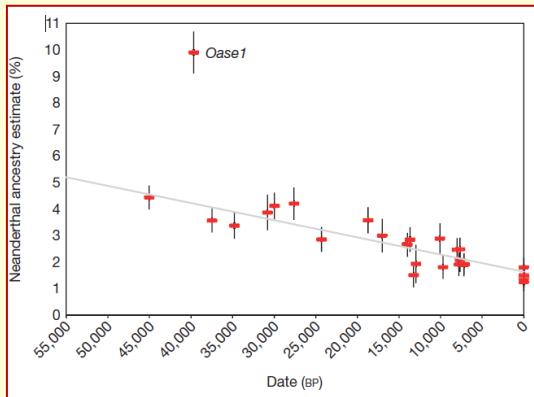
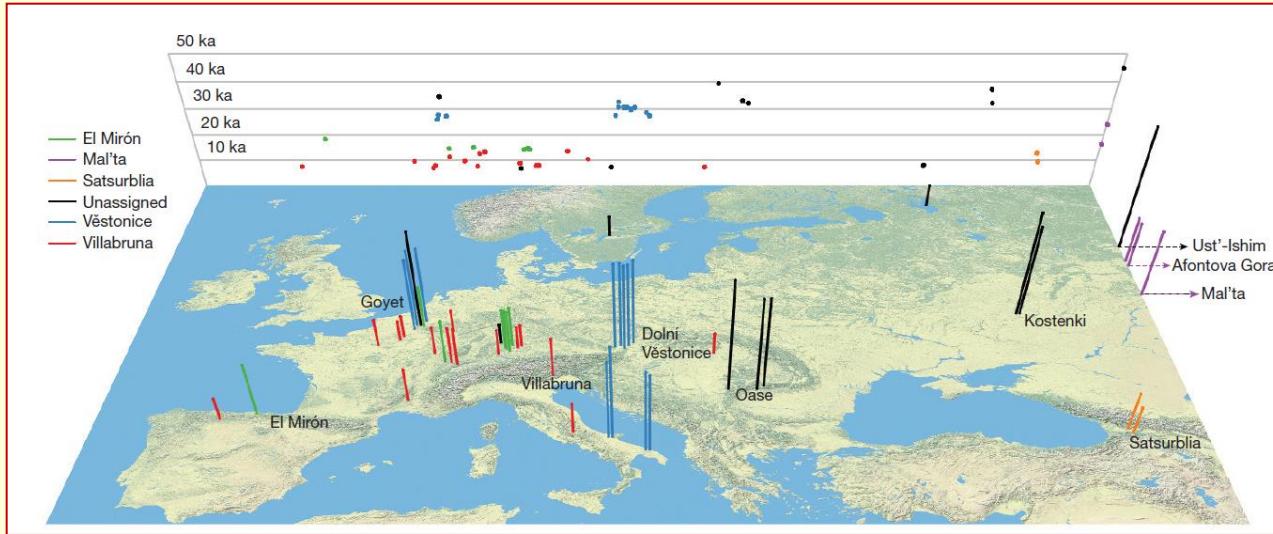
"Language was central to human expansion across the globe. It was our secret weapon, and as soon we got language we became a really dangerous species"

(Mark Pagel, NYT, April 14, 2011)

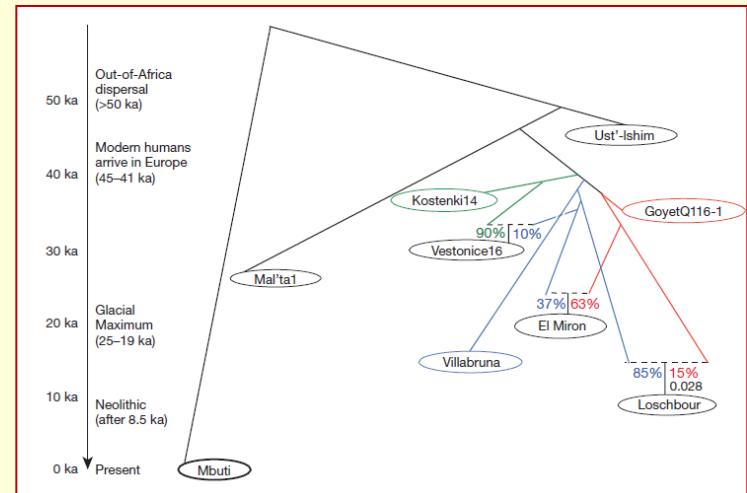
Not only one sapiens..

The genetic history of Ice Age Europe

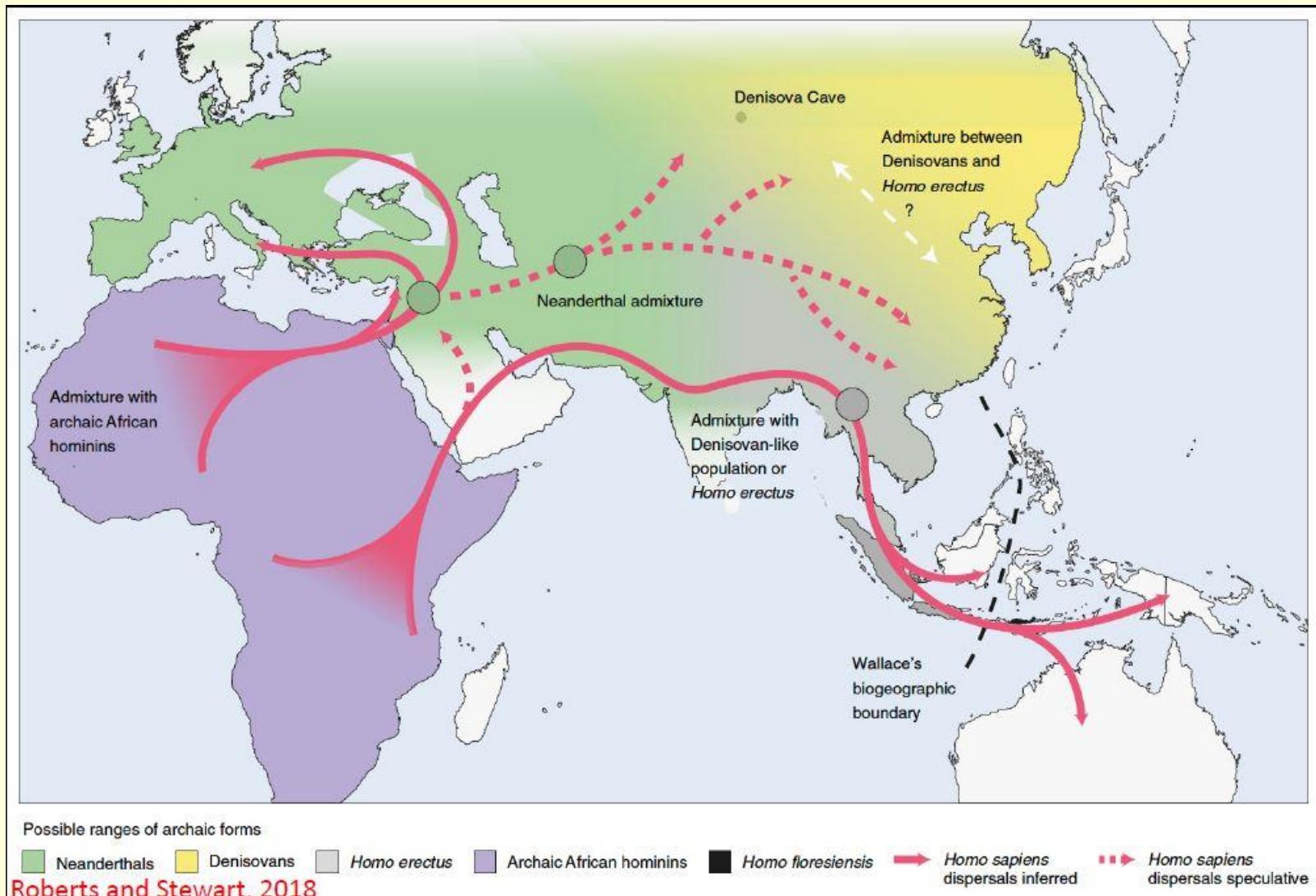
Qiaomei Fu^{1,2,3}, Cosimo Posth^{4,5*}, Mateja Hajdinjak^{3*}, Martin Petr³, Swapna Mallick^{2,6,7}, Daniel Fernandes^{8,9},



Decreasing Neandertal DNA in
Palaeolithic euroasiatic humanity



Sapiens e gli altri

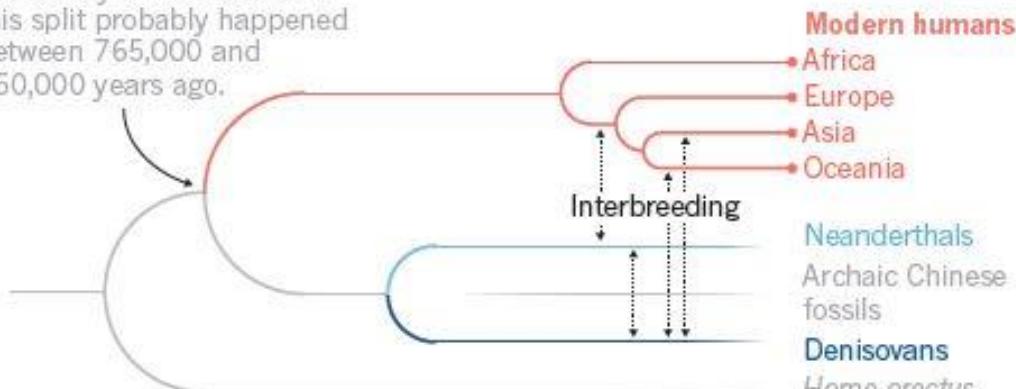




Ghosts in the cave

A mysterious group of ancient humans known as Denisovans is helping to rewrite our understanding of human evolution. Who were they?

DNA analysis indicates that this split probably happened between 765,000 and 550,000 years ago.



Current Biology

The Combined Landscape of Denisovan and Neanderthal Ancestry in Present-Day Humans

Highlights

- Denisovan admixture into modern humans occurred after Neanderthal admixture
- There is more Denisovan ancestry in South Asians than expected from current models
- Denisovan ancestry has been subject to positive and negative selection after admixture
- Male infertility most likely occurred after modern human interbreeding with Denisovans

Authors

Sriram Sankararaman,
Swapan Mallick, Nick Patterson,
David Reich

Correspondence

sriram@cs.ucla.edu (S.S.),
reich@genetics.med.harvard.edu (D.R.)

In Brief

Sankararaman et al. present a map of

Report

