

Human migration routes into the Urumqi region of Xinjiang, China: Perspectives from The Genographic Project and Google Earth

Carl Tape

May 8, 2006

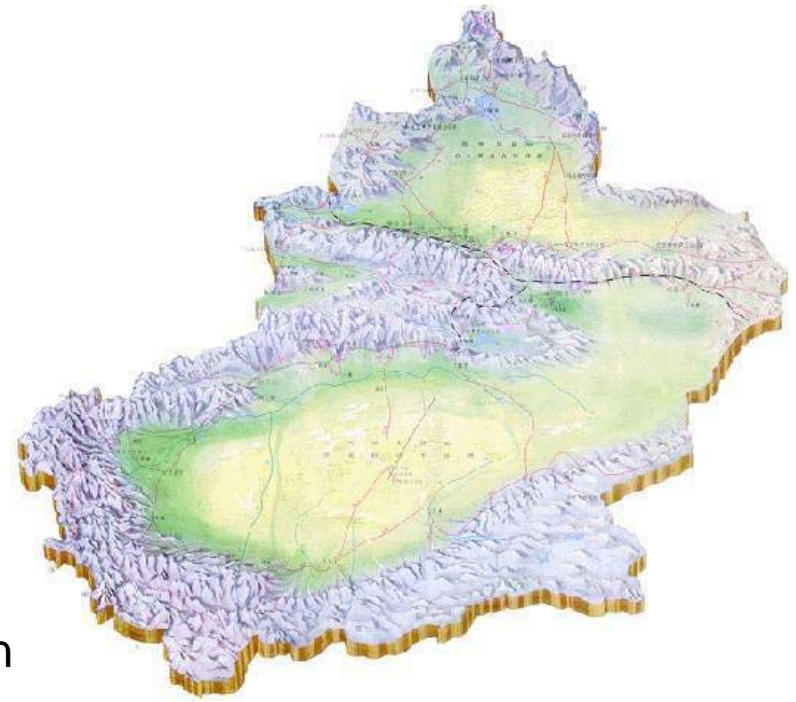
Caltech GPS trip to Tian Shan (June 2006)

The Genographic Project (National Geographic)

<https://www3.nationalgeographic.com/genographic/>

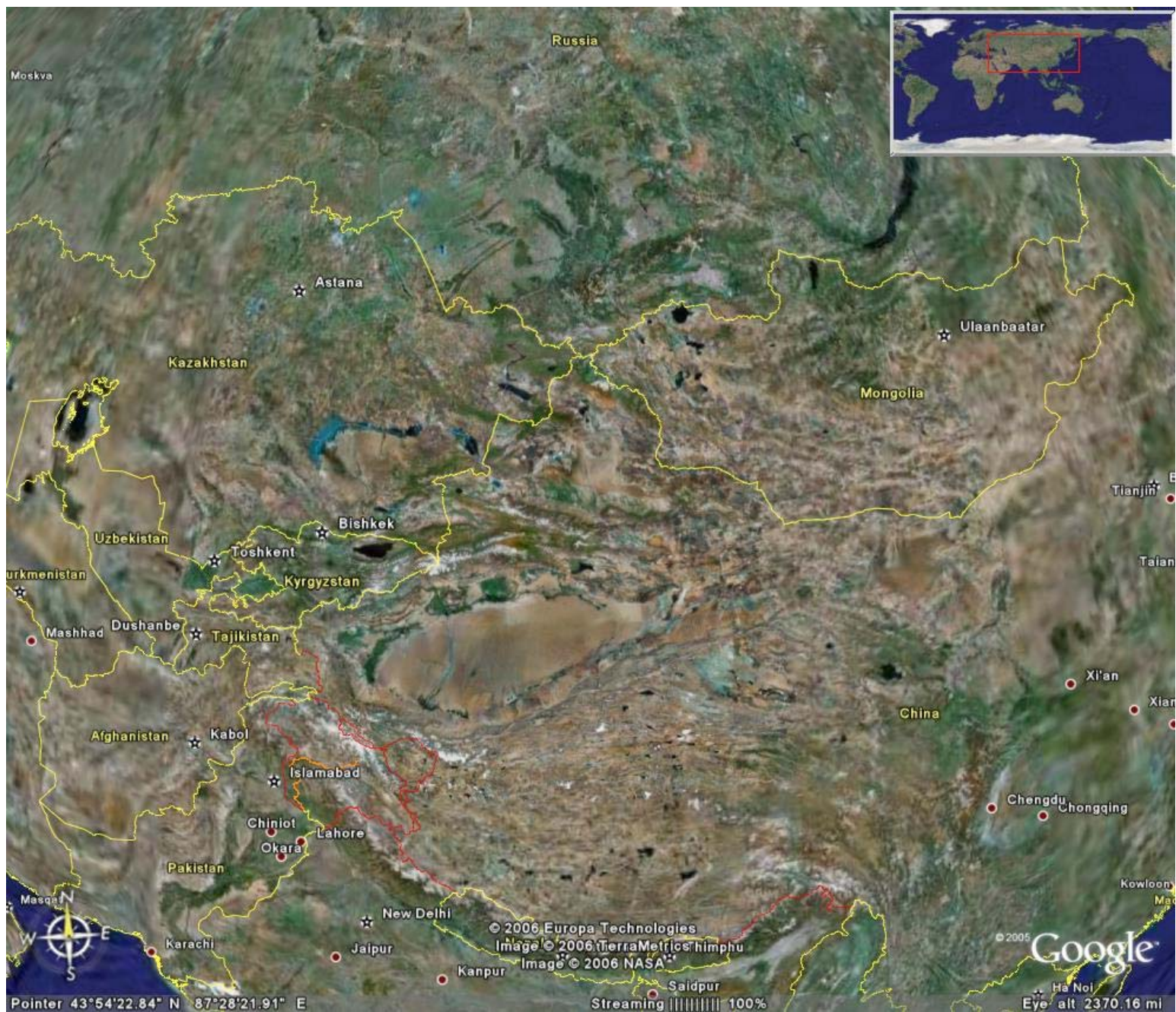
Google Earth

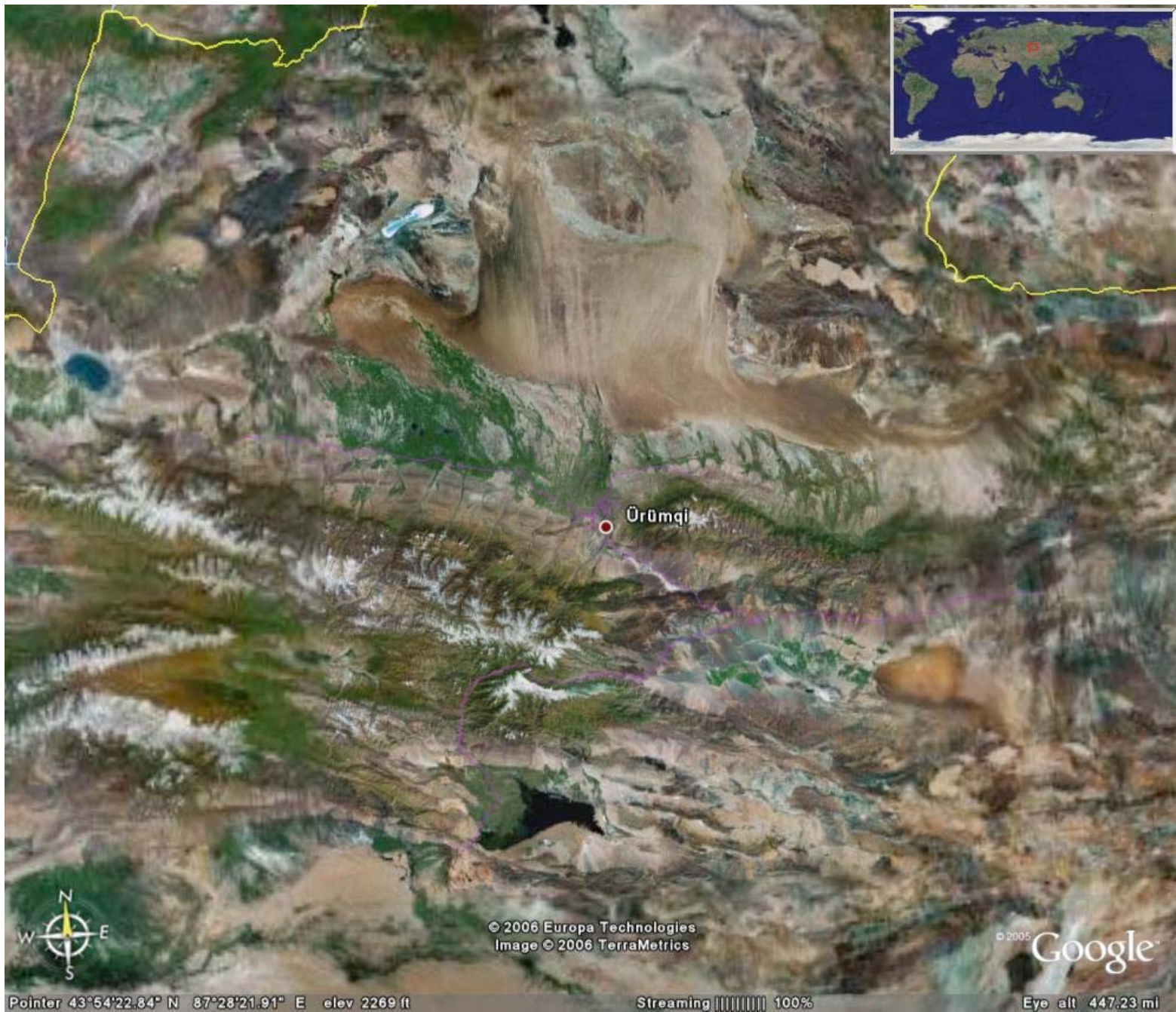
<http://earth.google.com>



1. **Name:** Xinjiang Uygur Autonomous Region
2. **Area:** 1.6604 million square kilometers
3. **Population:** 19.25 million (the 2000 population census)
4. **Capital:** Urumqi
5. **Geography:** Xinjiang is situated in the northeastern border area of China. It borders Mongolia, Russia, Kazakhstan, Kyrgyzstan, Tajikistan, Afghanistan, Pakistan, and India, among others. In China, it adjoins Tibet, Qinghai, Gansu and other provinces.







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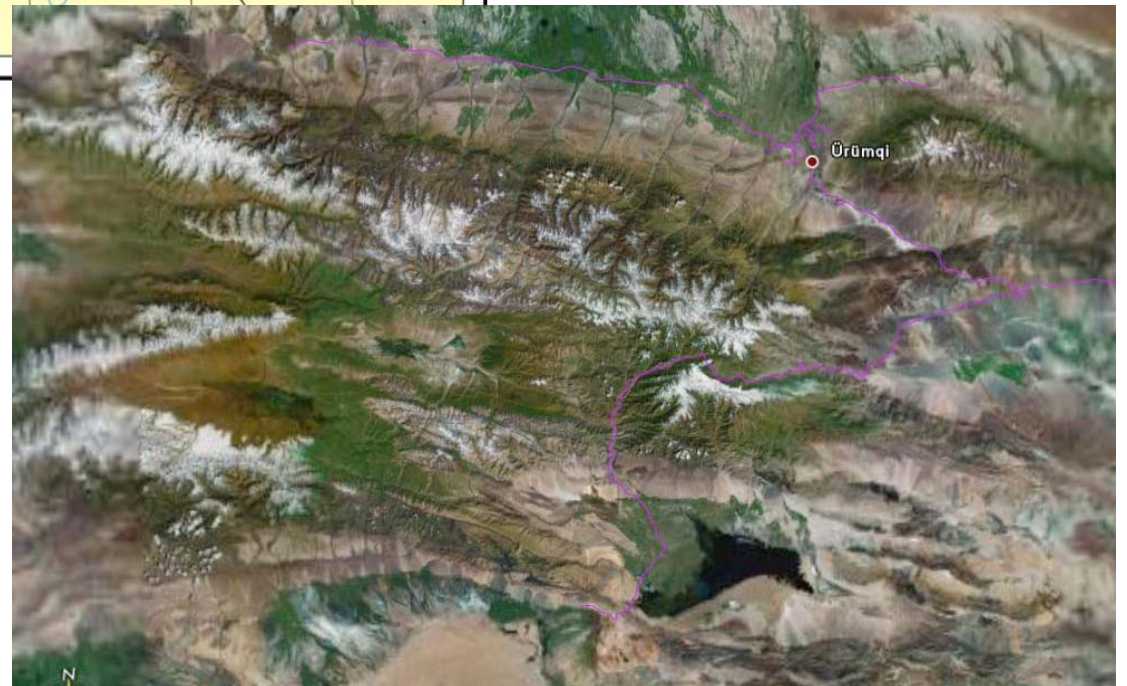
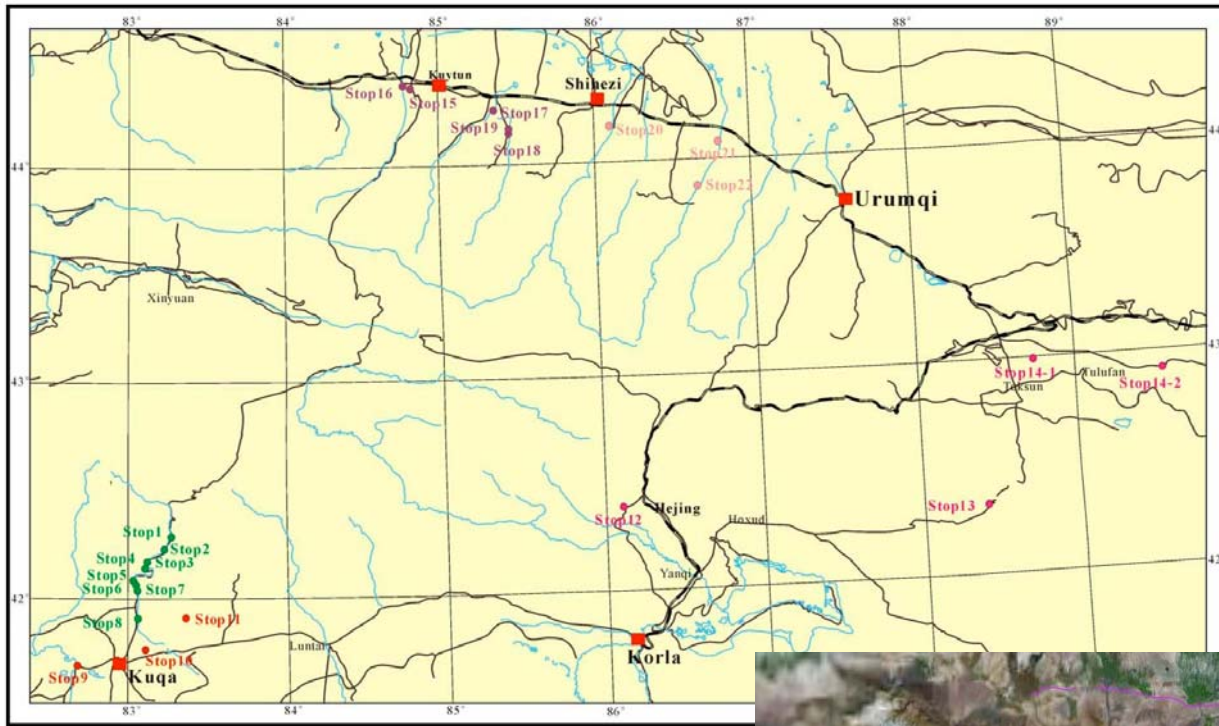
© 2005 Google

Pointer 43°54'22.84° N 87°28'21.91° E elev 2269 ft

Streaming ||||| 100%

Eye alt 447.23 mi

Field Stop Overview







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THE
GENOGRAPHIC
PROJECT

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MAIN
MENU

GENETICS
OVERVIEW

ATLAS OF THE
HUMAN JOURNEY

YOUR GENETIC
JOURNEY

Introduction

Introduction

200,000 B.C.

60,000

50,000

40,000

30,000

20,000

10,000 B.C.



SELECT AN ERA TO EXPLORE

ATLAS OF THE HUMAN JOURNEY

When humans first ventured out of Africa some 60,000 years ago, they left genetic footprints still visible today. By mapping the appearance and frequency of genetic markers in modern peoples, we create a picture of when and where ancient humans moved around the world. These great migrations eventually led the descendants of a small group of Africans to occupy even the farthest reaches of the Earth.



GO TO: [GENETIC MARKERS](#) + [JOURNEY HIGHLIGHTS](#) +

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THE HUMAN BODY

1 2 3 4

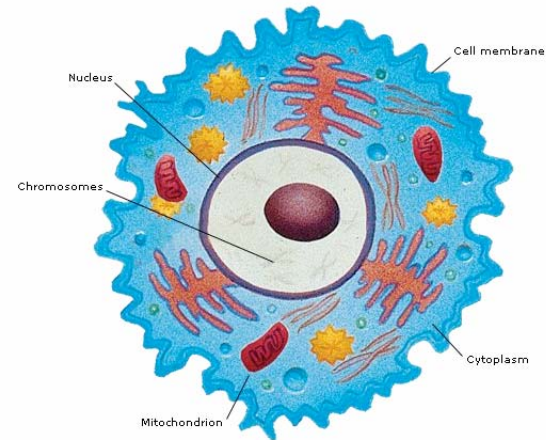
The human body is made of some 50 to 100 trillion cells, which form the basic units of life and combine to form more complex tissues and organs. Inside each cell, genes comprise a "blueprint" for protein production that determines how the cell will function. Genes also determine physical characteristics, or traits. The complete set of some



CELLS

1 2 3 4

"little organs," called organelles. The most important of these is the nucleus, which controls the cell and houses the genetic material in structures called chromosomes. Another type of organelle is the mitochondria. These "cellular power plants" have their own genome, and do not recombine during reproduction.



CHROMOSOMES

1 2 3 4

Chromosomes carry hereditary, genetic information in long strings of DNA called genes. Humans have 22 numbered pairs of chromosomes and a single pair of sex chromosomes; XX in females and XY in males. Each chromosomal pair includes one inherited from the father and one from the mother. If unwound, the microscopic DNA

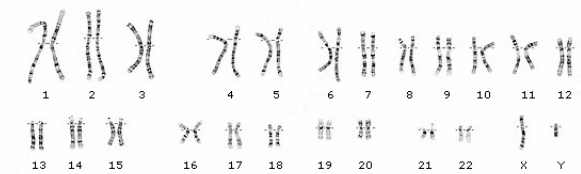
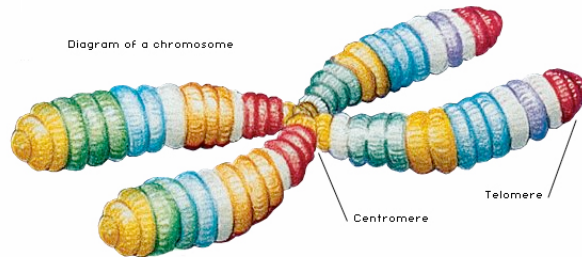


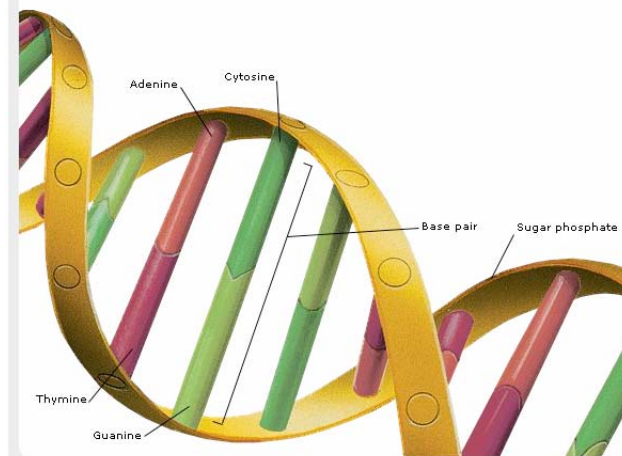
Diagram of a chromosome



DNA

1 2 3 4

DNA (deoxyribonucleic acid) is the set of genetic instructions for creating an organism. DNA molecules are shaped like a spiral staircase called a double helix. Each stair is composed of the DNA bases A, C, T, and G. Some segments of these bases contain sequences, like A-T-C-C-G-A-A-C-T-A-G, which constitute individual genes. Genes

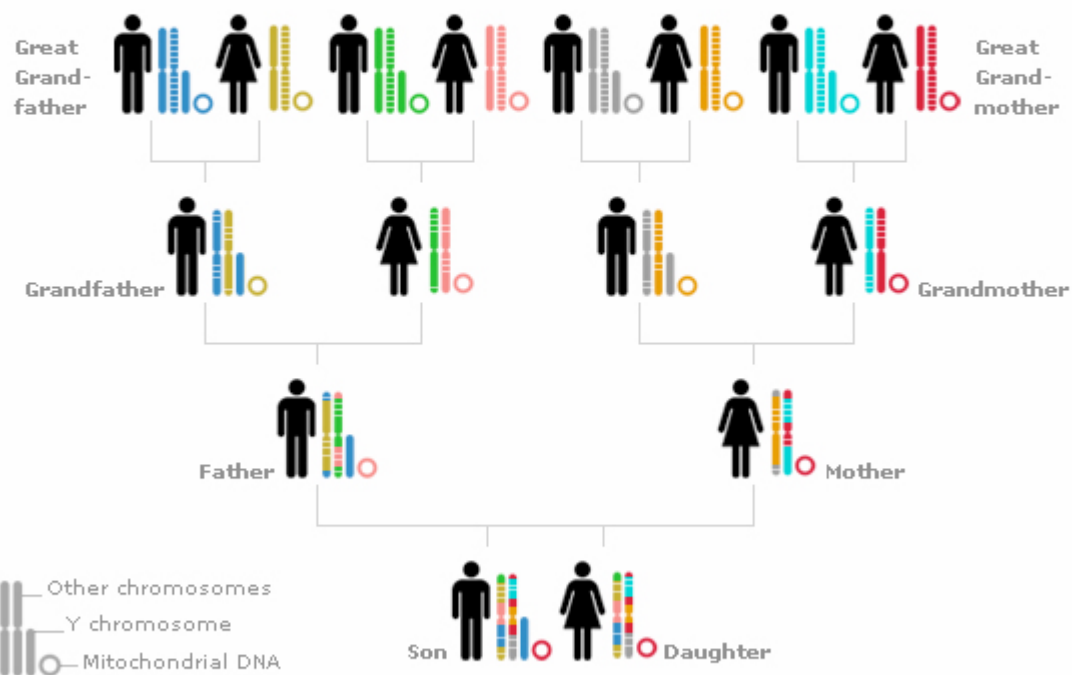


The Basic Idea

SHUFFLING THE DECK

1 2 3 ▶

For most of our genome we receive half of our genes from our father and from our mother. Each half represents a shuffled combination of DNA passed down to us from our ancestors. This recombination process makes it difficult to study lines of descent—it creates a genetic mix of everyone who has come before. Fortunately for



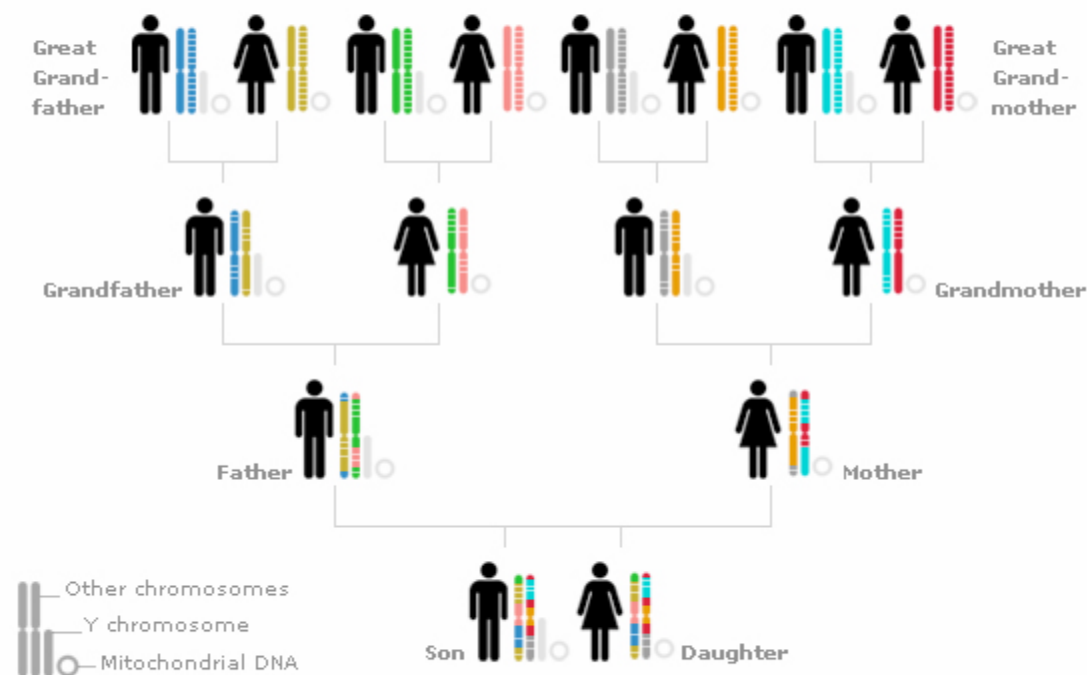
☒ Show both types of DNA ☐ Show recombined DNA ☐ Show non-recombined DNA

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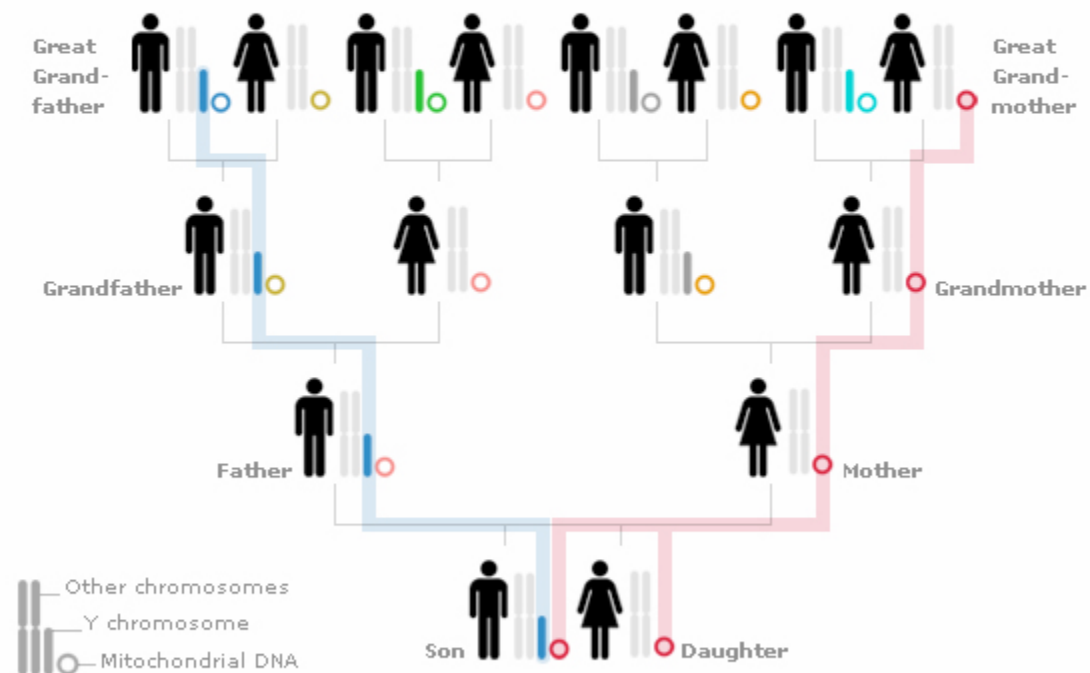
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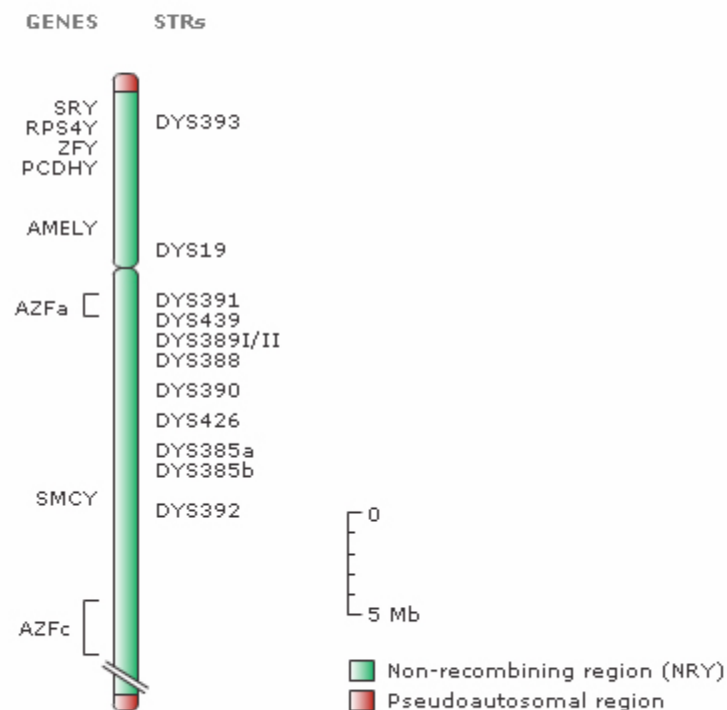
☐ Show both types of DNA ☐ Show recombined DNA ☒ Show non-recombined DNA

The Basic Idea

Y CHROMOSOME

◀ 1 2 3 ▶

The Y chromosome is the sex-determining chromosome in humans. While all other chromosomes are found in matching pairs, it is the mismatch of the Y with its partner, the X chromosome, that determines gender—men have a mismatched pair (Y and X), while women have two X chromosomes. Because the Y does not have a matching

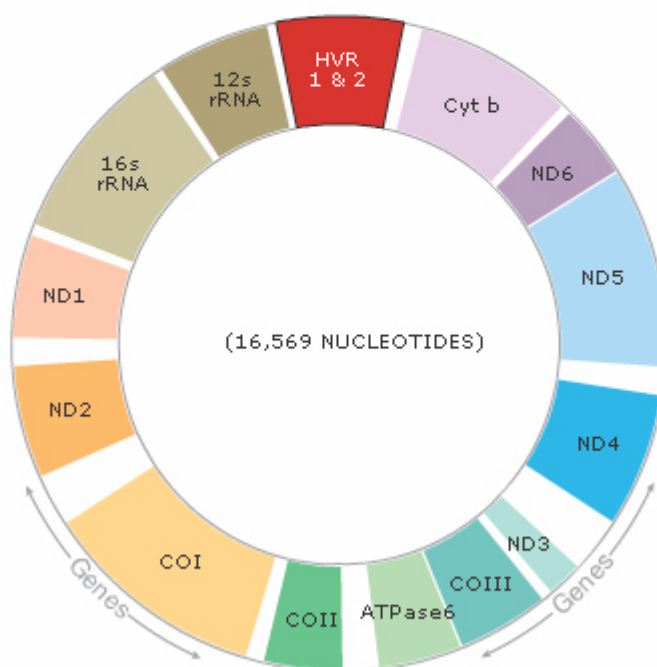


The Basic Idea

MITOCHONDRIAL DNA (mtDNA)

◀ 1 2 3

If the Y chromosome traces the male lineage back through history, then the mitochondrial genome (mtDNA) can be considered its female counterpart. Mitochondria are self-reproducing structures found inside the cells of all higher organisms, typically present in hundreds of copies per cell. They are responsible for




The Basic Idea

GENETIC DIVERSITY

◀ 1 2 3 4 5 ▶

As genetic markers are inherited, they are passed down through generations, forming a complex story that can be traced backward in time. The exact shape of this tree is affected by other evolutionary forces: natural selection, genetic drift, and migration.

Genetic marker

Follow the passage of genetic markers through one man's male descendants. Each generation inherits this man's unique marker.

NEXT GENERATION

Genetic Marker

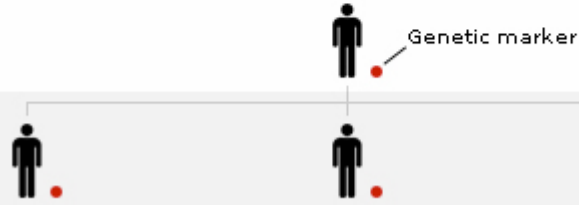
Random mutations in the DNA sequence which act as genetic milestones. Once markers have been identified they can be traced back in time to their origin—the most recent common ancestor of everyone who carries the marker.

The Basic Idea

GENETIC DIVERSITY

◀ 1 2 3 4 5 ▶

As genetic markers are inherited, they are passed down through generations, forming a complex story that can be traced backward in time. The exact shape of this tree is affected by other evolutionary forces: natural selection, genetic drift, and migration.



If this man has three sons, each of them would also inherit his marker, since it is passed through the Y-chromosome.

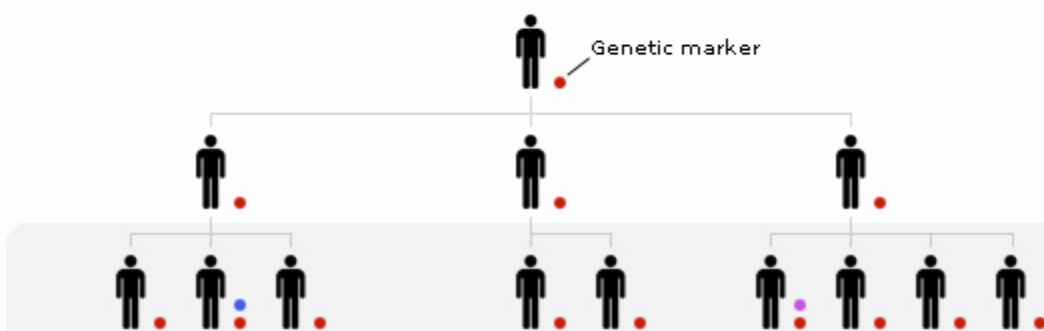
NEXT GENERATION

The Basic Idea

GENETIC DIVERSITY

◀ 1 2 3 4 5 ▶

As genetic markers are inherited, they are passed down through generations, forming a complex story that can be traced backward in time. The exact shape of this tree is affected by other evolutionary forces: natural selection, genetic drift, and migration.



Two new Y-chromosome markers appear in the third generation. These are caused by natural genetic mutations.

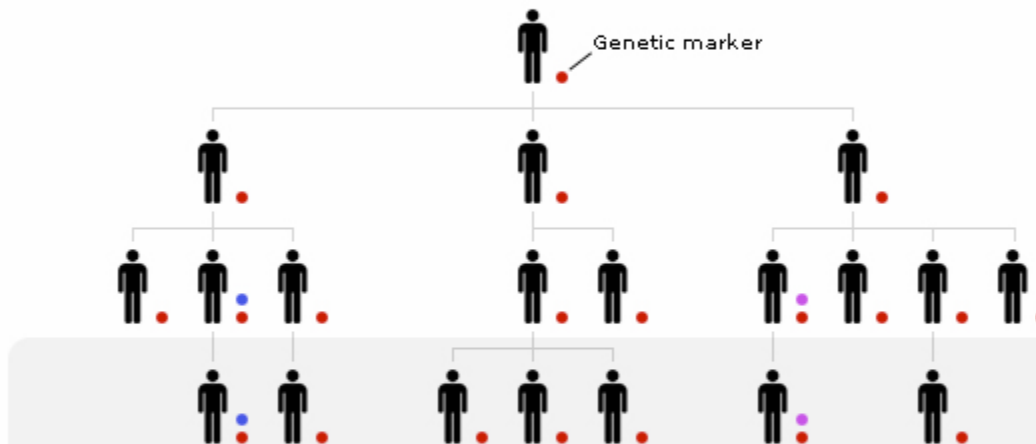
NEXT GENERATION

The Basic Idea

GENETIC DIVERSITY

◀ 1 2 3 4 5 ▶

As genetic markers are inherited, they are passed down through generations, forming a complex story that can be traced backward in time. The exact shape of this tree is affected by other evolutionary forces: natural selection, genetic drift, and migration.



Fourth generation offspring inherit the new markers that originated with their fathers.

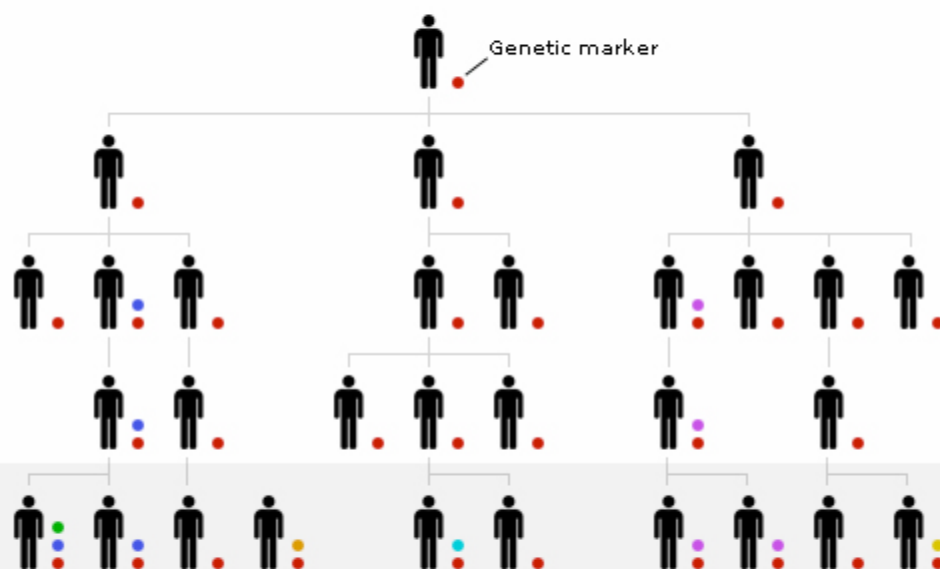
NEXT GENERATION

The Basic Idea

GENETIC DIVERSITY

◀ 1 2 3 4 5 ▶

As genetic markers are inherited, they are passed down through generations, forming a complex story that can be traced backward in time. The exact shape of this tree is affected by other evolutionary forces: natural selection, genetic drift, and migration.



New unique markers appear in the fifth generation. Note that even brothers do not share these individual mutations.

NEXT GENERATION

The Basic Idea

GENETIC DIVERSITY

As genetic markers are inherited, they are passed down through generations, forming a complex story that can be traced backward in time. The exact shape of this tree is affected by other evolutionary forces: natural selection, genetic drift, and migration.

After six generations, there are six surviving combinations of markers (called haplotypes) descended from one ancestor.

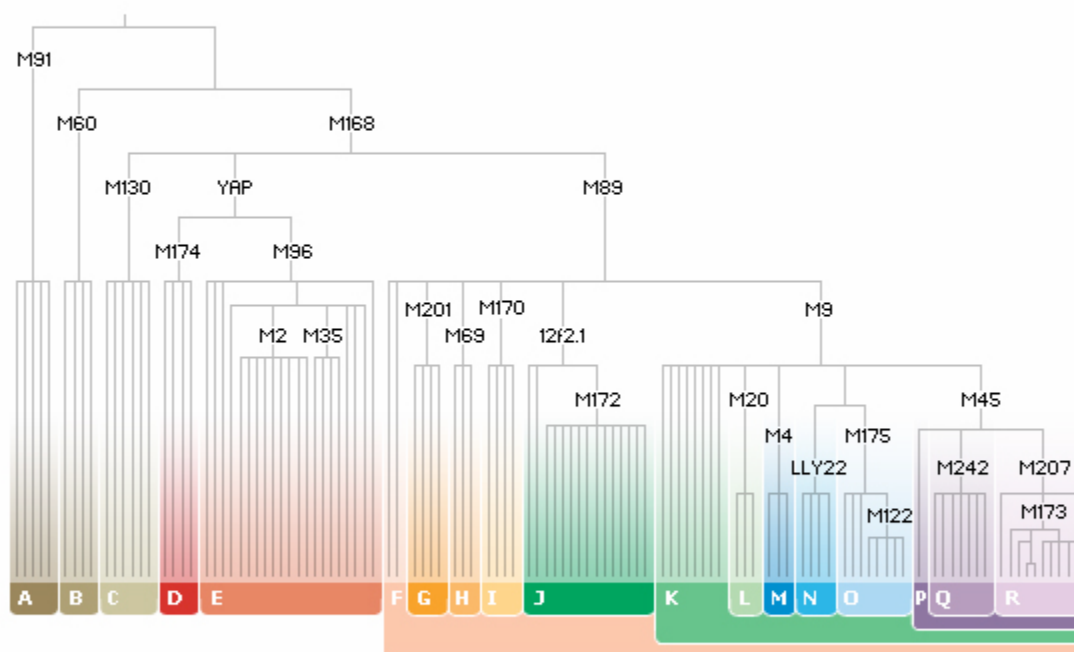
RESET

The Basic Idea

THE HUMAN FAMILY TREE

◀ 1 2 3 4 5

Y chromosome DNA, passed from father to son, and mitochondrial DNA, passed from mother to daughter, are varied through the generations only by occasional natural mutations called markers. These mutations, occurring in an otherwise continuous string of genetic replication, serve as genetic signposts for tracing human evolution.



Colors in tree denote particular haplogroups

☒ Y Chromosome tree ☐ Mitochondrial DNA Tree

The Basic Idea (for \$99.95)

YOUR GENETIC JOURNEY

Did you ever wonder about your most ancient ancestors? The Genographic Project will introduce you to them, and explain the genetic journeys that bond your personal lineage over tens of thousands of years.

Learn more
about the
results you
will receive



STEP 1: TEST

Once you have purchased your own **Genographic Project Public Participation Kit**, you can begin the exploration into your deep ancestry. The first step involves a painless cheek swab to acquire a DNA sample. Once you have completed the cheek swabbing process, you will secure the swabs inside the transport tubes and mail the tubes off to the lab using the supplied envelope. It's that simple, and guaranteed anonymous.



STEP 2: TRACK

The exploration continues here in the Genographic Project Web site where you can track your test kit, step by step, through the various stages of DNA sequencing and processing. Along the way, multimedia presentations explain how scientists actually decode the information found in the molecules of your DNA.



STEP 3: EXPLORE

When your results are ready Project Director Dr. Spencer Wells will introduce you to your earliest human relatives—the members of your specific haplogroup. You'll receive a personalized genetic analysis, including an online overview of your deep ancestral history. The analysis reveals where and when your haplogroup originated and how they lived. You'll also receive a dynamic map, specific to your lineage, on which to trace your relatives' journeys across the planet.

Learn more about the results you will receive

200,000 to 60,000 B.C.



60,000 to 55,000 B.C.



55,000 to 50,000 B.C.



50,000 to 45,000 B.C.



45,000 to 40,000 B.C.



40,000 to 35,000 B.C.



35,000 to 30,000 B.C.



30,000 to 25,000 B.C.



25,000 to 20,000 B.C.



20,000 to 15,000 B.C.



15,000 to 10,000 B.C.



10,000 to 5,000 B.C.





45-40,000 B.C.

Introduction

200,000 B.C.

60,000

50,000

40,000

30,000

20,000

10,000 B.C.



ZOOM OUT -

EUROPE

ASIA

AFRICA

NORTH PACIFIC OCEAN

SOUTH PACIFIC OCEAN

INDIAN OCEAN

AUSTRALIA

MAP TOOLS





GENETIC MARKERS

HAPLOGROUP B

First Appeared

Perhaps 60,000 years ago

Type mtDNA



[BACK TO WORLD MAP](#)

Your haplogroup likely arose on the high plains of Central Asia between the Caspian Sea and Lake Baikal. It is one of the founding East Asian lineages and, along with haplogroups *F* and *M*, comprises around three quarters of all mitochondrial lineages found there today.

Radiating out from the Central Asian homeland, haplogroup *B*-bearing individuals began migrating into the surrounding areas and quickly headed south, making their way throughout East Asia. Today haplogroup *B*



Peru



GENETIC MARKERS

HAPLOGROUP F

First Appeared

To be determined

Type mtDNA



[BACK TO WORLD MAP](#)

This haplogroup likely arose on the high plains of Central Asia between the Caspian Sea and Lake Baikal. It is one of the founding East Asian lineages and, along with haplogroups *B* and *M*, comprises around three-quarters of all mitochondrial lineages found there today.

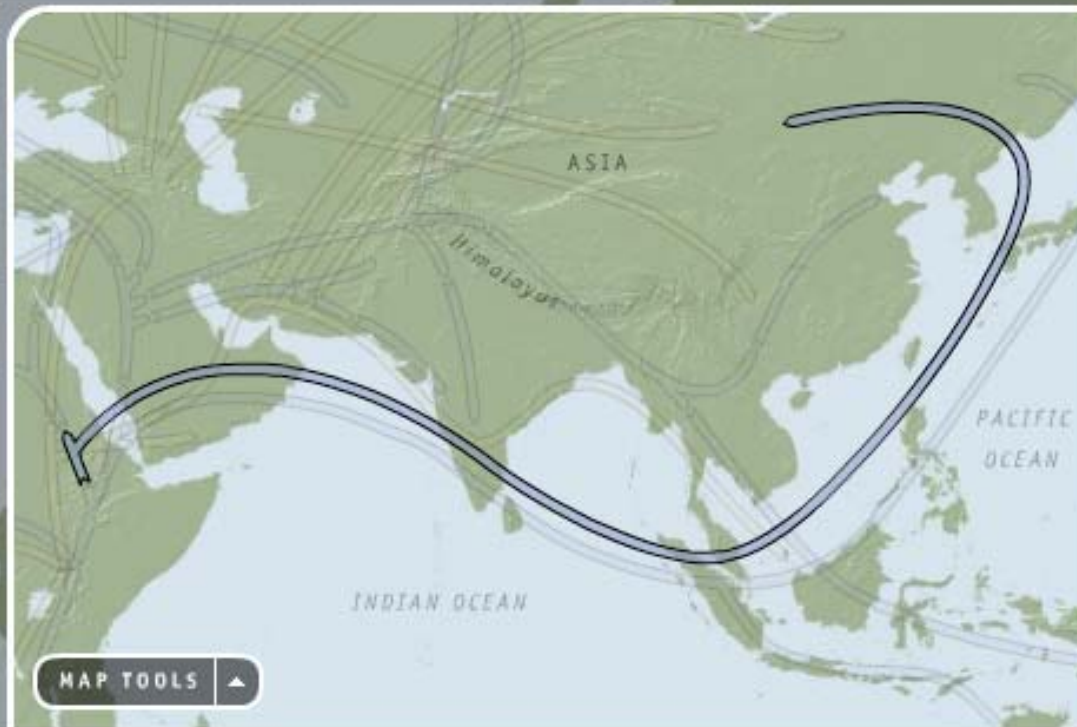
Radiating out from Central and Southeast Asia, haplogroup *F*-bearing individuals, your own distant ancestors, began migrating into the surrounding areas and quickly headed east. Today haplogroup *F* makes up over 25



China



Kashmir



GENETIC MARKERS

M174 (HAPLOGROUP D)

First Appeared

50,000 years ago

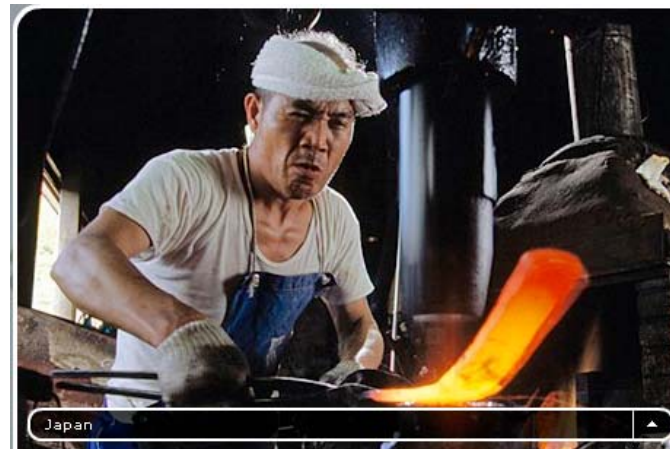
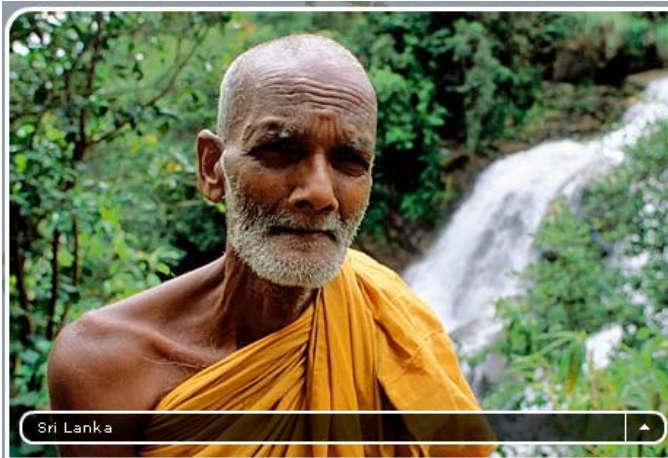
Type Y-Chromosome



Haplogroup *D* may have accompanied another group, the Coastal Clan (haplogroup *C*) on the first major wave of migration out of Africa around 50,000 years ago. Taking advantage of the plentiful seaside resources, these intrepid explorers followed the coastline of Africa through the southern Arabian Peninsula, India, Sri Lanka, and Southeast Asia.

Alternatively, they may have made the trek at a later time, following in the footsteps of the Coastal Clan. Pockets of these ancestors







GENETIC MARKERS

HAPLOGROUP Z

First Appeared

To be determined

Type mtDNA

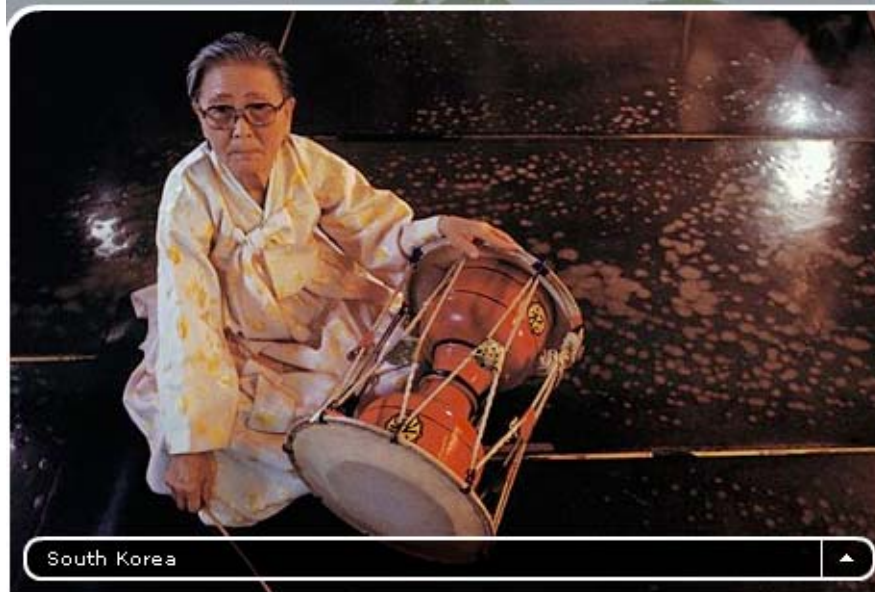


[BACK TO WORLD MAP](#)

Haplogroup Z arose on the high plains of Central Asia between the Caspian Sea and Lake Baikal. It is considered a characteristic Siberian lineage, and today accounts for around three percent of the entire mitochondrial gene pool found there. Because of its old age and frequency throughout northern Eurasia, it is widely accepted that this lineage was carried by the first humans to settle these remote areas.

Radiating out from the Siberian homeland,







GENETIC MARKERS

LLY22G (HAPLOGROUP N)

First Appeared
To be determined

Type Y-Chromosome



[BACK TO WORLD MAP](#)

One of the men in a group of Eurasian Clan peoples who traveled north through the Pamir Knot region gave rise to the LLY22G marker, which defines haplogroup N. He was probably born in Siberia within the last 10,000 years.

Today his descendants effectively trace a migration of Uralic-speaking peoples during the last several thousand years. This lineage has dispersed throughout the generations, and is now found in southern parts of Scandinavia as well as northeastern Eurasia.



Siberia, Russia



Saami man, Finland



JOURNEY HIGHLIGHTS



PAMIR KNOT

Dates

40,000 years ago

Type Natural World



[BACK TO WORLD MAP](#)

The Pamir is a high mountainous plateau about the size of Pennsylvania (45,600 square miles or 118,000 square kilometers). It is known as the Pamir "Knot" because of its location at the junction of four great Asian mountain ranges: the Himalaya, the Karakoram, the Hindu Kush, and the Tian Shan.

Even today these towering mountains pose a nearly insurmountable barrier to human travel. When Upper Paleolithic peoples first reached the area from the westward



