

# **DNA and its Place in Family History Research -Introduction-**

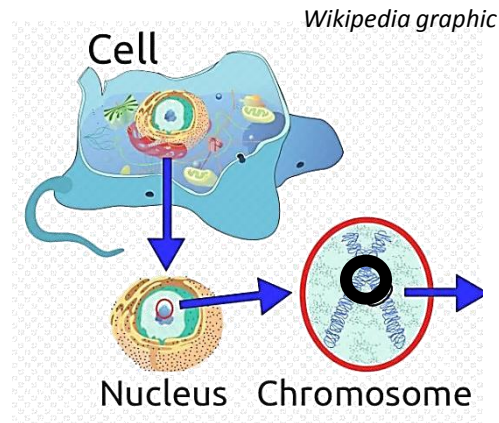
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***07 May 2022***

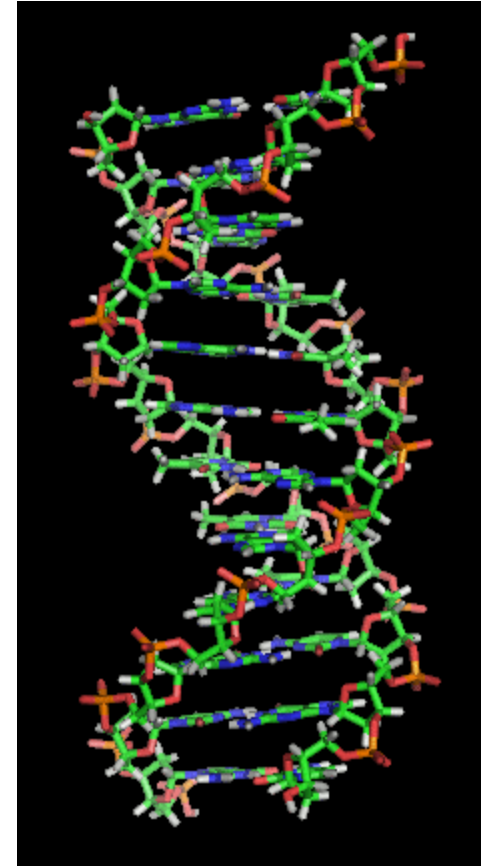
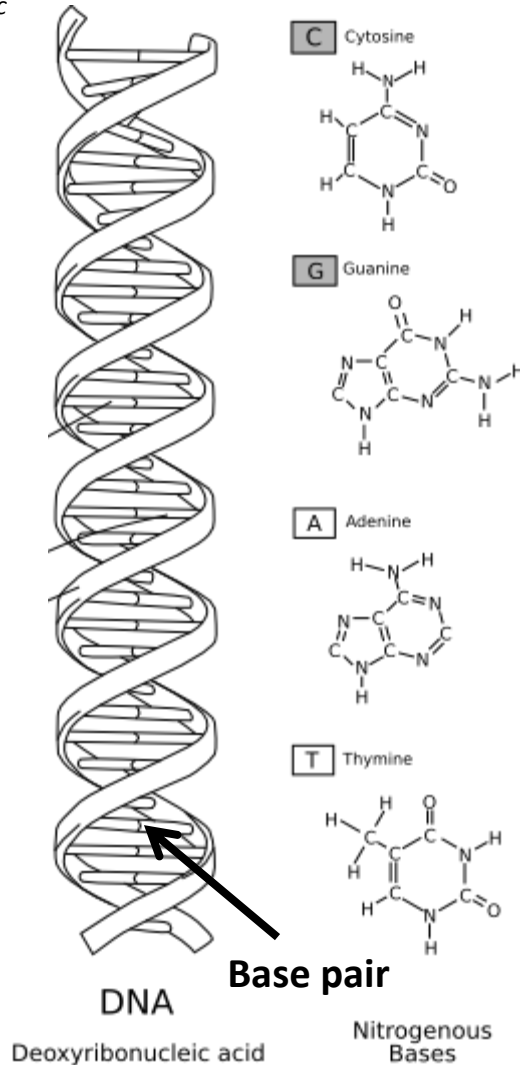
# Outline

- **What is DNA?**
- **DNA Terms (test result reporting and social)**
- **DNA in Genealogy Research**
- **DNA Testing**
  - SNP mapping examples
  - Test types in use today
  - DNA testing companies & comparison
  - Can company databases be compared?
  - Consanguinity & Correspondence with DNA
- **Conclusion**
- **Triangulation Examples (recent)**

# What is DNA? (1)

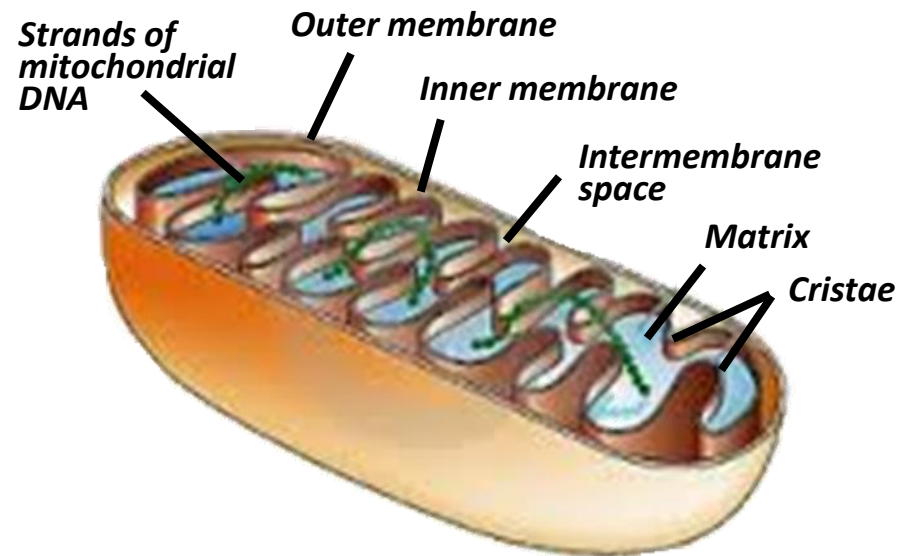


- DNA is what makes up each of the 23 Chromosome pairs within each nucleus in each cell in our bodies
- A pairs with T, C pairs with G



# What is DNA? (2)

- Within the cell nucleus:
  - 22 Chromosome pairs (autosomes)
  - 1 sex chromosome pair: (“normally” XY = male, XX = female)
    - ✓ Males get their Y from their father; X from mother
    - ✓ Females get an X from mother and father (crossover can occur)
- We also have mitochondria in our cells which have DNA (mtDNA)
  - we get all our mtDNA from our mothers



***A Mitochondrion within a cell***

<http://www.rajaha.com/description-structure-mitochondria-function>

# DNA Terms – Testing (1)

- MRCA – Most Recent Common Ancestor
  - “Estimated number of generations to MRCA”
- Y-STR – Standard Marker test for short tandem repeats (STRs) of a sequence in a region on the Y-Chromosome; “value” stated is the allele count for the marker (usually # repeats for simple markers)
  - *Two men matching 111 Y-STR markers may still be 8<sup>th</sup> to 9<sup>th</sup> cousins*
- SNP - Single Nucleotide Polymorphism (pronounced “snip”) are single mutations in a short sequences of base pair values within a section on a chromosome; e.g.: AAGCCTA to AAGCITA
  - *SNP testing is goal of large scale Genome testing*

# DNA Terms – Testing (2)

- **Haplotype** - A collection of SNPs per chromosome or a collection of Y-STR values (for Y-DNA analysis) that tend to be inherited together
- **Haplogroup** – a group of similar haplotypes with a common ancestor determined by specific SNPs; Primarily defined for paternal (Y-DNA) and maternal (mtDNA) descendance, e.g., “*Adam*” starts the paternal “A” haplogroup and “*Eve*” starts the maternal “L” haplogroup.
- **Illumina DNA sequencing/genotyping chips:**
  - AncestryDNA: Omniexpress v2 to 2019? Now GSA?
  - 23andMe: Global Screen Array (GSA) v5 Aug 2017+...
  - FamilyTreeDNA: changed to GSA April 2019 (they process MyHeritage kits as well)

# DNA Terms – Testing (3)

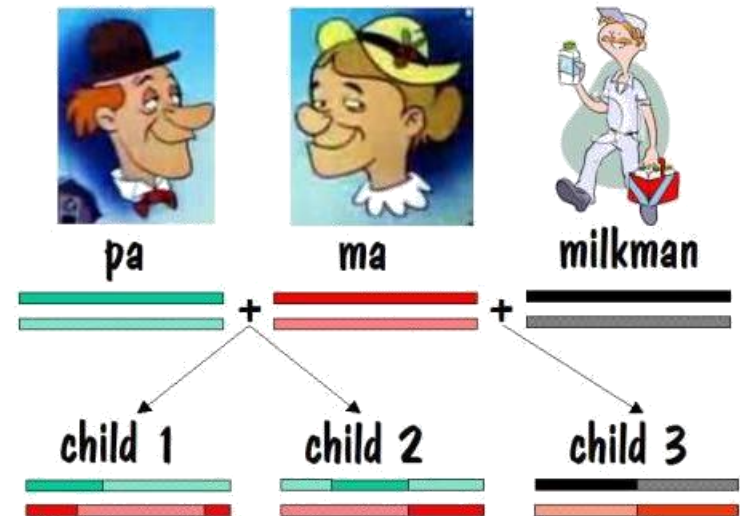
- **cM (centiMorgan)** – in Autosomal DNA research, a measure of how frequently a DNA segment is exchanged , or *recombined*, when a sperm/egg is created. Humans have 46 chromosomes in 23 pairs. When males produce sperm and women produce eggs, only 23 chromosomes (half) are passed to the sperm/egg by each parent. The child produced will receive 23 **recombined** chromosome segments from each parent. The length of each segment passed from the parent to the sperm/egg may be complete, half, etc..., or zeroed out completely.

***In practical terms, using current technology, there are ~7500 cM available to be shared (ancestryDNA says to use ~7000 cM for theirs) (e.g., full siblings share about 50% or 3750 cM)***

# DNA Terms - Social

- **NPE** – Non-Paternal Event; surname of child does not match Y-Chromosome biology – adoption, unmarried mother passing maiden name, extra-marital affair, etc... Estimate is 5-10% likelihood each generation
- **OOW** – Out Of Wedlock; child born to unmarried mother
  - 18% of US births in 1980, 30% 1990, ~40% 2007-2015  
<https://www.cdc.gov/nchs/data/databriefs/db162.htm>

*(NOTE: DNA Testing companies have a disproportionate number of NPE/OOW individuals in their databases!! – how much more than ~5-10% ??? )*



<http://www.stevemorse.org/genetealogy/beyond.htm>

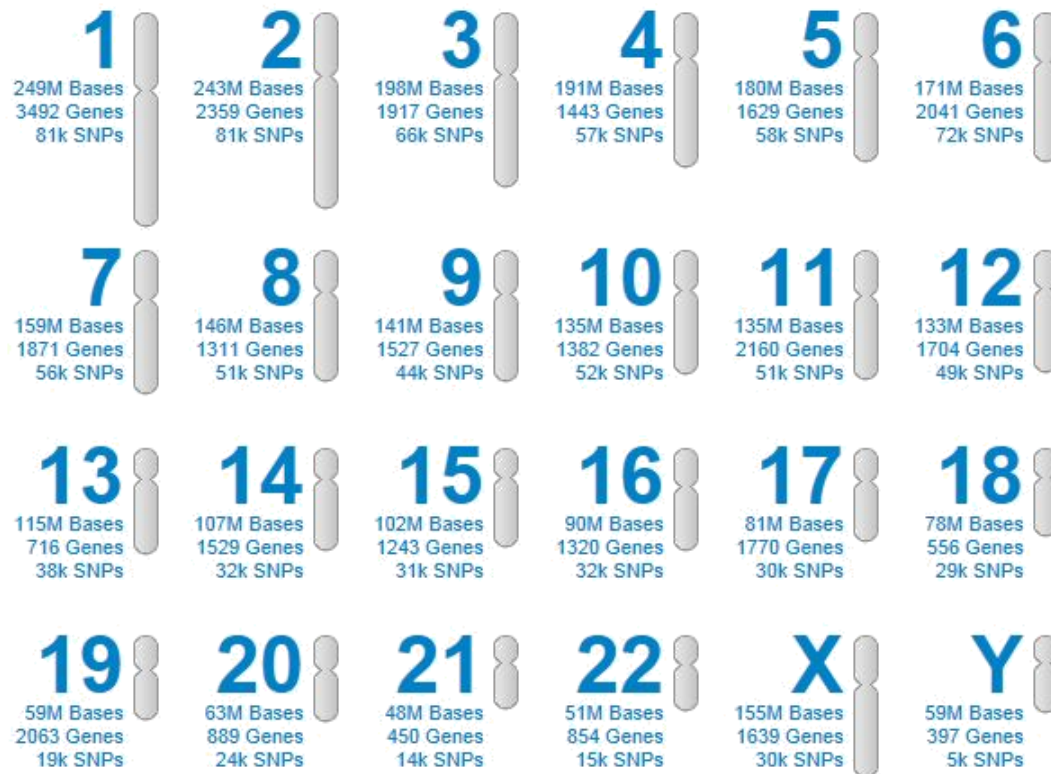


# DNA in Genealogy Research

- DNA directly stores information about our ancestry
  - Our chromosomes are a combination of our parents' chromosomes; their chromosomes are combinations of their parents, etc.
  - The combination is random (see cM description), however certain regions '*usually*' get passed down as a unit (genes are an example); Note: if a segment gets passed down exactly over multiple generations – MRCA wrong!
  - Mutations can occur during the copying/replicating process producing sperm/egg– this is critical to DNA testing!
    - ✓ Y Chromosome mutations: STR and SNP tests
    - ✓ Autosome and X chromosome mutations: SNP tests

# Chromosome SNPs (*pronounced “snips”*)

- Single Nucleotide Polymorphisms (SNPs) are short sequences of base pair values within a section on a chromosome



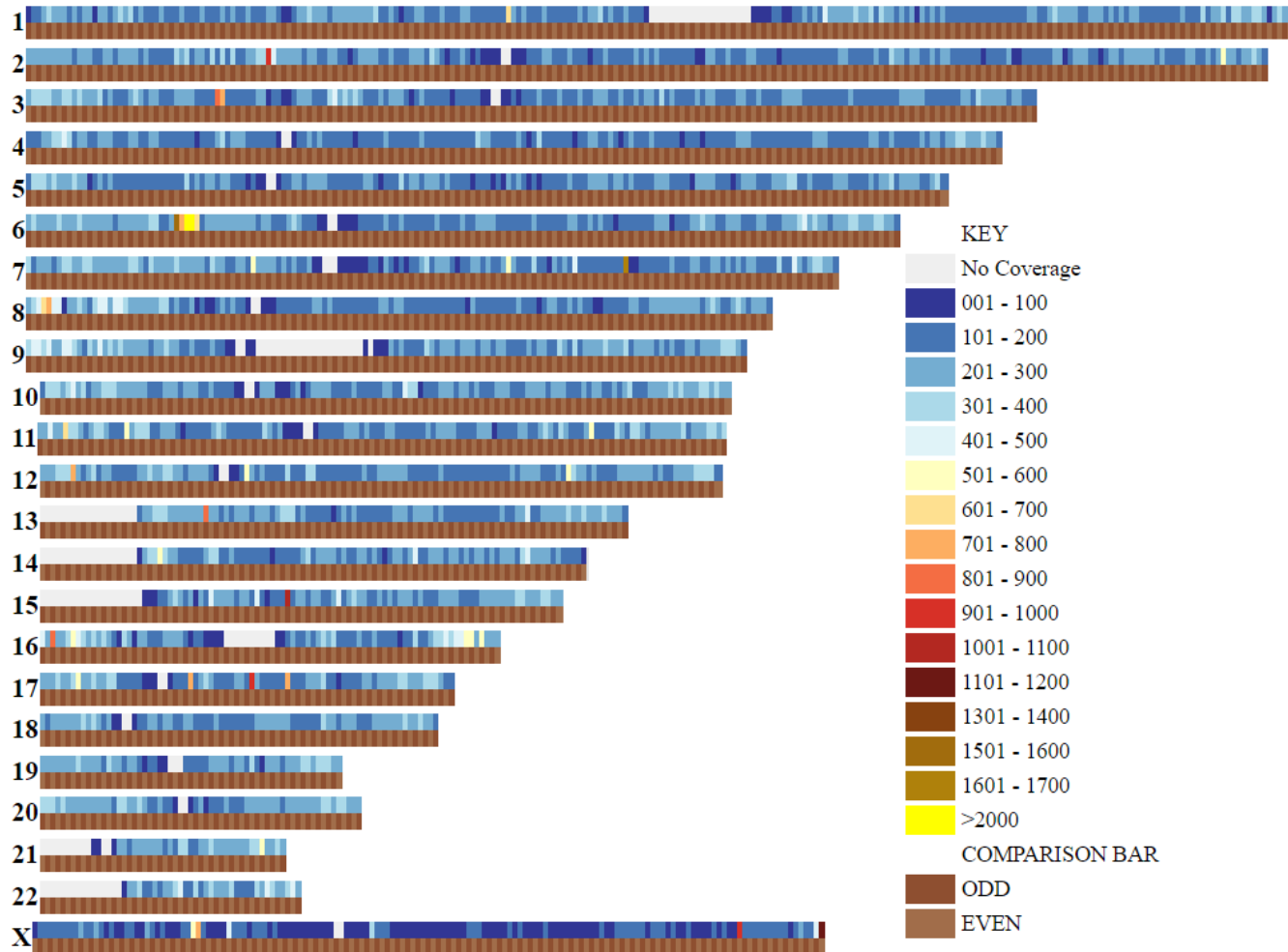
**MT** ○  
16k Bases  
37 Genes  
4k SNPs

Source: [23andme.com](http://23andme.com) (note: #SNPs is for older V3 chip)

Total Human Genome: ~3 Billion base pairs

# DNA Terms – SNP example

- Single Nucleotide Polymorphisms (SNPs) : 23andMe V4 chip (~602K SNPs)



Source: <http://haplogroup.org/23andme-v4-chip/>

Ranges are SNP counts per 1M base pairs; Total Human Genome: ~3B base pairs 11

# DNA Testing Types - Current

- **Y-DNA STR and SNP testing** can help determine ***paternal*** ancestry over many generations *for males only*
  - *STR testing usually done first with a standard set of markers; specific Y-SNP testing usually done next*
  - *Big-Y, Y Elite, ...*
- **mtDNA SNP testing** can determine ***maternal*** ancestry over many generations *for both males and females*
  - *due to low mutation rate, not very useful within 10+ generations*
- **Autosomal testing** (and actually X sex chromosome too), allows *both males and females* to determine ***relationship with ~6-7 generations***
  - *Current technology, i.e., number of DNA locations sampled and min segment length considered significant, limits the ability to match others to ~4<sup>th</sup>-5<sup>th</sup> cousinship at best*
- **Whole/Full Genome testing** (*how to handle 100 to 300 Gb data!*)
  - *Below \$1K, Nebula Genomics will do 30x sequencing depth for \$199!*

# Major DNA Testing Companies

- [FamilyTreeDNA.com](https://www.familytreeDNA.com) (FTDNA) - Y-STR (37/~~67~~/111), Y-SNP (Big-Y/Individual SNPs), mtDNA (~~HVR1+HVR2~~/Full), autosomal V3 chip (Family Finder – no Y-SNP), National Geographic Genographic Project
- [23andme.com](https://www.23andme.com) – autosomal “customized” GSA V5 chip (includes X, Y, and mtDNA SNP testing)
- [dna.ancestry.com](https://www.dna.ancestry.com) – autosomal; Blackstone private equity. Note: now, each kit must be registered under its own unique account (e-mail).
- [myheritagedna.com](https://www.myheritagedna.com) – autosomal, Israel based (not available in Israel(!), France, Poland, Alaska); V3 chip

# DNA Company Comparison- Autosomal

Company	23andMe ↕	Family Tree DNA's Family Finder test ↕	Ancestry.com's AncestryDNA test ↕
Website	<a href="http://www.23andme.com">www.23andme.com</a> <a href="#">↗</a> (56 countries)	<a href="http://www.familytreedna.com">www.familytreedna.com</a> <a href="#">↗</a> (worldwide)	<a href="http://dna.ancestry.com">http://dna.ancestry.com</a> <a href="#">↗</a> (~35 countries)
Price (as of May 2022)	US: \$99 ancestry; \$199 +health; xtras \$30/yr	\$79	\$99 (routinely reduced to \$59 now) in the U.S. (subscription required to access some features)
Upload of raw data file allowed from other companies	No	23andMe V3, V4 & V5; ancestryDNA Omniexpress V2 &? newer GSA? ; MyHeritage	No
SNP chip used for testing	Illumina GSA "V5"	Illumina Omniexpress V3, now GSA?	Illumina GSA (as of late 2019?)
Number of autosomal SNPs tested	630,132 V5 (pre-Dec 2013 ~967K V3, pre-Aug 2017 ~577K V4)	708,092 ← V3 chip – ~612K filtered	682,549 ← V2 chip; ~637K filtered
Number of Y chromosome SNPS	3733 V5 (pre-Dec 2013 3089 V3, pre-Aug 2017 2329 V4)	None	1691 (labeled chromosome 24)
Number of X chromosome SNPS	16,530 V5 (pre-Dec 2013 26,087 V3, pre-Aug 2017 19,487 V4)	18,091	28,892 (labeled as chromosome 23), plus 440 SNPs labeled as chromosome 25 that are either from chromosome X or from the pseudoautosomal regions of the Y chromosome
Number of mitochondrial DNA SNPS	4318 V5 (pre-Dec 2013 2737 V3, pre-Aug 2017 3154 V4)	None	263 (2022), 195(2018), used to be none
Number of people in the database (as of Feb 2019)	12,200,000+	~1,800,000	21,000,000+ -----

[www.myheritagedna.com](http://www.myheritagedna.com) also does testing (~5.6M people!!) – based in Israel, but not allowed in Israel ?

From: [http://www.isogg.org/wiki/Autosomal\\_DNA\\_testing\\_comparison\\_chart](http://www.isogg.org/wiki/Autosomal_DNA_testing_comparison_chart)

# DNA Testing - Comparing Databases

- Comparison web sites – Autosomal:
  - [www.GEDmatch.com](http://www.GEDmatch.com) – upload and compare autosomal data (T# (randomized #) = FTDNA FF; M# = 23andme (Note: V5 chip new testing requires a different algorithm); A# = AncestryDNA)
  - FTDNA will accept the older 23andme.com V3 (Nov 2010 - Nov 2013) raw data and they will accept AncestryDNA – into their database. Also MyHeritage because they process it.
- Comparison web sites – Y Chromosome:
  - Y-SNP & whole genome: [www.yfull.com](http://www.yfull.com) – accepts [www.familytreedna.com](http://www.familytreedna.com) Big-Y500 and Big-Y700; [www.fullgenomes.com](http://www.fullgenomes.com) Y-Elite; Nebula [www.nebula.org](http://www.nebula.org)
- Other helpful site:
  - [www.isogg.org](http://www.isogg.org) – International Society of Genetic Genealogy
  - (many, many others)

# Consanguinity (Blood Relationship)

Common Ancestor	Child	Grandchild	Great Grandchild	2 <sup>nd</sup> Great Grandchild	3 <sup>rd</sup> Great Grandchild	4 <sup>th</sup> Great Grandchild	5 <sup>th</sup> Great Grandchild	6 <sup>th</sup> Great Grandchild	7 <sup>th</sup> Great Grandchild
Child	Sibling	Niece/Nephew	Grandniece / Grandnephew	Great Grandniece / Grandnephew	2 <sup>nd</sup> Great Grandniece / Grandnephew	3 <sup>rd</sup> Great Grandniece / Grandnephew	4 <sup>th</sup> Great Grandniece / Grandnephew	5 <sup>th</sup> Great Grandniece / Grandnephew	6 <sup>th</sup> Great Grandniece / Grandnephew
Grandchild	Niece/Nephew	First Cousin	First Cousin, 1r	First Cousin, 2r	First Cousin, 3r	First Cousin, 4r	First Cousin, 5r	First Cousin, 6r	First Cousin, 7r
Great Grandchild	Grandniece / Grandnephew	First Cousin, 1r	Second Cousin	Second Cousin, 1r	Second Cousin, 2r	Second Cousin, 3r	Second Cousin, 4r	Second Cousin, 5r	Second Cousin, 6r
2 <sup>nd</sup> Great Grandchild	Great Grandniece / Grandnephew	First Cousin, 2r	Second Cousin, 1r	Third Cousin	Third Cousin, 1r	Third Cousin, 2r	Third Cousin, 3r	Third Cousin, 4r	Third Cousin, 5r
3 <sup>rd</sup> Great Grandchild	2 <sup>nd</sup> Great Grandniece / Grandnephew	First Cousin, 3r	Second Cousin, 2r	Third Cousin, 1r	Fourth Cousin	Fourth Cousin, 1r	Fourth Cousin, 2r	Fourth Cousin, 3r	Fourth Cousin, 4r
4 <sup>th</sup> Great Grandchild	3 <sup>rd</sup> Great Grandniece / Grandnephew	First Cousin, 4r	Second Cousin, 3r	Third Cousin, 2r	Fourth Cousin, 1r	Fifth Cousin	Fifth Cousin, 1r	Fifth Cousin, 2r	Fifth Cousin, 3r
5 <sup>th</sup> Great Grandchild	4 <sup>th</sup> Great Grandniece / Grandnephew	First Cousin, 5r	Second Cousin, 4r	Third Cousin, 3r	Fourth Cousin, 2r	Fifth Cousin, 1r	Sixth Cousin	Sixth Cousin, 1r	Sixth Cousin, 2r
6 <sup>th</sup> Great Grandchild	5 <sup>th</sup> Great Grandniece / Grandnephew	First Cousin, 6r	Second Cousin, 5r	Third Cousin, 4r	Fourth Cousin, 3r	Fifth Cousin, 2r	Sixth Cousin, 1r	Seventh Cousin	Seventh Cousin, 1r
7 <sup>th</sup> Great Grandchild	6 <sup>th</sup> Great Grandniece / Grandnephew	First Cousin, 7r	Second Cousin, 6r	Third Cousin, 5r	Fourth Cousin, 4r	Fifth Cousin, 3r	Sixth Cousin, 2r	Seventh Cousin, 1r	Eighth Cousin



# Consanguinity vs. % Autosomal DNA

- % Autosomal DNA in common between two people:
  - *NOTE: statistics are +/- these values; e.g., Siblings share ~40-~60% DNA (half-siblings are half this); 1<sup>st</sup> Cousins share ~7% to ~18%; 2C are ~1% to ~5%; 3C ~0% to ~2% (10-15% of 3<sup>rd</sup> cousins don't share any DNA!)*

Start	parent	grandparent	1st great-grandparent	2nd great-grandparent	3rd great-grandparent	4th great-grandparent	5th great-grandparent	6th great-grandparent	7th great-grandparent
parent	Sibling 50.000%	25.000%	12.500%	6.250%	3.125%	1.563%	0.781%	0.391%	0.195%
grandparent	25.000%	1 <sup>st</sup> Cousin 12.500%	6.250%	3.125%	1.563%	0.781%	0.391%	0.195%	0.098%
1st great-grandparent	12.500%	6.250%	2 <sup>nd</sup> Cousin 3.125%	1.563%	0.781%	0.391%	0.195%	0.098%	0.049%
2nd great-grandparent	6.250%	3.125%	1.563%	3 <sup>rd</sup> Cousin 0.781%	0.391%	0.195%	0.098%	0.049%	0.024%
3rd great-grandparent	3.125%	1.563%	0.781%	0.391%	4 <sup>th</sup> Cousin 0.195%	0.098%	0.049%	0.024%	0.012%
4th great-grandparent	1.563%	0.781%	0.391%	0.195%	0.098%	5 <sup>th</sup> Cousin 0.049%	0.024%	0.012%	0.006%
5th great-grandparent	0.781%	0.391%	0.195%	0.098%	0.049%	0.024%	6 <sup>th</sup> Cousin 0.012%	0.006%	0.003%
6th great-grandparent	0.391%	0.195%	0.098%	0.049%	0.024%	0.012%	0.006%	7 <sup>th</sup> Cousin 0.003%	0.002%
7th great-grandparent	0.195%	0.098%	0.049%	0.024%	0.012%	0.006%	0.003%	0.002%	8 <sup>th</sup> Cousin 0.001%

# Conclusion

- Y-DNA can be used to determine paternal ancestry via comparison with other males
  - watch-out for NPEs! – surname may be different!
- Autosomal DNA can be used to verify/validate close (~5<sup>th</sup> cousin or better) relationships; can be used for paternity/maternity/sibling verification
- Records based Genealogy still very important!
  - *DNA cannot really stand on its own; would need many, many samples to do so*
- DNA may be only hope for adoption/illegitimacy/etc. to obtain biological ancestry

# NPE Search Examples Using Triangulation

- If *paternal* line is unknown: Is *tester* :
  - Male? : 1<sup>st</sup>: Y-DNA to hopefully determine surname; 2<sup>nd</sup> Autosomal
  - Female? (or Y-DNA inconclusive) : Autosomal test
- If *maternal* line is unknown: Autosomal test
- Goal of triangulation is to separate matches into lines known (if any) and unknown – Ancestry finally makes this “easy”

AncestryDNA cM mapping: (Ancestry claims their test results in ~7000 cM per person)

Group A: 2800-4200 cM (40%-60%; 50% ctr) -> full siblings, parent-child

Group B: 1400-2100 cM (20%-30%; 25% ctr) -> half-sibling, Aunt/Uncle-Niece/Nephew; double 1<sup>st</sup> cousin; Grandparent-Grandchild

Group C: 490-1050 cM (~7% - ~18%; 12.5% ctr) -> 1C, half Aunt/Uncle-Niece/Nephew; GG-greatgrandchild; GAunt/GUncle-GNiece/GNephew

Group D: 210-630 cM (~3%-~9%; 6.25% ctr) -> 1C1R, 2GGrandparent-2GGrandchild, half 1<sup>st</sup> cousin; Half GAunt/GUncle-GNiece/GNephew

Group E: 70 – 350 cM (~1%-~5%; 3.125% ctr) -> 2C, 1C2R, ½ 1C1R, 3GGrandparent-3GGrandchild

Group F: 35 – 210 cM (~0.5%-~3%; 1.563% ctr) -> 2C1R, ½ 2C, 1C3R, ½ 1C2R, ...

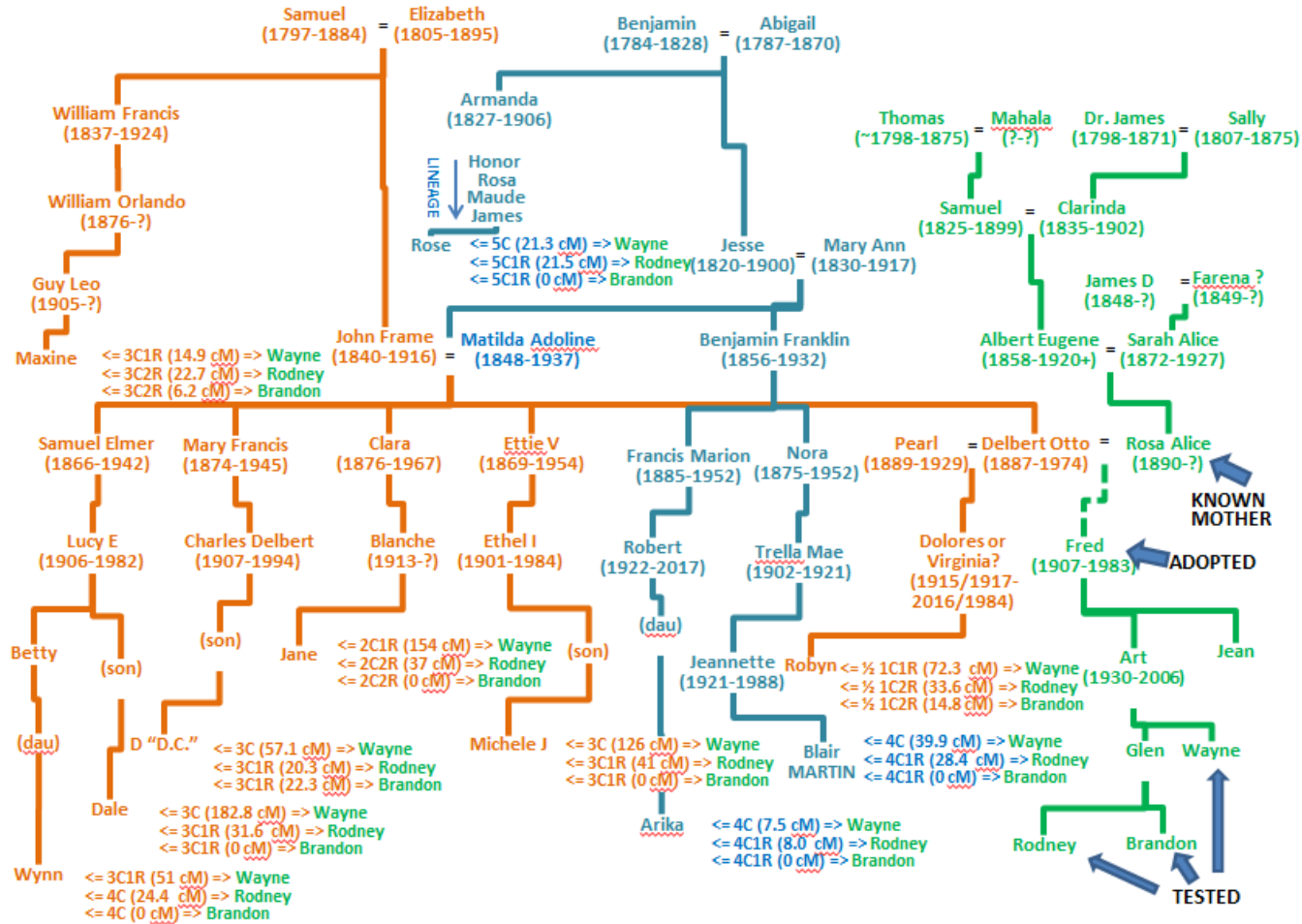
Group G: 0 – 140 cM (0% - ~2%; 0.781% ctr) -> 3C, 2C2R, 1C4R, ... (10-15% of 3<sup>rd</sup> cousins do not share any DNA)

Group H: 0 – 28 cM (0% - ~0.4%; 0.391% ctr) -> 3C1R, ... (~50% of 4<sup>th</sup> cousins do not share any DNA)

# Example: Paternal Search A (1)

- Fred was adopted in 1910 in Kansas – bio mother “known” (DNA matches not shown for her family)
- Rodney did Y-DNA and Autosomal; Brandon and Wayne did Autosomal tests
- *FTDNA Y-DNA test proved “surname” (very common name!)*
- AncestryDNA autosomal tests with quite a bit of work, proves who bio father “was” -> DNA matches on father’s father’s and mother’s sides...

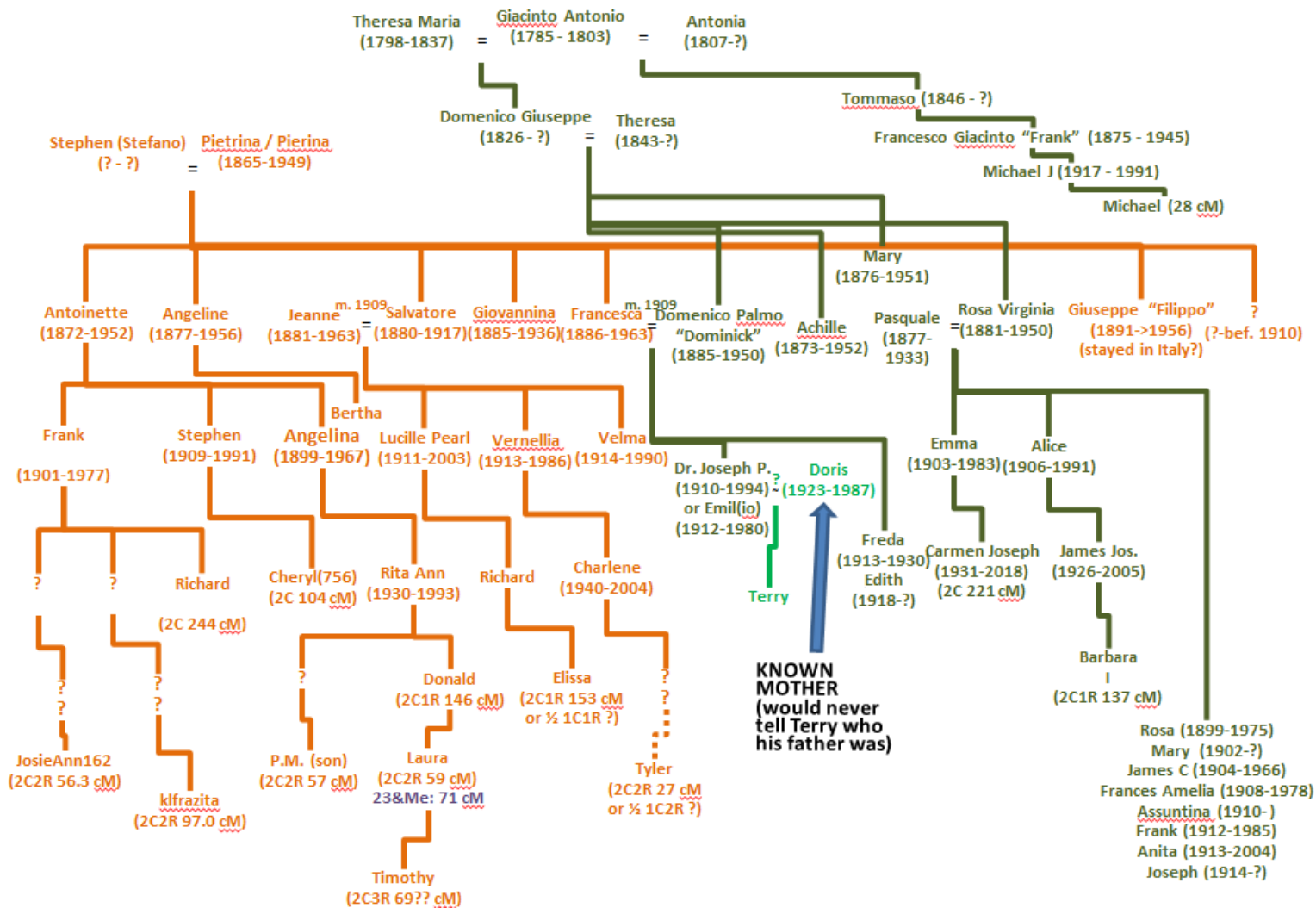
# Example: Paternal Search A (2)



# Example: Paternal Search B (1)

- Terry did Y-DNA and had his “brother” do Y-DNA and found out he wasn’t his dad’s son... (no match with “brother” and also family knowledge that he wasn’t surname)
- *Y-DNA totally inconclusive – closest matches were also NPEs. (Surname is Italian and “no male” has been tested... trying to get one to get tested!)*
- 23andMe – very confusing results
- AncestryDNA finally helped solve the mystery...at least down to one of two brothers was his father. Apparently “neither” had other children?

# Example: Paternal Search B (2)

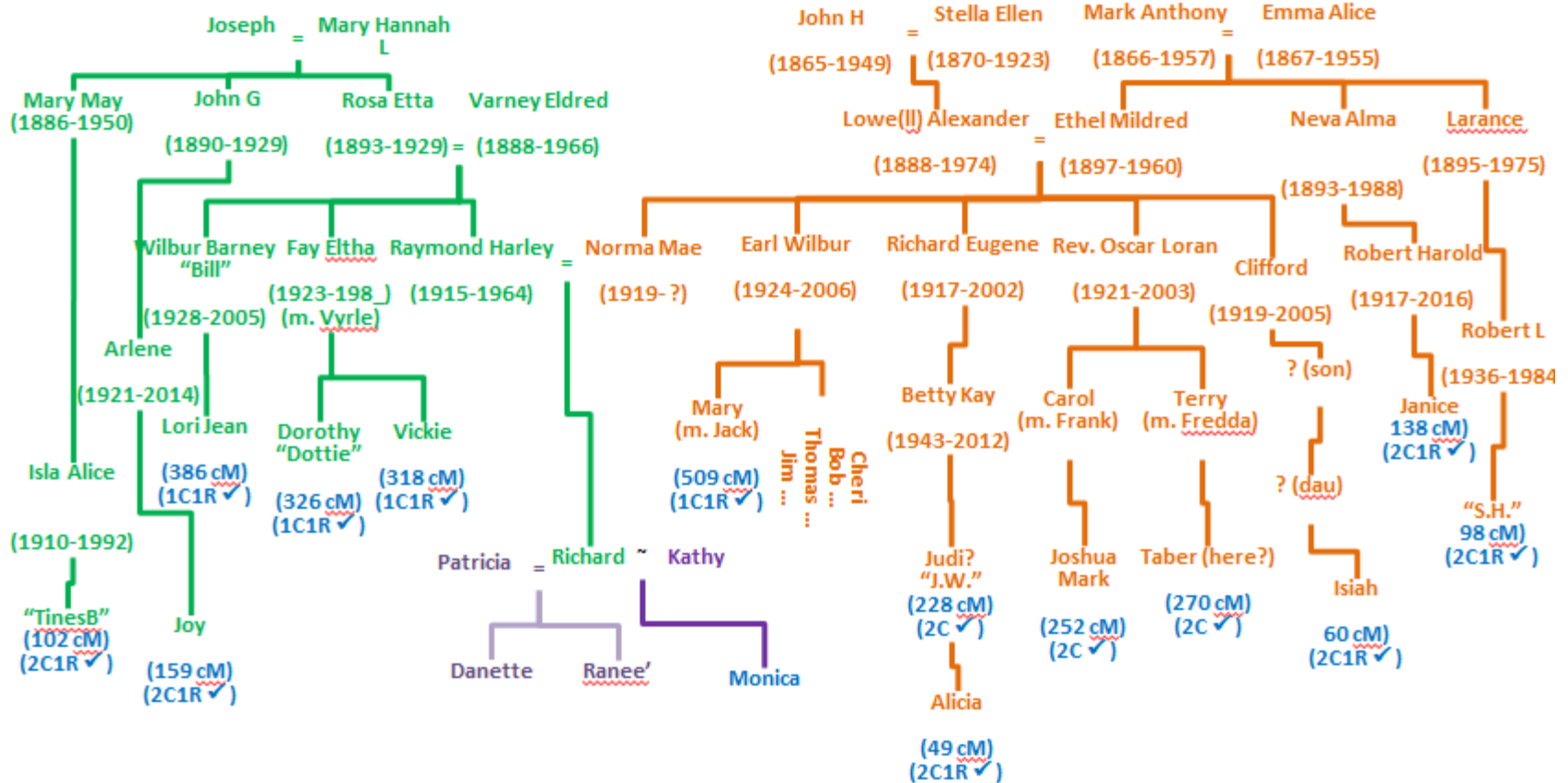


# Example: Paternal Search C (1)

- “We” have always known my niece, Monica, was not my brother’s... Finally did AncestryDNA to prove it and figure out who her bio father was. Her maternal Aunt and Uncle “know” – but family still won’t acknowledge that “we” now know who her father is..
- She is in touch with the two half-sisters we know about now... sadly, father has Parkinson’s
- Raneer’ asked how we could figure this out – without her father’s DNA !!



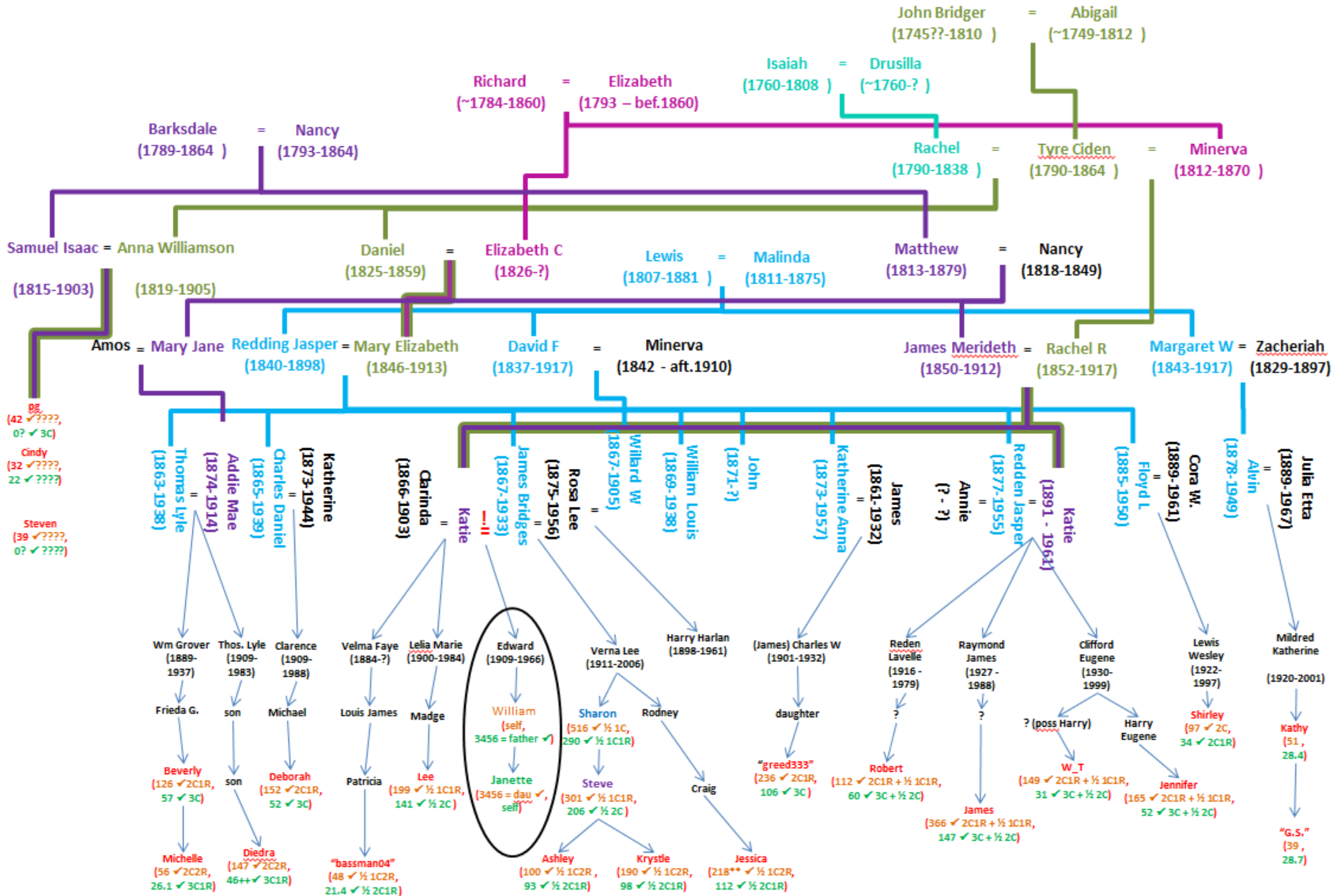
# Example: Paternal Search C (2)



# Example : Full Bio Parent Search (1)

- Very complicated – due to people related to each other multiple ways => you “add” up each cM range!
- Edward was Adopted in northern Illinois; his son killed him in 1966
- Another son, William, has a daughter, Janette, who wanted to figure out who Edward’s parents were – they were “told” never to look because if it ever came out – it would be big trouble...
- Turns out their roots are in southern Illinois...
- They have now met some of their cousins – all is well

# Example : Full Bio Parent Search (2)



# Example : Full Bio Parent Search (3)

