









Modern Human Origins

Hugo Reyes-Centeno, Yonatan Sahle, Christian Bentz







https://moodle02.zdv.uni-tuebingen.de/course/view.php?id=1932









- ☐ Species concepts and definitions
 How do we define a species and
 how does this relate to modern
 humans?
- ☐ Genomics of modern human origins
 What does (ancient) DNA tell us
 about the origins and evolution of
 modern humans?
- Models of anthropogeny
 What model of modern human
 origins is best supported with the
 current fossil and genomic
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- ☐ The serial founder effect

 What are the genetic signatures of the human expansion out of Africa?



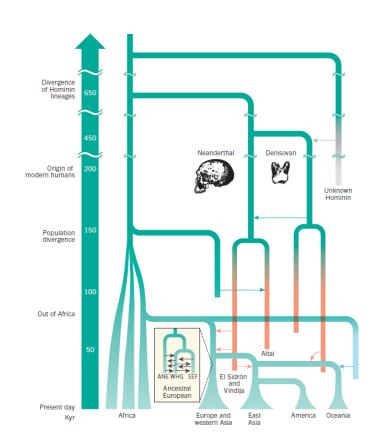




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Nielsen et al 2017

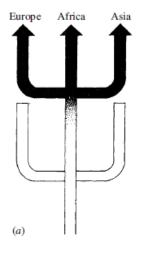


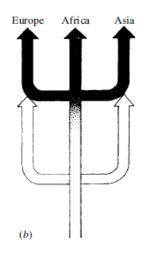


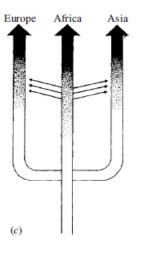


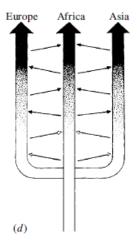
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Stringer 2001







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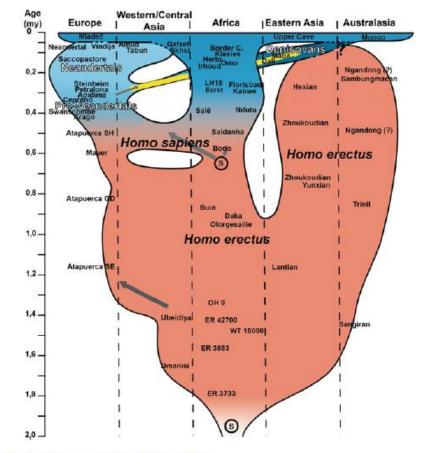


Fig. 7 Origin and evolution of Homo sapiens

Bräuer 2015



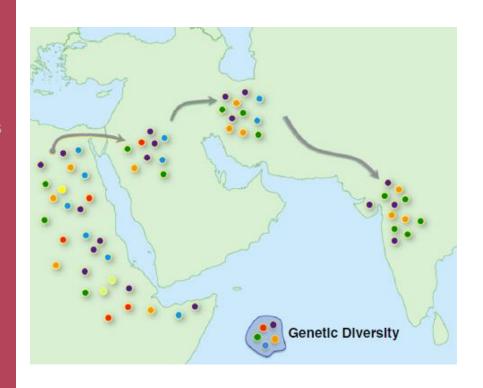




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Henn et al. 2012







This lecture:

- Modern human dispersals
 When, how, and why did
 anatomically modern humans
 - anatomically modern humans disperse out of Africa?
- Neutral and adaptive evolution
 What is the difference between
 evolution by chance and evolution
 under selection?
- ☐ Co-evolution

 Do the genotype and phenotype co-evolve?







Modern human dispersals

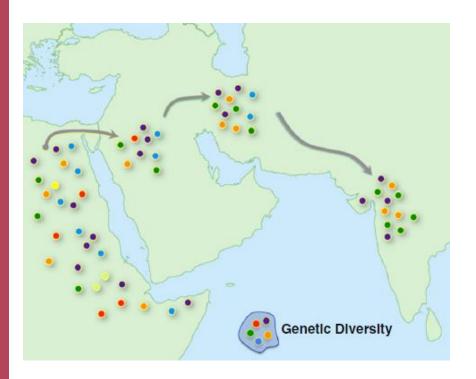
When, how, and why did anatomically modern humans disperse out of Africa?







- ☐ Timing:
 - ~130 ka
 - ~50 ka
- Mode: Number of dispersals
 - 1 Out-of-Africa event
 - 2 Out-of-Africa events
- Mode: Route of dispersal
 - Southern (to Arabian Peninsula)
 - Northern (to Levant)



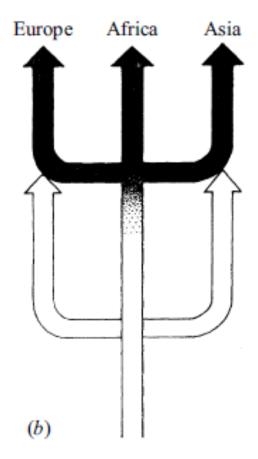
Henn et al. 2012







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- ☐ Classical Out-of-Africa and replacemnet view: Out-of-Africa at ~50ka and replacement of other hominins (e.g. Neanderthals in Europe and *erectus* in Asia)



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 - Skhul and Qazeh represent anatomically modern humans (AMH) outside of Africa between ~80-130 ka

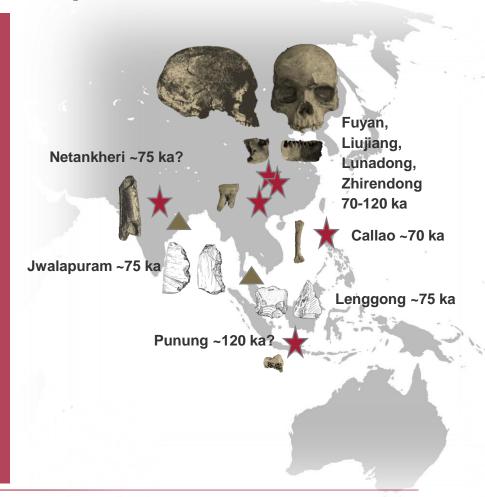








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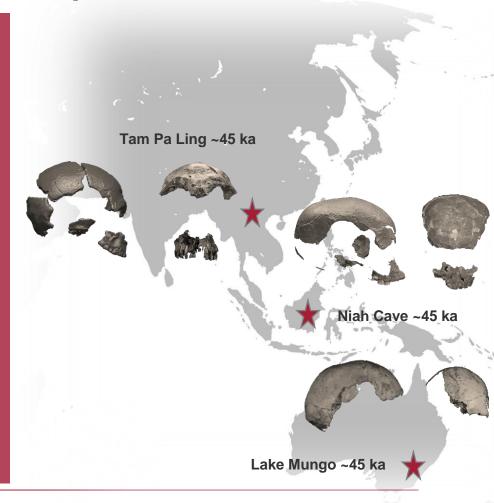








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 - Occupation of Eurasia >45 ka









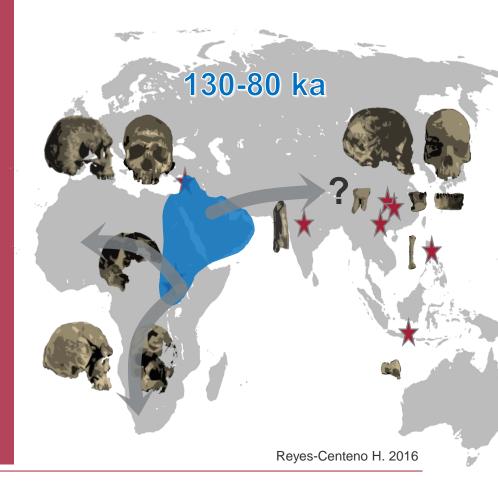
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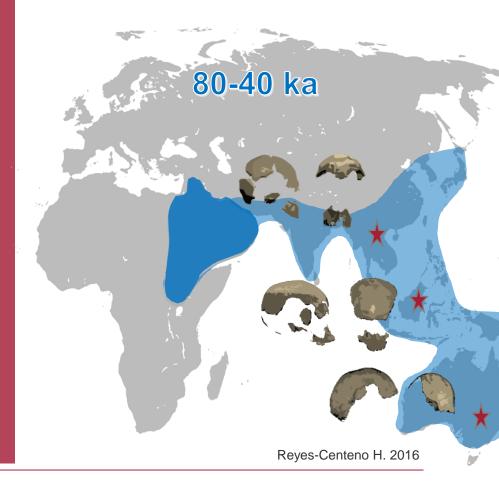








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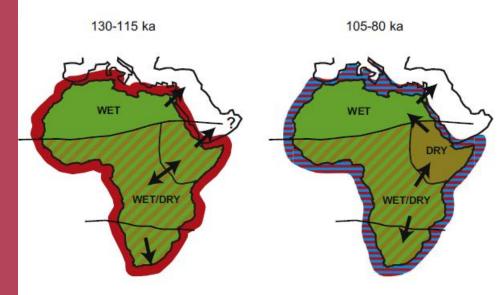








- Mode: Route of dispersal
 - Southern (to Arabian Peninsula)
 - Paleoenvironment: passage more likely between 145-115 ka and again between 80-65
 - Northern (to Levant)
 - Paleoenvironment: passage more likely between 140-75 ka



Blome et al. 2012

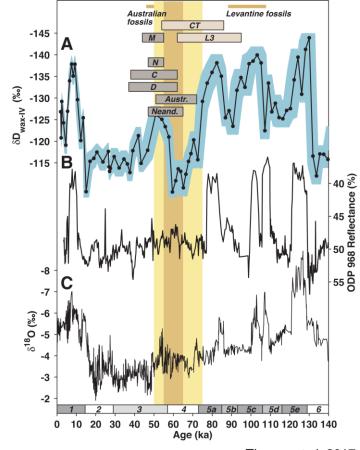






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Tierney et al. 2017

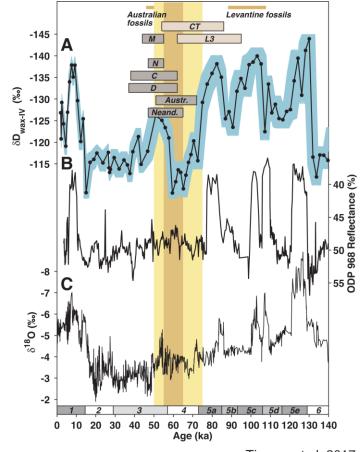






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 - Why disperse during arid conditions time period?
 - Lag in genomic estimates?









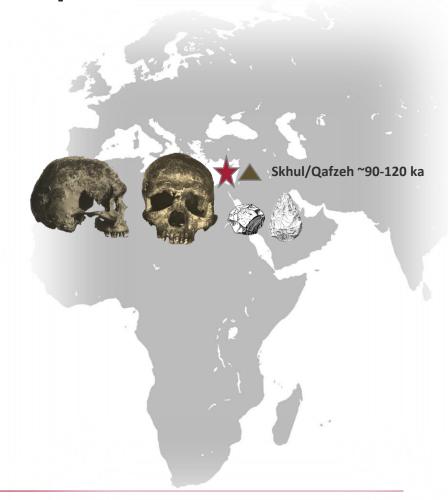
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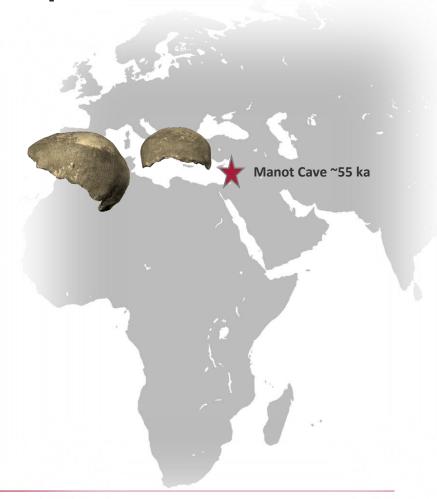








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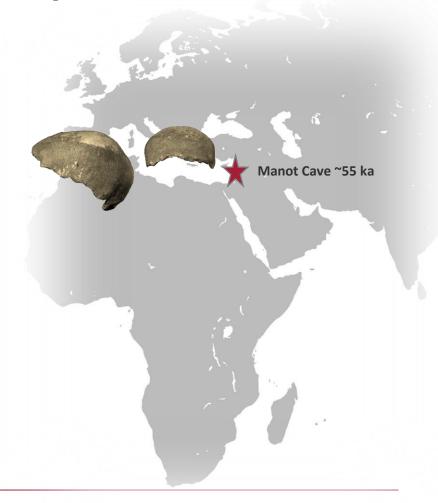








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 - No fossils yet reported in southern Arabian Peninsula



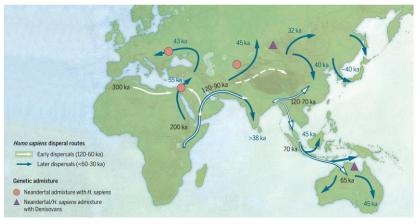


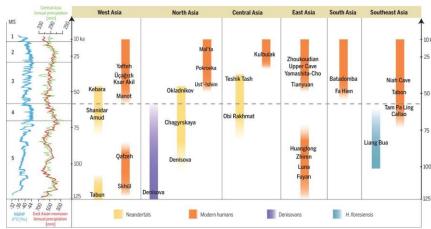




☐ Revised view:

- Expansion ~130 ka to the Levant and possibly further into Eurasia, followed by extinction
 - Small populations?
 - Contributions to Neanderthals?
 - Competition with other hominins?
 - Middle Paleolithic / MSA toolkit?
- Major dispersal between ~80-50 ka





Bae et al 2017







Neutral & adaptive evolution

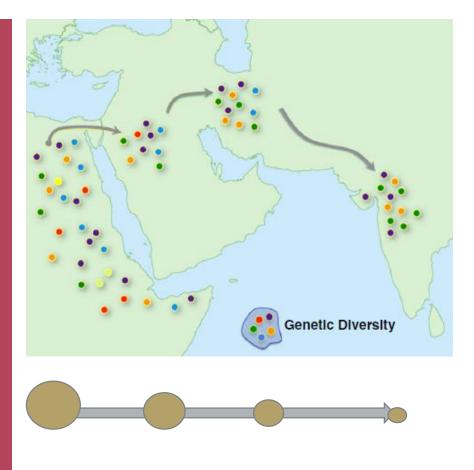
What is the difference between evolution by chance and evolution under selection?







- ☐ Serial founding effect (or cascading bottlenecks)
 - Decreasing intra-population diversity with geographical distance from Africa



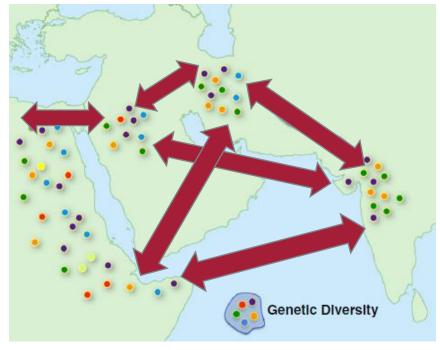
Henn et al. 2012

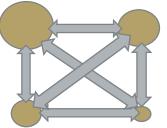






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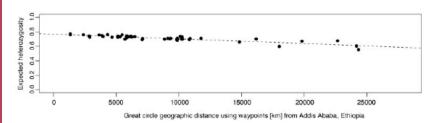


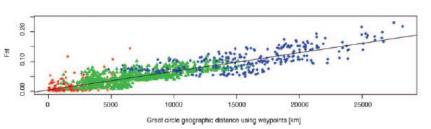






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Ramachandran et al. 2005

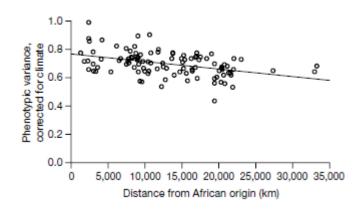


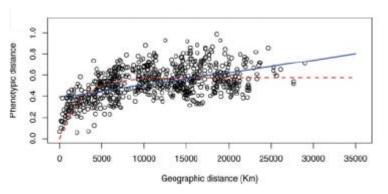




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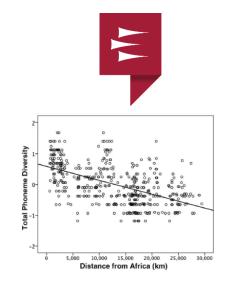
Manica et al. 2007; Betti et al. 2011

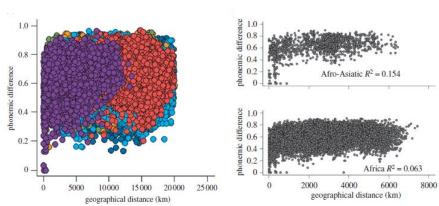






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Atkinson 2011; Hunley et al. 2012

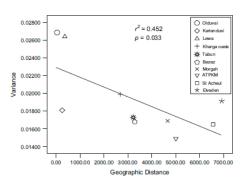






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Lycett & von Cramon-Taubadel 2008

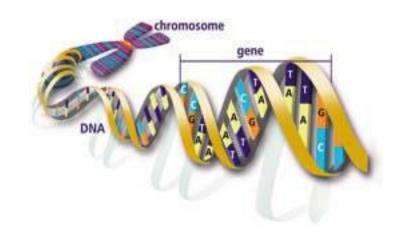






Modes of evolution

■ Neutral evolution (or drift) refers to changes in the genotype and phenotype that refer to chance events or <u>stochastic processes</u> (e.g. random mutations)



US Department of Energy

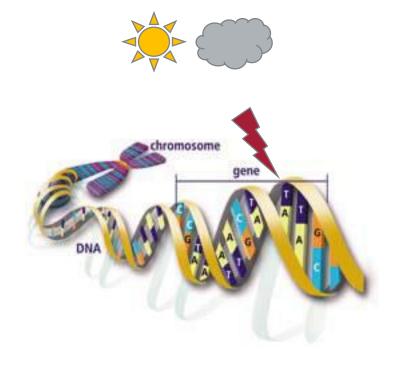






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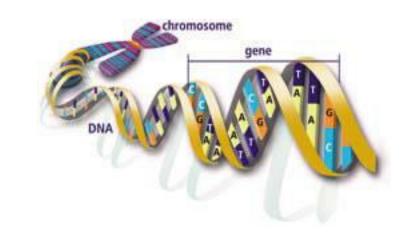




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Note: changes in genotype and phenotype can occur during an organisms lifetime, but they will not be inherited to the next generation unless they occur along the germ line or an epigenetic process.



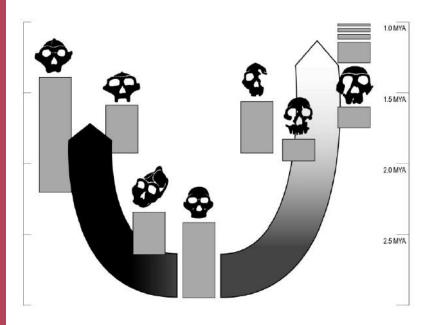
US Department of Energy







Adaptive evolution is thought to have played an important role in early human evolution



Rogers Ackermann & Cheverud 2004







- Adaptive evolution is thought to have played an important role in early human evolution
- Netural evolution is thought to play the primary role in recent human evolution



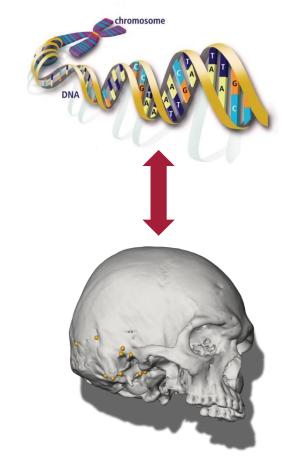
Weaver et al 2007







- Adaptive evolution is thought to have played an important role in early human evolution
- Netural evolution is thought to play the primary role in recent human evolution
 - Direct comparison of netural genomic and phenotypic variation between populations is significant



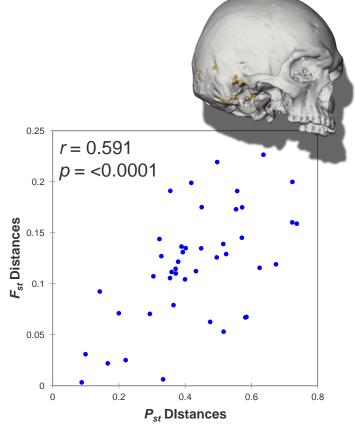
US Department of Energy; Reyes-Centeno et al 2017







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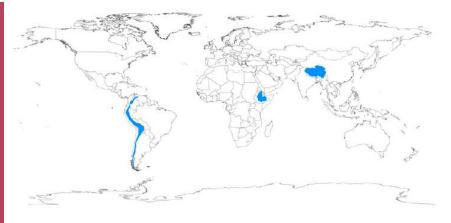
Reyes-Centeno et al 2017

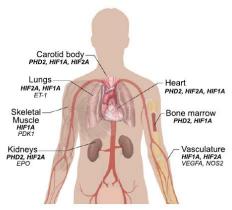






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- ☐ Adaptation still plays an important role
 - e.g. adaptation to high altitude environments: independent (convergent) evolution and possible hominin introgression in Tibet





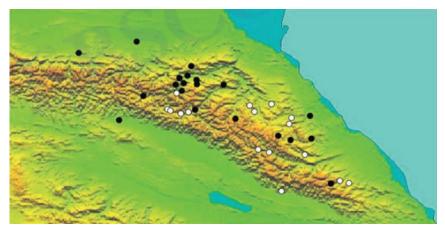
Bingham & Lee 2014







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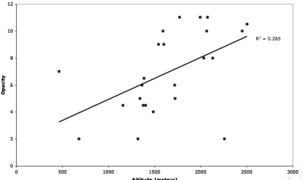


Fig. 2. Altitude (x-axis) by opacity (y-axis). Number of languages: N = 26. Shared variance: $R^2 = 0.27$

Nichols 2013







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 - Multiple candidate genes that are associated with phenotypes

Genes	Function or phenotype
LCT, MAN2A1, SI, SLC27A4, PPARD, SLC25A20, NCOA1, LEPR, LEPR, ADAMTS19, ADAMTS20, APEH, PLAU, HDAC8, UBR1, USP26, SCP2, NKX2-2, AMY1, ADH, NPY1R, NPY5R	Digestion of milk and dairy products; metabolism of carbohydrates, starch, proteins, lipids and phosphates; alcohol metabolism
Cytochrome P450 genes (CYP3A5, CYP2E1, CYP1A2 and CYP2D6)	Detoxification of plant secondary compounds
CD58, APOBEC3F, CD72, FCRL2, TSLP, RAG1, RAG2, CD226, IGJ, TJP1, VPS37C, CSF2, CCNT2, DEFB118, STAB1, SP1, ZAP70, BIRC6, CUGBP1, DLG3, HMGCR, STS, XRN2, ATRN, G6PD, TNFSF5, HbC, HbE, HbS, Duffy, α -globin	Immunity, pathogen response; resistance to malaria and other crowd diseases
LEPR, PON1, RAPTOR, MAPK14, CD36, DSCR1, FABP2, SOD1, CETP, EGFR, NPPA, EPHX2, MAPK1, UCP3, LPA, MMRN1	Energy metabolism, hot or cold tolerance; heat-shock genes
SLC24A5, SLC25A2, EDAR, EDA2R, SLC24A4, KITLG, TYR, 6p25.3, OCA2, MC1R, MYO5A, DTNBP1, TYRP1, RAB27A, MATP, MC2R, ATRN, TRPM1, SILV, KRTAPs, DCT	The externally visible phenotype (skin pigmentation, hair thickness, eye and hair colour, and freckles)
CDK5RAP2, CENPJ, GABRA4, PSEN1, SYT1, SLC6A4, SNTG1, GRM3, GRM1, GLRA2, OR4C13, OR2B6, RAPSN, ASPM, RNT1, SV2B, SKP1A, DAB1, APPBP2, APBA2, PCDH15, PHACTR1, ALG10, PREP, GPM6A, DGKI, ASPM, MCPH1, FOXP2	Nervous system, brain function and development; language skills and vocal learning
BMP3, BMPR2, BMP5, GDF5	Skeletal development
MYH16, ENAM	Jaw muscle fibres; tooth-enamel thickness

Laland et al 2010







Co-evolution

Do the genotype and phenotype co-evolve?







 Darwin's hypothesized the co-evolution of languages and genes

"If we possessed a perfect pedigree of mankind, a genealogical arrangement of the races of man would afford the best classification of the various languages now spoken throughout the world...."



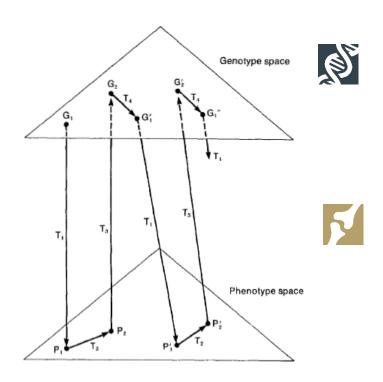
Charles Darwin, 1859







- Darwin's hypothesized the co-evolution of languages and genes
- Theoretical framework: one can trace how the genotype and phenotype affect each other
 - Selection acts on phenotypes and gene frequency changes as a consequence



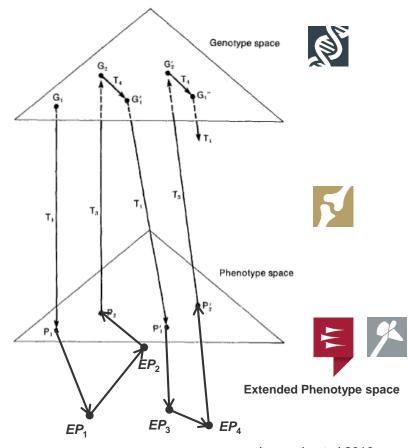
Lewontin et al 2010







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- Darwin's hypothesized the co-evolution of languages and genes
- Theoretical framework: one can trace how the genotype and phenotype affect each other
 - Selection acts on phenotypes and gene frequency changes as a consequence
 - Language and behavior can be considered as an extended phenotype
 - Language and behavior tend to change much faster than the genotype

"Organic evolution is not the only sort of evolution in the sense of a process of cumulative change. When making level of intelligence symbolic speech possible was reached in the anthropoid line, a new evolutionary process emerged, enormously more rapid than organic evolution."

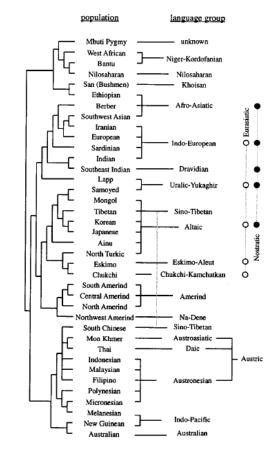
Sewall Wright, 1950







- Darwin's hypothesized the co-evolution of languages and genes
 - Language and gene trees sometimes mirror each other, but other times do not
 - At the level of language families, there is a high correlation but not always at lower language levels

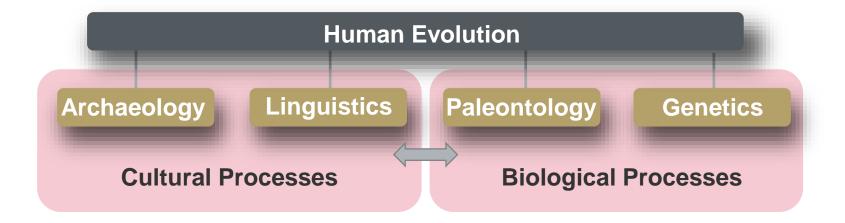


Cavalli-Sforza 1997





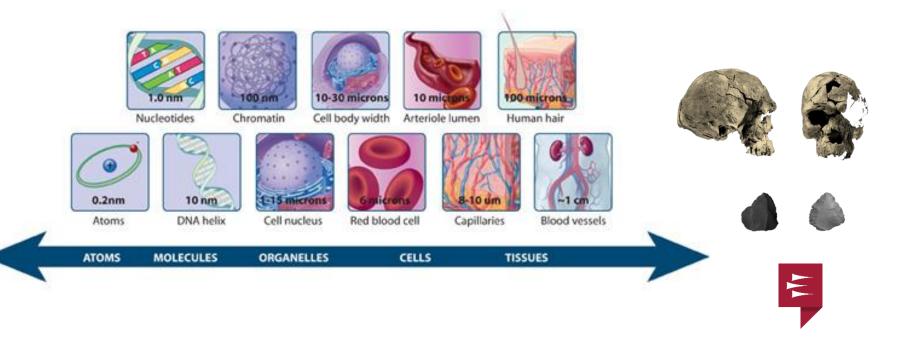












Nature Education 2010; White et al. 2003; Beyin 2013







