yourDNA Portal – Native American Ancestry Analysis of Each Chromosome for DKF

The following results from "yourDNAportal.com" purport to provide a detailed fine grained exploration of the ancestral results for each chromosome using different algorithms and reference samples. The author's assessment of this data will be provided at the end of the Results section below.

CHROMOSOME ANALYSIS

A) MODERN NATIVE AMERICAN REFERENCE GROUPS - RESULTS:

Chromosynthesis

Native American Modern Chromosynthesis

Explore your DNA
via the power of Chromosynthesis!

By utilising chromosomal analysis, you can dig deeper into the more obscure areas of your ancestry. The fine detail of a chromosomal test, looks at each chromosome individually, allowing you to capture ancestry that may be missed by other tests.

Chromosynthesis is an excellent tool for capturing even the most distant ancestry. Each chromosome contains unique information and can have a very different autosomal profile. Usually the first 10 chromosomes are the most relevant, while the last 2 are less informative. A small percentage (1% of a chromosome) equals approximately 0.045% of the entire autosome. By analysing each chromosome, we can reveal both a deeper and more detailed ancestry than other tests.

Chromosome 6

Speculative results

Population	Value
Basque	50%
North-Western European	37.5%
Central Mediterranean	6.25%
Chipewyan Saskatchewan Canada	6.25%

Conservative results

Population	Value
Basque	75%
North-Western European	25%

Chromosome 18

Speculative results

Population	Value
● Basque	75%
● South Asian	18.75%
● Shipibo Peru	6.25%

Conservative results

Population	Value
● Basque	75%
● South Asian	25%

Chromosome 22

Speculative results

Population	Value
● East-Central European	68.75%
● Central European	18.75%
● Huambisa Peru	6.25%
● Uros Titicaca Peru	6.25%

Conservative results

Population	Value
● East-Central European	50%
● Balkan	25%
● Central European	25%

B) <u>GENOMEWIDE ANALYSIS USING MODERN NA REFERENCE SAMPLES</u>:

Chromosynthesis

Native American Genomewide Chromosynthesis

Explore your DNA
via the power of Chromosynthesis!

By utilising chromosomal analysis, you can dig deeper into the more obscure areas of your ancestry. The fine detail of a chromosomal test, looks at each chromosome individually, allowing you to capture ancestry that may be missed by other tests.

Chromosynthesis is an excellent tool for capturing even the most distant ancestry. Each chromosome contains unique information and can have a very different autosomal profile. Usually the first 10 chromosomes are the most relevant, while the last 2 are less informative. A small percentage (1% of a chromosome) equals approximately 0.045% of the entire autosome. By analysing each chromosome, we can reveal both a deeper and more detailed ancestry than other tests.

Chromosome 6

Speculative results

Population	Value
Basque	50%
North-Western European	37.5%
Central Mediterranean	6.25%
Native American	6.25%

Conservative results

Population	Value
Basque	75%
North-Western European	25%

Chromosome 18

Speculative results

Population	Value
● Basque	75%
● South Asian	18.75%
● Native American	6.25%

Conservative results

Population	Value
● Basque	75%
● South Asian	25%

Chromosome 22

Speculative results

Population	Value
● East-Central European	81.25%
● Native American	12.5%
● South-Central Asian	6.25%

Conservative results

Population	Value
● East-Central European	50%
● Balkan	25%
● Central European	25%

C) ANCIENT NA REFERENCE GROUPS - RESULTS:

Chromosynthesis

Native American Ancient Chromosynthesis

Explore your DNA
via the power of Chromosynthesis!

By utilising chromosomal analysis, you can dig deeper into the more obscure areas of your ancestry. The fine detail of a chromosomal test, looks at each chromosome individually, allowing you to capture ancestry that may be missed by other tests.

Chromosynthesis is an excellent tool for capturing even the most distant ancestry. Each chromosome contains unique information and can have a very different autosomal profile. Usually the first 10 chromosomes are the most relevant, while the last 2 are less informative. A small percentage (1% of a chromosome) equals approximately 0.045% of the entire autosome. By analysing each chromosome, we can reveal both a deeper and more detailed ancestry than other tests.

Chromosome 1

Speculative results

Population	Value
Insular Celtic	62.5%
Ancient Caucasus	18.75%
Ancient West Finnic	12.5%
Ancient Central American	6.25%

Conservative results

Population	Value
Insular Celtic	50%
Ancient Caucasus	25%
Ancient West Finnic	25%

Chromosome 6

Speculative results

Population	Value
● Ancient West Mediterranean	75%
● Insular Celtic	18.75%
● Ancient North American	6.25%

Conservative results

Population	Value
● Ancient West Mediterranean	75%
● Insular Celtic	25%

Chromosome 7

Speculative results

Population	Value
● Insular Celtic	37.5%
● Mediterranean Neolithic Farmer	37.5%
● Ancient Caucasus	12.5%
● Ancient Patagonian	6.25%
● Ancient West Mediterranean	6.25%

Conservative results

Population	Value
● Ancient Caucasus	25%
● Ancient West Mediterranean	25%
● Insular Celtic	25%
● Mediterranean Neolithic Farmer	25%

Chromosome 14

Speculative results

Population	Value
● Mediterranean Neolithic Farmer	50%
● Ancient Central American	12.5%
● Ancient East Mediterranean	12.5%
● Ancient West Mediterranean	12.5%
● Ancient Oceanian	6.25%
● Sub-Saharan African	6.25%

Conservative results

Population	Value
● Mediterranean Neolithic Farmer	50%
● Ancient Central American	25%
● Ancient West Mediterranean	25%

Chromosome 16

Speculative results

Population	Value
● Ancient West Finnic	50%
● Ancient East Mediterranean	37.5%
● Ancient North American	6.25%
● Ancient South-Central Asian	6.25%

Conservative results

Population	Value
● Ancient East Mediterranean	50%
● Ancient West Finnic	50%

Chromosome 17

Speculative results

Population	Value
● Ancient West Mediterranean 🐾	56.25%
● Ancient Caucasus 🐾	37.5%
● Ancient North American 🐾	6.25%

Conservative results

Population	Value
● Ancient Caucasus 🐾	50%
● Ancient West Mediterranean 🐾	50%

Chromosome 18

Speculative results

Population	Value
● Ancient West Mediterranean 🐾	75%
● Ancient South Asian 🐾	18.75%
● Ancient West Amazonian 🐾	6.25%

Conservative results

Population	Value
● Ancient West Mediterranean 🐾	75%
● Ancient South Asian	25%

Chromosome 22

Speculative results

Population	Value
● Balto-Slavic 🐾	81.25%
● Ancient Andean 🐾	6.25%
● Ancient Central American 🐾	6.25%
● Ancient South-Central Asian 🐾	6.25%

Conservative results

Population	Value
● Balto-Slavic 🐾	75%
● Ancient Central American 🐾	25%

D) <u>LANIAKEA ANALYSIS</u>:

MODERN

Laniakea Global
Chromosynthesis

This test utilises Laniakea Global to give you a chromosome by chromosome analysis of your genetic identity. For each chromosome, you will get:

1. Admixture results
2. The populations most genetically similar
3. An estimation of ethnicity

Chromosome 1

Admixture Results

Your admixture results are derived directly from the SNPs present in your file. They reveal the percentages that you share with the principal components of the populations present in the calculator, based on the frequency of alleles.

Admixture Population	Percentage
European Northatlantic	30.69%
European Fennoscandian	27.94%
Mediterranean East	21.2%
European Germanic	12.64%
Asian Far-East	3.2%
Eurasian Alanic	1.67%
Native American South	1.54%
Native American Central	0.84%
European Atlantic	0.29%

Asian Far-East : 3.2%

The Asian Far-East component has its peak frequency in Japan, Korea and in eastern and north-eastern China. The component is fairly widespread and common throughout East Asia from the Philippines to Mongolia.

Native American South : 1.54%

The American South component has its frequency peak among the indigenous peoples of the Andes but is widespread and common throughout South America and more generally all the Americas.

Native American Central : 0.84%

The Native American Central component has its frequency peak in the descendants of the Mayan and Aztec peoples, however it is widespread and common in all the Americas.

In-depth Origins results

Population	Value
● Northern Celtogermanic 🦶	54%
● Continental Celtogermanic 🦶	22.5%
● Finnish 🦶	19%
● Eastern North African 🦶	2%
● Arabic 🦶	1.5%
● Japanese 🦶	0.5%
● Polynesian 🦶	0.5%

Chromosome 6

Admixture Results

Your admixture results are derived directly from the SNPs present in your file. They reveal the percentages that you share with the principal components of the populations present in the calculator, based on the frequency of alleles.

Admixture Population	Percentage
● European Fennoscandian	44.69%
● European Atlantic	25.02%
● European Northatlantic	16.82%
● Mediterranean West	9.67%
● Native American North	2.0%
● African Rainforest	1.81%

Native American North : 2.0%

The Nativce American North component has its peak of frequency among the indigenous populations of Canada and the USA, but it is widespread and common in all the Americas, to a lesser extent also in Greenland and Siberia.

Chromosome 7

Admixture Results

Your admixture results are derived directly from the SNPs present in your file. They reveal the percentages that you share with the principal components of the populations present in the calculator, based on the frequency of alleles.

Admixture Population	Percentage
● European Fennoscandian	27.33%
● European Central	25.34%
● Mediterranean West	25.16%
● Mediterranean East	6.46%
● European Atlantic	5.56%
● MENA Maghrebi	3.59%
● MENA Mesopotamian	3.46%
● European Northatlantic	1.59%
● Asian Miao-Yi-Dai	1.52%

Asian Miao-Yi-Dai : 1.52%

The Miao-Yi-Dai component has its frequency peak in southern China but is widespread and common also in northern and western China, Taiwan and Japan, a good part of the Austronesian spectrum.

Chromosome 10

Admixture Results

Your admixture results are derived directly from the SNPs present in your file. They reveal the percentages that you share with the principal components of the populations present in the calculator, based on the frequency of alleles.

Admixture Population	Percentage
● European Northatlantic	37.67%
● European Germanic	28.52%
● European Central	14.33%
● European Eastern	10.17%
● Eurasian Alanic	5.09%
● Native American North	3.32%
● African Khoisan	0.52%
● Oceanian	0.38%

In-depth Origins results

Population	Value
● Northern Celtogermanic	99.5%
● Northern Amerindian	0.5%

Chromosome 12

Admixture Results

Your admixture results are derived directly from the SNPs present in your file. They reveal the percentages that you share with the principal components of the populations present in the calculator, based on the frequency of alleles.

Admixture Population	Percentage
● European Germanic	42.7%
● European Central	17.66%
● European Fennoscandian	12.91%
● European Atlantic	9.62%
● European Northatlantic	6.8%
● Native American South	5.48%
● Mediterranean East	2.46%
● Asian Gedrosia	2.36%

Population	Value
● Scandinavian 🏹	57%
● Continental Celtogermanic 🏹	34.5%
● Central Mediterranean 🏹	8%
● Central Amerindian 🏹	0.5%

Chromosome 15

Admixture Results

Your admixture results are derived directly from the SNPs present in your file. They reveal the percentages that you share with the principal components of the populations present in the calculator, based on the frequency of alleles.

| Admixture Population	Percentage
● European Central	49.75%
● European Northatlantic	20.09%
● European Atlantic	13.52%
● European Fennoscandian	6.19%
● Oceanian	3.46%
● Asian Miao-Yi-Dai	3.07%
● MENA Armenian	2.91%
● European Germanic	1.01%

Chromosome 16

Admixture Results

Your admixture results are derived directly from the SNPs present in your file. They reveal the percentages that you share with the principal components of the populations present in the calculator, based on the frequency of alleles.

Admixture Population	Percentage
● European Northatlantic	28.59%
● European Central	12.78%
● European Atlantic	12.38%
● European Fennoscandian	10.43%
● MENA Mesopotamian	10.26%
● Mediterranean East	9.74%
● Native American North	6.99%
● Eurasian Alanic	3.45%
● Asian Miao-Yi-Dai	2.15%
● Asian Tuvan	1.39%
● Asian Siberian	1.34%
● African Khoisan	0.48%

In-depth Origins results

Population	Value
● Northern Celtogermanic 🐾	73.5%
● Southern Italian 🐾	21%
● Arctic 🐾	2.5%
● Northern Amerindian 🐾	1%
● Levantine 🐾	0.5%
● Mongolian 🐾	0.5%
● Northern Chinese 🐾	0.5%
● Southern Chinese 🐾	0.5%

● Arctic 🐾

2.5%

Arctic

The Arctic group represents the genetic admixture of the Inuit people.

Chromosome 17

Admixture Results

Your admixture results are derived directly from the SNPs present in your file. They reveal the percentages that you share with the principal components of the populations present in the calculator, based on the frequency of alleles.

Admixture Population	Percentage
● European Atlantic	27.08%
● European Germanic	23.77%
● MENA Mesopotamian	19.18%
● MENA Armenian	13.31%
● European Northatlantic	10.85%
● Native American North	5.81%

In-depth Origins results

Population	Value
● Northern Celtogermanic	43%
● Iberian	17.5%
● Central Mediterranean	14%
● Levantine	10%
● Northern Italian	9.5%
● Arctic	2%
● Iranic	1.5%
● Northern Amerindian	1.5%
● Eastern North African	1%

Chromosome 18

Admixture Results

Your admixture results are derived directly from the SNPs present in your file. They reveal the percentages that you share with the principal components of the populations present in the calculator, based on the frequency of alleles.

Admixture Population	Percentage
European Atlantic	37.2%
European Germanic	21.56%
Mediterranean East	12.14%
Asian Dravidian	9.37%
Oceanian	7.33%
Native American South	4.66%
European Northatlantic	3.41%
Asian Miao-Yi-Dai	2.23%
European Fennoscandian	2.11%

In-depth Origins results

Population	Value
Iberian	76.5%
Northern Celtogermanic	8%
Northeastern Indian & Bangladeshi	5%
Australian	3%
Southern Indian	2.5%
Continental Celtogermanic	2%
Melanesian	1.5%
Polynesian	1%
Central Amerindian	0.5%

Chromosome 22

Admixture Results

Your admixture results are derived directly from the SNPs present in your file. They reveal the percentages that you share with the principal components of the populations present in the calculator, based on the frequency of alleles.

Admixture Population	Percentage
● European Germanic	34.94%
● European Fennoscandian	29.82%
● European Northatlantic	17.25%
● MENA Armenian	12.29%
● Native American Central	3.14%
● Mediterranean West	2.46%
● MENA Arabian	0.09%

Analysis: The present author is unsure of how to interpret these results such that they make a valid and trustworthy contribution to the study of the ancestral composition of each chromosome (recalling that both the maternal and the paternal chromosomes are being combined here). Consistency (internal and external) is critical for arriving at a conclusion as to the merits of the data above. Clearly the results are somewhat "scattered" and vary from test procedure to test procedure, and are difficult to reconcile with data from other calculators (e.g., 23andMe, Gedmatch). For example, the array of ancestries (e.g., various Siberian groups, Southeast Asian) on chromosome 16 with the Laniakea calculator does not seem to be duplicated elsewhere. If those results were associated with chromosome 18, the results would make a lot more sense based on testing using different methodologies. It would seem that there is some tweaking of the components of their test procedures that are needed before any of the above results can be "taken to the bank".

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