

Supplementary Material for: Towards a new history and geography of human genes informed by ancient DNA

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23 -em 0.001475 3 1 36363.6363636 -ej 0.0015 3 23 -em 0.001475 4 1
34545.4545455 -ej 0.0015 4 23 -em 0.001475 5 1 32727.2727273 -ej 0.0015
5 23 -em 0.001475 6 1 30909.0909091 -ej 0.0015 6 23 -em 0.001475 7 1
29090.9090909 -ej 0.0015 7 23 -em 0.001475 8 1 27272.7272727 -ej 0.0015
8 23 -em 0.001475 9 1 25454.5454545 -ej 0.0015 9 23 -em 0.001475 10 1
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7272.72727273 -ej 0.0015 41 45 -em 0.001475 42 23 5454.54545455 -ej
0.0015 42 45 -em 0.001475 43 23 3636.36363636 -ej 0.0015 43 45 -em
0.001475 44 23 1818.18181818 -ej 0.0015 44 45 -en 0.02725 45 0.0025 -ej
0.0275 45 23 -en 0.05475 23 0.0025 -ej 0.055 23 1

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In the demographic model in Figure 1C, there are three ancestral populations, two of which have admixed with an archaic population in different proportions. All observed populations are mixtures (in different proportions) of two of these populations. To simulate this scenario, we used the following command. The output of this command is 20 haplotypes from 46 populations (the archaic population, the three ancestral populations and the 42 observed populations). For plotting purposes, we ignore the archaic population and the three ancestral populations (populations 0, 22, 44 and 45 in the command):

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ms 920 1 -t 100 -r 10 100000 -I 46 20 20 20 20 20 20 20 20 20 20 20 20 20
20 20 20 20 20 20 20 20 20 20 20 20 20 20 20 20 20 20 20 20 20 20 20
20 20 20 20 20 20 20 20 20 20 -em 0.001475 2 1 38181.8181818 -ej 0.0015
2 23 -em 0.001475 3 1 36363.6363636 -ej 0.0015 3 23 -em 0.001475 4 1
34545.4545455 -ej 0.0015 4 23 -em 0.001475 5 1 32727.2727273 -ej 0.0015
5 23 -em 0.001475 6 1 30909.0909091 -ej 0.0015 6 23 -em 0.001475 7 1
29090.9090909 -ej 0.0015 7 23 -em 0.001475 8 1 27272.7272727 -ej 0.0015
8 23 -em 0.001475 9 1 25454.5454545 -ej 0.0015 9 23 -em 0.001475 10 1
23636.3636364 -ej 0.0015 10 23 -em 0.001475 11 1 21818.1818182 -ej
0.0015 11 23 -em 0.001475 12 1 20000.0 -ej 0.0015 12 23 -em 0.001475 13
1 18181.8181818 -ej 0.0015 13 23 -em 0.001475 14 1 16363.6363636 -ej
0.0015 14 23 -em 0.001475 15 1 14545.4545455 -ej 0.0015 15 23 -em
0.001475 16 1 12727.2727273 -ej 0.0015 16 23 -em 0.001475 17 1
10909.0909091 -ej 0.0015 17 23 -em 0.001475 18 1 9090.90909091 -ej

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0.0015 18 23 -em 0.001475 19 1 7272.72727273 -ej 0.0015 19 23 -em
0.001475 20 1 5454.54545455 -ej 0.0015 20 23 -em 0.001475 21 1
3636.36363636 -ej 0.0015 21 23 -em 0.001475 22 1 1818.18181818 -ej
0.0015 22 23 -em 0.001475 24 23 38181.81818181 -ej 0.0015 24 45 -em
0.001475 25 23 36363.63636363 -ej 0.0015 25 45 -em 0.001475 26 23
34545.4545455 -ej 0.0015 26 45 -em 0.001475 27 23 32727.2727273 -ej
0.0015 27 45 -em 0.001475 28 23 30909.0909091 -ej 0.0015 28 45 -em
0.001475 29 23 29090.9090909 -ej 0.0015 29 45 -em 0.001475 30 23
27272.7272727 -ej 0.0015 30 45 -em 0.001475 31 23 25454.5454545 -ej
0.0015 31 45 -em 0.001475 32 23 23636.3636364 -ej 0.0015 32 45 -em
0.001475 33 23 21818.1818182 -ej 0.0015 33 45 -em 0.001475 34 23
20000.0 -ej 0.0015 34 45 -em 0.001475 35 23 18181.8181818 -ej 0.0015 35
45 -em 0.001475 36 23 16363.6363636 -ej 0.0015 36 45 -em 0.001475 37 23
14545.4545455 -ej 0.0015 37 45 -em 0.001475 38 23 12727.2727273 -ej
0.0015 38 45 -em 0.001475 39 23 10909.0909091 -ej 0.0015 39 45 -em
0.001475 40 23 9090.90909091 -ej 0.0015 40 45 -em 0.001475 41 23
7272.72727273 -ej 0.0015 41 45 -em 0.001475 42 23 5454.54545455 -ej
0.0015 42 45 -em 0.001475 43 23 3636.36363636 -ej 0.0015 43 45 -em
0.001475 44 23 1818.18181818 -ej 0.0015 44 45 -em 0.02 23 46 4000 -em
0.020025 23 46 0 -em 0.02 1 46 20000 -em 0.020025 1 46 0 -ej 0.0275 45
23 -ej 0.055 23 1 -ej 1 46 1

Admixture tests. To generate Figure 2, we combined SNP data generated on Illumina chips from a number of sources (Li et al., 2008; Altshuler et al., 2010; Behar et al., 2010; Henn et al., 2011; Schlebusch et al., 2012). We excluded the Jewish populations from Behar et al. 2010. In total, the data set consisted of 103 populations and 256,540 SNPs (Supplementary Table 1).

We used ADMIXTOOLS (Patterson et al., 2012) to compute all possible f_3 -statistics of the form $f_3(A; B, C)$ on these populations. We considered an f_3 -statistic to be significant evidence for admixture in population A if it was at least three standard errors less than zero (corresponding to a P-value of about 0.001). In Supplementary Table 1, we list all populations, their approximate latitudes and longitudes, and the representatives of the admixing populations (if any). These representatives were chosen as the population pair B and C that give the minimum f_3 -statistic.

Table 1: Populations used in tests for admixture

Panel	Population	Country	Region	Lat	Long	P1	P2
Pagani	AFAR	Ethiopia	EASTAF	12	41	Sardinian	SUDANESE
Pagani	AMHARA	Ethiopia	EASTAF	10	39	Sardinian	SUDANESE
Pagani	ANUAK	Ethiopia	EASTAF	8	34	ARIBLACKSMITH	SUDANESE
Pagani	ARIBLACKSMITH						
Pagani	H	Ethiopia	EASTAF	6	37	NA	NA
Pagani	ARICULTIVATO						
Pagani	R	Ethiopia	EASTAF	6	39	Sardinian	Juhoansi
Pagani	ESOMALI	Ethiopia	EASTAF	9	42	Sardinian	SUDANESE
Pagani	GUMUZ	Ethiopia	EASTAF	10.78	35.57	NA	NA
Pagani	OROMO	Ethiopia	EASTAF	8	37	Sardinian	SUDANESE
Pagani	SOMALI	Ethiopia	EASTAF	5.15	46.2	Sardinian	SUDANESE
Pagani	SUDANESE	Sudan	EASTAF	12.86	30.21	NA	NA
Pagani	TYGRAY	Ethiopia	EASTAF	9	38	Sardinian	SUDANESE
Pagani	WOLAYTA	Ethiopia	EASTAF	6	37	MbutiPygmy	Sardinian

Henn	Khomani	SouthAfrica	SAF	26.97	20.79	Juhoansi	Belorussians
KSP	Khwe	Angola	SAF	-17.4	22.95	Mozabite	Juhoansi
KSP	Xun	Angola	SAF	-14.6	17.66	Yoruba	Juhoansi
KSP	GuiGana	Botswana	SAF	-23.6	24.66	Yoruba	Juhoansi
KSP	Juhoansi	Namibia	SAF	19.59	20.49	NA	NA
KSP	Nama	Namibia	SAF	22.55	17.07	Basque	Juhoansi
KSP	Karretjie	SouthAfrica	SAF	30.71	25.10	Russian	Juhoansi
Hgdp	BiakaPygmy	CAR	CAF	4	17	NA	NA
Hgdp	MbutiPygmy	Congo	CAF	1	29	NA	NA
Henn	HADZA	Tanzania	EASTAF	-3.38	36.68	NA	NA
Henn	SANDAWÉ	Tanzania	EASTAF	-6.18	35.74	Sardinian	Juhoansi
HapMap	LWK	Kenya	EASTAF	0.61	34.76	BiakaPygmy	Sardinian
Hgdp	BantuKenya	Kenya	EASTAF	-3	37	BiakaPygmy	Samaritians
Hgdp	BantuSouthAfrica	SouthAfrica	SAF	-21	18.7	YRI	Juhoansi
HapMap	YRI	Nigeria	WAF	8	5	NA	NA
Hgdp	Yoruba	Nigeria	WAF	8	5	NA	NA
Hgdp	Mandenka	Senegal	WAF	12	-12	NA	NA
HapMap	MKK	Kenya	EASTAF	-1.32	36.82	MbutiPygmy	Sardinian
KSP	ColouredWellington	SouthAfrica	SAF	-33.6	19.01	Russian	Juhoansi
Hgdp	Mozabite	Algeria-Mزاب	MIDDLEEAST	32	3	YRI	Sardinian
Hgdp	Druze	Israel-Carmel	MIDDLEEAST	32	35	NA	NA
Hgdp	Palestinian	Israel-Central	MIDDLEEAST	32	35	Yoruba	Sardinian
Hgdp	Bedouin	Israel-Negev	MIDDLEEAST	31	35	Sardinian	SUDANESE
HapMap	GIH	India	CSASIA	23	72	Paniya	Georgians
Hgdp	Uyгур	China	CSASIA	44	81	Italian	Japanese
Hgdp	Xibo	China	CSASIA	43.5	81.5	Lithuanians	Japanese
Hgdp	Balochi	Pakistan	CSASIA	30.5	66.5	Cypriots	Paniya
Hgdp	Brahui	Pakistan	CSASIA	30.5	66.5	Cypriots	Paniya
Hgdp	Buruscho	Pakistan	CSASIA	36.5	74	Georgians	Naxi
Hgdp	Hazara	Pakistan	CSASIA	33.5	70	Italian	Japanese
Hgdp	Kalash	Pakistan	CSASIA	36	71.5	NA	NA
Hgdp	Makrani	Pakistan	CSASIA	26	64	Legzins	BantuSouthAfrica
Hgdp	Pathan	Pakistan	CSASIA	33.5	70.5	Samaritians	Karitiana
Hgdp	Sindhi	Pakistan	CSASIA	25.5	69	Paniya	Georgians
HapMap	CEU	CEPH	NEUROPE	55	-3	Karitiana	Sardinian
HapMap	TSI	Italy	SEUROPE	43	11	Karitiana	Sardinian
Hgdp	French	France	NEUROPE	46	2	Karitiana	Sardinian
Hgdp	Basque	France	SEUROPE	43	0	NA	NA
Hgdp	Italian	Italy	SEUROPE	46	10	Karitiana	Sardinian
Hgdp	Sardinian	Italy	SEUROPE	40	9	NA	NA
Hgdp	Orcadian	Orkney	NEUROPE	59	-3	Karitiana	Sardinian
Hgdp	Russian	Russia	NEUROPE	61	40	Karitiana	Sardinian
Hgdp	Adygei	Russia-Caucasus	NEUROPE	44	39	Karitiana	Sardinian
HapMap	CHB	China	EASTASIA	32.5	114	Dai	Daur

HapMap	JPT	Japan	NEASTASIA	38	138	NA	NA
Hgdp	Cambodian	Cambodia	EASTASIA	12	105	Dai	Samaritians
Hgdp	Dai	China	EASTASIA	21	100	NA	NA
Hgdp	Daur	China	NEASTASIA	48.5	124	Han	Yakut
Hgdp	Han	China	EASTASIA	32.5	114	Dai	Daur
Hgdp	Hezhen	China	NEASTASIA	47.5	133.5	She	Yakut
Hgdp	Lahu	China	EASTASIA	22	100	NA	NA
Hgdp	Miaozu	China	EASTASIA	28	109	NA	NA
Hgdp	Mongola	China	NEASTASIA	48.5	119	Lithuanians	Japanese
Hgdp	Naxi	China	EASTASIA	26	100	NA	NA
Hgdp	Oroqen	China	NEASTASIA	50.5	126.5	Han	Yakut
Hgdp	She	China	EASTASIA	27	119	NA	NA
Hgdp	Tu	China	EASTASIA	36	101	CEU	Tujia
Hgdp	Tujia	China	EASTASIA	29	109	Dai	Hezhen
Hgdp	Yizu	China	EASTASIA	28	103	NA	NA
Hgdp	Japanese	Japan	EASTASIA	38	138	NA	NA
Hgdp	Yakut	Siberia	NEASTASIA	70	129.5	NA	NA
Hgdp	Melanesian	Bougainville	OCEANIA	-6	155	NA	NA
Hgdp	Papuan	NewGuinea	OCEANIA	-4	143	NA	NA
Hgdp	Karitiana	Brazil	AMERICA	-10	-63	NA	NA
Hgdp	Surui	Brazil	AMERICA	-11	-62	NA	NA
Hgdp	Colombian	Colombia	AMERICA	3	-68	NA	NA
Hgdp	Maya	Mexico	AMERICA	19	-91	Moroccans	Surui
Hgdp	Pima	Mexico	AMERICA	29	-108	NA	NA
Behar	Armenians	Armenia	CAUCASUS	40.07	45.04	GIH	Sardinian
Behar	Belorussians	Belarus	NEUROPE	53.71	27.95	Karitiana	Sardinian
Behar	Chuvaths	Russia	NEUROPE	55.56	46.93	Oroqen	Lithuanians
Behar	Cypriots	Cyprus	SEUROPE	35.13	33.43	Sardinian	ANUAK
Behar	Egyptans	Egypt	NAFRICA	26.82	30.8	Yoruba	Sardinian
Behar	Georgians	Georgia	CAUCASUS	42.32	43.36	NA	NA
Behar	Hungarians	Hungary	NEUROPE	47.16	19.5	Karitiana	Sardinian
Behar	Iranians	Iran	MIDDLEEAST	32.43	53.69	Samaritians	Colombian
Behar	Jordanians	Jordan	MIDDLEEAST	30.59	36.24	Sardinian	SUDANESE
Behar	Legzins	Russia	CAUCASUS	42.14	47.09	Samaritians	Colombian
Behar	Lithuanians	Lithuania	NEUROPE	55.17	23.88	NA	NA
Behar	Moroccans	Morocco	NAFRICA	31.79	-7.09	Yoruba	Sardinian
Behar	North_Kannadi	India	CSASIA	10.85	76.27	NA	NA
Behar	Paniya	India	CSASIA	11.71	76.1	NA	NA
Behar	Romanians	Romania	NEUROPE	45.94	24.97	Karitiana	Sardinian
Behar	Sakilli	India	CSASIA	11.13	78.66	NA	NA
Behar	Samaritians	Israel	MIDDLEEAST	32.07	34.78	NA	NA
Behar	Saudis	SaudiArabia	MIDDLEEAST	23.89	45.08	GuiGana	Sardinian
Behar	Spaniards	Spain	SEUROPE	40.46	-3.75	Karitiana	Sardinian
Behar	Syrians	Syria	MIDDLEEAST	34.8	39	MbutiPygmy	Sardinian
Behar	Turks	Turkey	MIDDLEEAST	38.96	35.24	Karitiana	Sardinian
Behar	Uzbeks	Uzbekistan	CAUCASUS	41.38	64.59	Oroqen	Sardinian
Behar	Yemenese	Yemen	MIDDLEEAST	15.55	48.52	MbutiPygmy	Sardinian