

# DNA and Family History

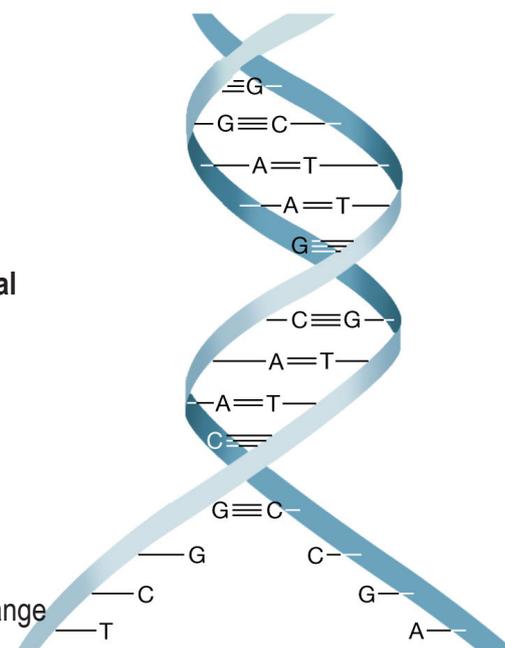
## Brief summary

### Tree Building - Genealogy Basics

- Build your tree through documents like BDMs
- Make sure you have evidence to support your tree
- DNA tests are a tool to help confirm or extend your tree

### DNA Basics (see glossary)

- We have two very different types of DNA, **Nuclear** and **Mitochondrial**
- Nuclear DNA is coiled up into 23 pairs of chromosomes
- One set came from father, the other from mother
- 22 pairs - autosomal, the other pair are sex cells (X and Y)
- Males have YX sex cells, females XX



### Inheritance

- Mitochondrial DNA passes from **mother to all children** with little change
- Y DNA passes from **father to boys only** with little change
- X DNA has a specific inheritance pathway
- Autosomal DNA is 'shuffled' during the making of sperm and egg cells
- This 'shuffling' (recombination) is random, but allows large segments of DNA to pass on unchanged

### DNA testing

- 3 key tests (Y-DNA, mt-DNA, at-DNA)
- Multiple suppliers of autosomal tests; only one does Y-DNA and mt-DNA tests
- Each supplier has large database of potential matches
- Consider 'fishing in all ponds' for matches

## The tests in summary...

TEST	USED TO –	STRENGTHS	LIMITS
<b>Y - DNA</b>	Find male relatives that share common ancestor in <b>paternal</b> line	Close matches share paternal ancestor. If close male relatives don't match, proves there's been a mistake in the tree (a NPE?)	Only males can be tested Only checks paternal line Common ancestors may have lived 100s of years ago. Doesn't 'prove' paternal relationship
<b>MT-DNA</b>	Find relatives that share common ancestor in <b>maternal</b> line	Close matches share maternal ancestor. If close female relatives don't match, proves there's been a mistake in the tree	Only checks maternal line Common ancestors may have lived many 100s of years ago Doesn't 'prove' maternal relationship
<b>AUTOSOMAL</b>	Find relatives that share common ancestor in <b>any branch</b> of your tree	Most likely will find many cousins X-matches can be useful Tools available to do significant analysis	Some relationships (3rd cousins and beyond) may not show up Hard to tell which branch Multiple companies; your relatives may not be in their databases

# Where to buy the tests...

TEST	AVAILABLE FROM	COMMENTS	COSTS (EXCHANGE RATE 1 JAN16) CHECK PRICES BEFORE YOU BUY (DISCOUNTS COMMON)
Y - DNA	FamilyTreeDNA	Number of markers is important; the more markers the more \$\$\$	Y37 = \$US169 ( <b>\$A242</b> ) Y67 = \$US268 ( <b>\$A385</b> ) Y111 = \$US359 ( <b>\$A515</b> )
MT-DNA	FamilyTreeDNA	Full sequence test can be very helpful	Full = \$US199 ( <b>\$A286</b> )
AUTOSOMAL	FamilyTreeDNA	"Family finder" test popular with UK and Australian customers	<b>\$US79 plus shipping</b>
AUTOSOMAL	Ancestry	Largest database by far (over 10 million)	<b>\$A129 plus shipping</b>
AUTOSOMAL	23andme	Many customers use it for medical information, rather than genealogy	<b>\$US99 plus shipping</b>
AUTOSOMAL	Myheritage	Recent arrival; offering discounts today	<b>\$A79 plus shipping</b>

## Test results

- Each site will display 'matches' with an estimate of closeness
- Each site has tools to help understand and analyse results
- Each site gives an estimate of 'ethnicity' – not helpful for genealogy
- You can download your 'raw data' from each site
- You can upload your raw data to other sites with different analytical tools

## Useful websites

- International Society of Genetic Genealogy - good for background, great Wiki
- Louise Coakley (Genie1) - Australian, also has a great Blog
- **FamilyTreeDNA** - testing company: only source of Y-DNA and mt-DNA tests
- **Ancestry** - testing company, source of genealogical info and published trees
- **23andme** - testing company, mostly focused on health
- **MyHeritage** - testing company, source of genealogical info
- **Gedmatch** - upload your raw data: free analytical tools (need to pay for Tier One tools)
- **DNA Land** - upload your raw data: free analytical tools
- **DNAGedcom** - upload Gedmatch processed data: free ADSA report
- **DNAPainter** – upload your raw data: free mapping tools
- <http://isogg.org>
- <http://genie1.com.au/>
- <https://www.familytreedna.com>
- <http://dna.ancestry.com.au>
- <https://www.23andme.com/en-int/>
- <https://www.myheritage.com>
- <http://v2.gedmatch.com/login1.php>
- <https://dna.land>
- <https://dnagedcom.com>
- [dnainter.com](http://dnainter.com)

# Glossary of useful terms

(adapted from John Chandler's glossary, Edmund Rice (1638) Association, <http://www.edmund-rice.org/dnagloss.htm>)

<b>allele</b>	variant of a <a href="#">gene</a> or genetic <a href="#">marker</a> . For <a href="#">STR</a> markers, each allele is designated by the number of repeats of the short <a href="#">base</a> sequence.
<b>autosome</b>	one of the non-sex-determining <a href="#">chromosomes</a> . Autosomes occur in nearly identical pairs. See also <a href="#">X</a> and <a href="#">Y</a> .
<b>base</b>	the building block of <a href="#">DNA</a> , one of four molecules that link up to make a DNA chain. The four are named adenine, cytosine, guanine, and thymine (A, C, G, T for short). These are also called <a href="#">nucleotides</a> .
<b>centiMorgan</b>	a unit of <a href="#">recombinant</a> frequency which is used to measure <a href="#">genetic distance</a> . It is often used to imply distance along a <a href="#">chromosome</a> , and takes into account how often recombination occurs in a region. A region with few cMs undergoes relatively less recombination.
<b>chromosome</b>	one of the <a href="#">DNA</a> macromolecules found in the cell nucleus. Humans have 46 chromosomes. See also <a href="#">X</a> and <a href="#">Y</a> .
<b>CRS</b>	the Cambridge Reference Sequence for <a href="#">mtDNA</a> , used for convenience to compare with all human mtDNA sequences, such that any sequence can be expressed concisely as a relatively short list of difference from the reference
<b>DNA</b>	deoxyribonucleic acid. The chemical constituent of genes and chromosomes. DNA has four different <a href="#">base</a> units, designated A, C, G, and T, which are connected in long double chains, and the sequence of these bases encodes the genetic information.
<b>DYS</b>	<a href="#">DNA Y-chromosome</a> Segment. A label for <a href="#">loci</a> or genetic <a href="#">markers</a> on the Y chromosome. Each marker is designated by a number, according to international conventions. Virtually all DYS designations discussed in the context of genetic genealogy are those of <a href="#">STR</a> markers (which are useful for genealogy because of their relatively high <a href="#">mutation</a> rate).
<b>gene</b>	a meaningful sub-unit of <a href="#">DNA</a> , encoding a protein.
<b>gene conversion</b>	a process replacing one <a href="#">allele</a> of a pair with a copy of the other. This term is used by extension for all kinds of <a href="#">loci</a> , not just <a href="#">genes</a> . See also <a href="#">recLOH</a> .
<b>genome</b>	the entire inventory of nuclear <a href="#">DNA</a> in an organism.
<b>genotype</b>	a set of <a href="#">allele</a> values of one or more <a href="#">markers</a> for one individual. Since most of the <a href="#">DNA</a> consists of nearly identical pairs of <a href="#">chromosomes</a> , a genotype typically has two alleles for each marker, one from each parent. See also <a href="#">haplotype</a> .
<b>haplogroup</b>	a classification comprising many different <a href="#">haplotypes</a> thought to be related. Haplogroups are defined in terms of <a href="#">markers</a> that mutate so slowly they are treated as if they have occurred only once in all time. The major haplogroups originated thousands or tens of thousands of years ago.
<b>haplotype</b>	a set of <a href="#">allele</a> values of one or more <a href="#">markers</a> for one individual and identified as coming just from one parent, particularly markers on the <a href="#">Y chromosome</a> or in <a href="#">mitochondrial</a> DNA. See also <a href="#">genotype</a> .
<b>heterozygous</b>	having two different <a href="#">alleles</a> in the two copies of the same <a href="#">marker</a> or <a href="#">locus</a> existing on a pair of <a href="#">chromosomes</a> . By extension, this term is sometimes applied to the two copies of a locus found on opposite arms of a <a href="#">palindrome</a> on the <a href="#">Y</a> chromosome. See also <a href="#">homozygous</a> .
<b>homozygous</b>	having the same <a href="#">allele</a> in the two copies of the same <a href="#">marker</a> or <a href="#">locus</a> existing on a pair of <a href="#">chromosomes</a> . By extension, this term is sometimes applied to the two copies of a locus found on opposite arms of a <a href="#">palindrome</a> on the <a href="#">Y</a> chromosome. See also <a href="#">heterozygous</a> .
<b>HVR1</b>	Hypervariable Region 1. A portion of the <a href="#">mtDNA</a> molecule noted for its especially high <a href="#">mutation</a> rate, consisting roughly of locations 16024-16569.

<b>HVR2</b>	Hypervariable Region 2. (See <a href="#">HVR1</a> .) The limits of HVR2 are even more vague than for HVR1. HVR2 is generally said to start at location 1 and to extend for a few hundred <a href="#">bases</a> , but part of this region is often called HVR3.
<b>locus</b>	(plural: loci) specific site on a <a href="#">DNA</a> chain.
<b>marker</b>	a distinctive sub-unit of <a href="#">DNA</a> , often not part of a <a href="#">gene</a> . Often used interchangeably with <a href="#">locus</a> , but referring to the <b>contents</b> of the site, rather than the <b>site</b> itself.
<b>microsatellite</b>	See <a href="#">STR</a> .
<b>mitochondria</b>	<a href="#">organelles</a> within the cell responsible for converting food into usable energy. Each mitochondrion has its own <a href="#">DNA</a> . The mitochondria in a child come entirely from the mother, and so mitochondrial DNA ("mtDNA" for short) is a tracer of female-line ancestry, just as <a href="#">Y</a> DNA is a tracer for male-line ancestry.
<b>mtDNA</b>	See <a href="#">mitochondria</a> .
<b>mutation</b>	An event in which the <a href="#">DNA</a> chain alters. In the case of <a href="#">STR</a> markers, a mutation is almost invariably the gain or loss of one repeat of the basic short sequence (or, rarely, two repeats). Other types of mutations include the substitution of one <a href="#">base</a> for another (known as a <a href="#">SNP</a> ) and the insertion or deletion of a whole segment (known as an <a href="#">indel</a> ). See also <a href="#">recombination</a> .
<b>nucleotide</b>	a unit of <a href="#">DNA</a> . See also <a href="#">base</a> .
<b>organelle</b>	any small, compact, and cohesive entity within a cell. An organelle is to a cell what an organ is to a multi-celled creature.
<b>PCR</b>	Polymerase Chain Reaction. A chemical process that replicates a given sample of <a href="#">DNA</a> many times, in imitation of natural replication. The process cycles between two stages: splitting the two strands of DNA apart and then forming new double strands by adding a mixture of the enzyme polymerase and the four DNA <a href="#">bases</a> . By adding <a href="#">primers</a> as well, the process can be used to replicate just the one or more DNA segments of interest.
<b>polymorphism</b>	The occurrence of more than one form of <a href="#">DNA</a> in different individuals, or even in different cells within one individual. Such diversity arises through the occurrence of <a href="#">mutations</a> .
<b>recLOH</b>	<a href="#">recombinant</a> loss of <a href="#">heterozygosity</a> . See also <a href="#">gene conversion</a> .
<b>recombination</b>	a process of "mixing and matching" of paired <a href="#">chromosomes</a> that takes place at cell division. One or more segments may be swapped between the two chromosomes, or occasionally a segment may replace the corresponding segment on the other chromosome. This process can also occur on <a href="#">palindromic</a> segments of the <a href="#">Y</a> chromosome and may affect paired <a href="#">loci</a> , such as <a href="#">DYS385a</a> and <a href="#">DYS385b</a> .
<b>SNP (pronounced "snip")</b>	Single- <a href="#">Nucleotide Polymorphism</a> . A type of <a href="#">mutation</a> characterized by the substitution of one <a href="#">base</a> for another, or the outright loss of a <a href="#">base</a> .
<b>STR</b>	Short Tandem Repeat. Also known as microsatellite. This is a genetic <a href="#">marker</a> consisting of multiple copies of a short <a href="#">motif</a> , (a sequence of <a href="#">DNA bases</a> ). Occasionally, a microsatellite will mutate by the gain or loss of one repeat. So-called "simple" STRs have just one contiguous set of repeats; "complex" STRs may have multiple sets of repeats separated by short patches of non-repeating DNA and may even have repeats of more than one motif.
<b>X</b>	one of the two sex-determining <a href="#">chromosomes</a> , See also <a href="#">Y</a> .
<b>Y</b>	one of the two sex-determining <a href="#">chromosomes</a> . A person with the combination XX is female, while a person with XY is male. Most of the Y chromosome, unlike the others, does not trade <a href="#">DNA</a> with a "partner chromosome," and it therefore passes essentially intact from father to son. This property leads to a minimum of ambiguity in interpreting the results of Y DNA analysis.