



AA

Indexing View Papers Aims & Scope Editorial Board Guideline Article Processing Charges

AA > Vol.2 No.4, November 2012

OPEN ACCESS

Re-Examining the Out-of-Africa Theory and the Origin of Europeoids (Caucasoids). Part 2. SNPs, Haplogroups and Haplotypes in the Y Chromosome of Chimpanzee and Humans

PDF (Size:596KB) PP. 198-213 DOI: 10.4236/aa.2012.24022

Author(s)

Anatole A. Klyosov, Igor L. Rozhanskii, Lyudmila E. Ryabchenko

ABSTRACT

Our consideration of human haplogroups, and our analysis of the dynamics of the Y-chromosome nucleotide flow from primates to humans during the evolution of genus *Homo* has shown that a common ancestor of the majority of present day human males, both African and non-African, lived approximately 160,000 years ago. The haplogroup of this common ancestor has been identified as the α -haplogroup, which is equivalent or close to haplogroups A1/A1b in the current phylogeny. The archaic lineages (currently summarily designated A0) descend from an ancestor who lived no later than 180,000 years ago, and probably much earlier. The α -haplogroup and the A0 lineages have significantly different nucleotide patterns, and they certainly did not descend one from another. Furthermore, our research points up the areas of mutations in Y-chromosome in *H. sapiens*, which allows us to use chimpanzee MSY (the male-specific region of the Y-chromosome) as a proxy for genus *Homo*'s common α -haplogroup ancestor. When we studied slow mutating 16-marker haplotypes, we discovered that chimpanzees and present day humans had a common ancestor 5.5 ± 0.9 million years before the present. It is clear that, when they are compared to loci in other primates, such as gorillas, orangutans, and macaques, many human Y-chromosome loci have been conserved from our common ancestor. Results of our analysis of haplotypes, conserved (ancestral) nucleotides, and SNPs suggest that there is no reason to believe that ancestors of non-Africans (β -haplogroup, i.e. haplogroup BT and its downstream haplogroups) descended from haplogroups A0, A1a, or any other African haplogroup. The data are adequately described by a model which shows that the African lineages and non-African lineages diverged from the α -haplogroup approximately 160,000 years before the present and that the Y-chromosomes of the two groups have evolved independently (in terms of Y-chromosome) since then. We have no indication of where the common ancestor of the α -haplogroup lived; he could just as easily have lived in Europe, in Asia, or in the Middle East, as in (less likely) Africa. We believe that all the presuppositions posited in support of the Out-of-Africa hypothesis fail to hold up under simple scrutiny. This study shows that the Out-of-Africa hypothesis has not been adequately substantiated. The common assertion that "anatomically modern humans came out of Africa some 70,000 years ago" has never been convincingly calculated or determined otherwise; our research suggests that it is incorrect.

KEYWORDS

Y-Chromosome; Mutations; Haplotypes; Haplogroups; Primates; Chimpanzees; SNP; Out-of-Africa

Cite this paper

Klyosov, A. , Rozhanskii, I. & Ryabchenko, L. (2012). Re-Examining the Out-of-Africa Theory and the Origin of Europeoids (Caucasoids). Part 2. SNPs, Haplogroups and Haplotypes in the Y Chromosome of Chimpanzee and Humans. *Advances in Anthropology*, 2, 198-213. doi: 10.4236/aa.2012.24022.

References

- [1] Atkinson, Q. D. (2011). Phonemic diversity supports a serial founder effect model of language expansion from Africa. *Science*, 332, 346-349. doi:10.1126/science.1199295
- [2] Behar, D. M., Villem, R., Soodyall, H., Blue-Smith, J., Pereira, L., Metspalu, E., Scozzari, R. et al. (2008). The dawn of human matrilineal diversity. *The American Journal of Human Genetics*, 82, 1130-

• Open Special Issues

• Published Special Issues

• Special Issues Guideline

AA Subscription

Most popular papers in AA

About AA News

Frequently Asked Questions

Recommend to Peers

Recommend to Library

Contact Us

Downloads: 24,046

Visits: 113,311

Sponsors >>

- [3] Benazzi, S., Douka, K., Fornai, C., Bauer, C. C., Kullmer, O., Svoboda, J. et al. (2011). Early dispersal of modern humans in Europe and implications for Neanderthal behaviour. *Nature*, 479, 525-528. doi:10.1038/nature10617
- [4] Campbell, M. C., & Tishkoff, S. A. (2010). The evolution of human genetic and phenotypic variation in Africa. *Current Biology*, 20, R166-R173. doi:10.1016/j.cub.2009.11.050
- [5] Carrigan, D., & Hammer, M. F. (2006). Reconstructing human origins in the genomic era. *Nature Reviews*, 7, 669-680. doi:10.1038/nrg1941
- [6] Cavalli-Sforza, L. L., Piazza, A., Menozzi, P., Mountain, J. (1988). Reconstruction of human evolution: Bringing together genetic, archaeological, and linguistic data. *Proceedings of the National Academy of Sciences*, 85, 6002-6006. doi:10.1073/pnas.85.16.6002
- [7] Chiaroni, J., Underhill, P. A., & Cavalli-Sforza, L. L. (2009). Y-chromosome diversity, human expansion, drift, and cultural evolution. *Proceedings of the National Academy of Sciences*, 106, 20174-20179. doi:10.1073/pnas.0910803106
- [8] Chromosomal Laboratories (2005). Y chromosomal haplogroup and ancient origins. <http://www.chromosomal-labs.com/ancestry/yhaplogroup.pdf>
- [9] Cruciani, F., Santolamazza, P., Shen, P., Macaulay, V., Moral, P., Olckers, A., Modiano, D. et al. (2002). A back migration from Asia to Sub-Saharan Africa is supported by high-resolution analysis of human Y-chromosome haplotypes. *The American Journal of Human Genetics*, 70, 1197-1214. doi:10.1086/340257
- [10] Cruciani, F., Trombetta, B., Sellitto, D., Mas-Saia, A., Destro-Bisol, G., Watson, E. et al. (2010). Human Y-chromosome haplogroup R-V88: A paternal genetic record of early mid Holocene trans-Saharan connections and the spread of Chadic languages. *European Journal of Human Genetics*, 23, 1-8.
- [11] Cruciani, F., Trombetta, B., Massaia, A., Destro-Bisol, G., Sellitto, D., & Scozzari, R. (2011). A revised root for the human Y chromosomal phylogenetic tree: The origin of patrilineal diversity in Africa. *The American Journal of Human Genetics*, 88, 1-5. doi:10.1016/j.ajhg.2011.05.002
- [12] Curnoe, D., Xueping, J., Herries, A. I. R., Kanning, B., Tacon, P. S. C., Zhende, B., Fink, D. et al. (2012). Human remains from the Pleistocene-holocene transition of Southwest China suggest a complex evolutionary history for East Asians. *PLOS One*, 7, e31918. doi:10.1371/journal.pone.0031918
- [13] Edmonds, C. A., Lillie, A. S., & Cavalli-Sforza, L. L. (2004) Mutations arising in the wave front of an expanding population. *Proceedings of the National Academy of Sciences*, 101, 975-979. doi:10.1073/pnas.0308064100
- [14] Fernandes, V., Alshamali, F., Alves, M., Costa, M. D., Pereira, J. B., Silva, N. M. et al. (2012). The Arabian cradle: Mitochondrial relicts of the first steps along the Southern route out of Africa. *The American Journal of Human Genetics*, 90, 347-355.
- [15] Green, R. E., Krause, J., Briggs, A. W., Maricic, T., Stenzel, U., Kircher, M., Patterson, N. et al. (2010). A draft sequence of the Neandertal genome. *Science*, 328, 710-722. doi:10.1126/science.1188021
- [16] Grine, F. E., Bailey, R. M., Harvati, K., Nathan, R. P., Morris, A. G., Henderson, G. M., Robot, I. et al. (2007). Late Pleistocene human skull from Hofmeyr, South Africa, and modern human origins. *Science*, 315, 226-229. doi:10.1126/science.1136294
- [17] Grine, F. E., Gunz, P., Betti-Nash, L., Neubauer, S., & Morris, A. G. (2010). Reconstruction of the late Pleistocene human skull from Hofmeyr, South Africa. *Journal of Human Evolution*, 59, 1-15. doi:10.1016/j.jhevol.2010.02.007
- [18] Gusmao, L., Gonzales-Neira, A., Alves, C., Lareu M, Costa S, Amorim A, & Carracedo A. (2002). Chimpanzee homologous of human Y specific STRs. A comparative study and a proposal for nomenclature. *Forensic Science Internships*, 126, 129-136.
- [19] Habgood, P. J. (1989). An examination of regional features on middle and early late Pleistocene Sub-Saharan African hominids. *South African Archaeological Bulletin*, 44, 17-22. doi:10.2307/3888315
- [20] Hanihara, T., Ishida, H., & Dodo, Y. (2003). Characterization of biological diversity through analysis of discrete cranial traits. *American Journal of Physical Anthropology*, 121, 241-251. doi:10.1002/ajpa.10233

- [21] Hellenthal, G., Auton, A., & Falush, D. (2008). Inferring human colonization history using a copying model. *PLOS Genetics*, 4, e1000078. doi:10.1371/journal.pgen.1000078
- [22] Henn, B. M., Gignoux, C. R., Jobin, M., Granka, J. M., Macpherson, J. M., Kidd, J. M., Rodríguez-Botigué, L. et al. (2011). Huntergatherer genomic diversity suggests a southern African origin for modern humans. *Proceedings of the National Academy of Sciences*, 108, 5154-5162. doi:10.1073/pnas.1017511108
- [23] Higham, T., Compton, T., Stringer, C., Jacobi, R., Shapiro, B., Trinkaus, E., Chandler, B. et al. (2011). The earliest evidence for anatomically modern humans in North-Western Europe. *Nature*, 479, 521-524. doi:10.1038/nature10484
- [24] Hoffecker, J. F. (2011). The early upper Paleolithic of Eastern Europe reconsidered. *Evolutionary Anthropology*, 20, 24-39. doi:10.1002/evan.20284
- [25] Hublin, J.-J. (2011). African origin. *Nature*, 476, 395. doi:10.1038/476395a
- [26] Hudhes, J. F., Skaletsky, H., Pyntikova, T., Graves, T. A., van Daalen, S. K. M., Minx, P. J., Fulton, R. S. et al. (2010). Chimpanzee and human Y-chromosomes are remarkably divergent in structure and gene content. *Nature*, 463, 536-539. doi:10.1038/nature08700
- [27] Hudhes, J. F., Skaletsky, H., Brown, L. G., Pyntikova, T., Graves, T. A., Fulton, R. S., Dugan, S. et al. (2012). Strict evolutionary conservation followed rapid gene loss on human and rhesus Y chromosomes. *Nature*, 483, 82-86. doi:10.1038/nature10843
- [28] Hudjashov, G., Kivisild, T., Underhill, P. A., Endicott, P., Sanchez, J. J., Lin, A. A., Shen, P. et al. (2007). Revealing the prehistoric settlement of Australia by Y-chromosome and mtDNA analysis. *Proceedings of the National Academy of Sciences*, 104, 8726-8730. doi:10.1073/pnas.0702928104
- [29] Jobling, M. A., & Tyler-Smith, C. (2003) The human Y-chromosome: an evolutionary marker comes of age. *Nature Reviews*, 4, 598-612. doi:10.1038/nrg1124
- [30] Klyosov, A. A. (2009). DNA genealogy, mutation rates, and some historical evidences written in Y-chromosome. I. Basic principles and the method. *Journal of Genetic Genealogy*, 5, 186-216.
- [31] Klyosov, A. A., & Rozhanskii, I. L. (2011). An archaic lineage of haplogroup A. *Proceedings of the Russian Academy of DNA Genealogy*, 4, 1495-1502.
- [32] Klyosov, A. A. (2012). Ancient history of the Arbins, bearers of haplogroup R1b, from central Asia to Europe, 16,000 to 1500 years before present. *Advances in Anthropology*, 2, 87-105. doi:10.4236/aa.2012.22010
- [33] Klyosov, A. A., & Rozhanskii, I. L. (2012a). Re-examining the out of Africa theory and the origin of Europeoids (Caucasoids) in light of DNA genealogy. *Advances in Anthropology*, 2, 80-86. doi:10.4236/aa.2012.22009
- [34] Klyosov, A. A., & Rozhanskii, I. L. (2012b). Haplogroup R1a as the proto indo-Europeans and the legendary Aryans as witnessed by the DNA of their current descendants. *Advances in Anthropology*, 2, 1-13. doi:10.4236/aa.2012.21001
- [35] Li, H., & Durbin, R. (2011). Inference of human population history from individual whole-genome sequences. *Nature*, 475, 493-496. doi:10.1038/nature10231
- [36] Majumder, P. P. (2010). The human genetic history of South Asia. *Current Biology*, 20, R184-R187. doi:10.1016/j.cub.2009.11.053
- [37] Mellars, P., & French, J. C. (2011). Tenfold population increase in Western Europe at the Neandertal-to-modern human transition. *Science*, 333, 623-627. doi:10.1126/science.1206930
- [38] Mellars, P. (2011). The earliest modern humans in Europe. *Nature*, 479, 483-485. doi:10.1038/479483a
- [39] Moorjani, P., Patterson, N., Hirschhorn, J. N., Keinan, A., Hao, L., Atzmon, G., Burns, E. et al. (2011). The history of African gene flow into Southern Europeans, Levantines, and Jews. *PLOS Genetics*, 7, e1001373. doi:10.1371/journal.pgen.1001373
- [40] Nei, M., & Takezaki, N. (1996). The root of the phylogenetic tree of human populations. *Molecular Biology and Evolution*, 13, 170-177. doi:10.1093/oxfordjournals.molbev.a025553
- [41] Ottoni, C., Primativo, G., Kashani, B. H., Achilli, A., Martinez-Labarga, C., Biondi, G., Torroni, A., &

Rickards, O. (2010). Mitochondrial haplogroup H1 in North Africa: An early holocene arrival from Iberia. *Plos One*, 5, e13378. doi:10.1371/journal.pone.0013378

- [42] Patin, E., Laval, G., Barreiro, L. B., Salas, A., Semino, O., Santachiara-Benerecetti, S., Kidd, K. K. et al. (2009). Inferring the demographic history of African farmers and Pygmy hunter-gatherers using a multilocus resequencing data set. *PLOS Genetics*, 5, 1-13.
- [43] Perelman, P., Johnson, W. E., Roos, C., Seuanez, H. N., Horvath, J. E., Moreira, M. A. M., & Kessing, B. (2011). A molecular phylogeny of living primates. *PLOS Genetics*, 7, e1001342. doi:10.1371/journal.pgen.1001342
- [44] Pinhasi, R., Higham, T. F. G., Golovanova, L. V., & Doronichev, V. B. (2011). Revised age of late Neanderthal occupation and the end of the middle Paleolithic in the northern Caucasus. *Proceedings of the National Academy of Sciences*, 108, 8611-8616.
- [45] Prat, S., Pean, S. C., Crepin, L., Drucker, D. G., Puaud, S. J., Valladas, H., Lasnickova-Galetova, M. et al. (2011). The oldest anatomically modern humans from far South-East Europe: Direct dating, culture and behavior. *PLOS One*, 6, e20843. doi:10.1371/journal.pone.0020834
- [46] Prüfer, K., Munch, K., Hellmann, I., Akagi, K., Miller, J. R., Walenz, B., Koren, S. et al. (2012). The bonobo genome compared with the chimpanzee and human genomes. *Nature*, 486, 527-531.
- [47] Redd, A. J., Agellon, A. B., Kearney, V. A., Contreras, V. A., Karafet, T., Park, H., de Knijff, P. et al. (2002). Forensic value of 14 novel STRs on the human Y-chromosome. *Forensic Science Internships*, 130, 97-111. doi:10.1016/S0379-0738(02)00347-X
- [48] Rightmire, G. P. (2009) Middle and later Pleistocene hominins in Africa and Southwest Asia. *Proceedings of the National Academy of Sciences*, 106, 16046-16050. doi:10.1073/pnas.0903930106
- [49] Rogers, A. R., & Jorde, L. B. (1995). Genetic evidence of modern human origins. *Human Biology*, 67, 1-36.
- [50] Rozhanskii, I. L., & Klyosov, A. A. (2011). Mutation rate constants in DNA genealogy (Y chromosome). *Advances in Anthropology*, 1, 26-34. doi:10.4236/aa.2011.12005
- [51] Sahoo, S., Singh, A., Himabindu, G., Banerjee, J., Sitalaximi, T., Gaikwad, S., Trivedi, R. et al. (2006). A prehistory of Indian Y-chromosomes: Evaluating demic diffusion scenarios. *Proceedings of the National Academy of Sciences*, 103, 843-848. doi:10.1073/pnas.0507714103
- [52] Scally, A., Dutheil, J. Y., Hillier, L. W., Jordan, G. E., Goodhead, I., Herrero, J., Hobolth, A., et al. (2012). Insights into hominid evolution from the gorilla genome sequence. *Nature*, 483, 169-175. doi:10.1038/nature10842
- [53] Shi, W., Ayub, Q., Vermeulen, M., Shao, R.-G., Zuniga, S., van der Gaag, K., de Knijff, P. et al. (2010). A worldwide survey of human male demographic history based on Y-STR and Y-SNP data from the HGDP-CEPH populations. *Molecular Biology and Evolution*, 27, 385-393. doi:10.1093/molbev/msp243
- [54] Simms, T. M., Martinez, E., Herrera, K. J., Wright, M. R., Perez, O. A., Hernandez, M., Ramirez, E. C., McCartney, Q., & Herrera, R. J. (2011). Paternal lineages signal distinct genetic contributions from British loyalists and continental Africans among different Bahamian islands. *American Journal of Physical Anthropology*, 146, 4594-608. doi:10.1002/ajpa.21616
- [55] Soares, P., Ermini, L., Thompson, N., Normina, M., Rito, T., Rohr, A., Salas, A., Oppenheimer, S., Macaulay, V., & Richards, M. B. (2009). Correcting for purifying selection: an improved human mitochondrial molecular clock. *Journal of Human Genetics*, 84, 740-759. doi:10.1016/j.ajhg.2009.05.001
- [56] Stewart, J. R., & Stringer, C. B. (2012). Human evolution out of Africa: The role of refugia and climate change. *Science*, 335, 1317-1321. doi:10.1126/science.1215627
- [57] Stoneking, M., & Delfin, F. (2010). The human genetic history of East Asia: Weaving a complex tapestry. *Current Biology*, 20, R188-R193. doi:10.1016/j.cub.2009.11.052
- [58] Stringer, C. B., & Andrews, P. (1988). Genetic and fossil evidence for the origin of modern humans. *Science*, 239, 1263-1268. doi:10.1126/science.3125610
- [59] Sun, J. X., Helgason, A., Masson, G., Ebenesersdóttir, S. S., Li, H., Mallick, S., Patterson, N. et al. (2011). A direct characterization of human mutation. 61st Annual Meeting, American Society of Human Genetics/ICHG, 11-15 October 2011, Montreal.

- [60] Tattersall, I. (2009). Human origins: Out of Africa. *Proceedings of the National Academy of Sciences*, 106, 16018-16031. doi:10.1073/pnas.0903207106
- [61] Underhill, P. A., Shen, P., Lin, A. A., Jin, L., Passarino, G., Yang, W. H., Kauffman, E. et al. (2000). Y-chromosome sequence variation and the history of human populations. *Nature Genetics*, 26, 358-361. doi:10.1038/81685
- [62] Xue, Y., Zerjal, T., Bao, W., Zhu, S., Lim, S.-K., Shu, Q. et al. (2005). Recent spread of a Y-chromosomal lineage in Northern China and Mongolia. *The American Journal of Human Genetics*, 77, 1112-1116. doi:10.1086/498583
- [63] Yotova, V., Lefebvre, J.-F., Moreau, C., Gbeha, E., Hovhannesyan, K., Bourgeois, S., & Bedarida, S. (2011). An X-linked haplotype of Neandertal origin is present among all non-African populations. *Molecular Biology and Evolution*, 28, 1957-1962

[Home](#) | [About SCIRP](#) | [Sitemap](#) | [Contact Us](#)

Copyright © 2006-2013 Scientific Research Publishing Inc. All rights reserved.