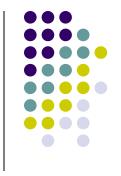
Family History Research

Using Genetic Genealogy



- DNA Testing Companies
- Three most common types of testing using DNA
 - Y-DNA
 - mtDNA
 - Autosomal DNA (atDNA) (including X-dna)
- DNA Analysis Tools

- Main testing companies to choose from:
- Family Tree DNA <u>www.familytreedna.com</u>
 - Y-dna (\$139USD), mtDNA (\$79-199USD), atDNA (\$79, includes ethnicity, not medically focussed)
 - Accepts transfers from some other testing companies (possibly free, or ~\$19USD)
- 23 and ME <u>www.23andme.com</u>
 - atDNA (\$249CDN, includes ethnicity, medically focussed)
- Ancestry <u>www.dna.ancestry.com</u>
 - atDNA (\$149CDN, includes ethnicity, not medically focussed)
- MyHeritage <u>www.myheritage.com</u>
 - atDNA \$79USD, accepts dna transfers for free
- LivingDNA www.livingdna.com
 - atDNA \$143CDN, plans to accept transfers in the near future
- *https://isogg.org/wiki/Autosomal_DNA_testing_comparison_chart (comparison details)

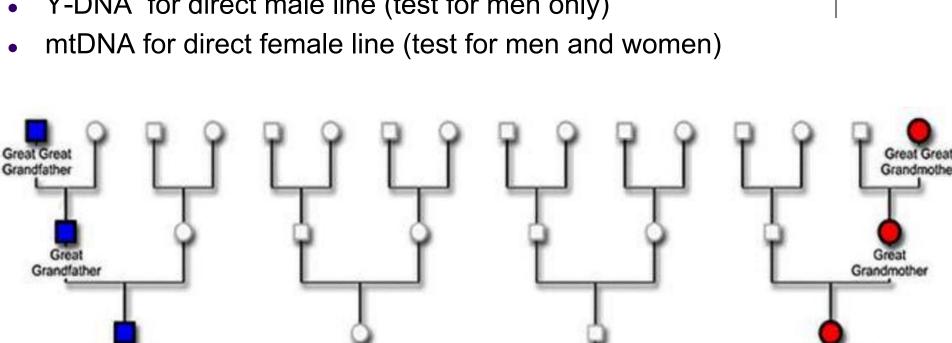


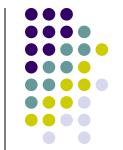


- Three most common types of testing using DNA
 - Y-DNA
 - mtDNA
 - Autosomal DNA (atDNA) (including X-dna)
 - Each tests a different type of dna and they CANNOT be compared to each other! Don't compare apples to oranges!

Y-DNA for direct male line (test for men only)

Grandfather



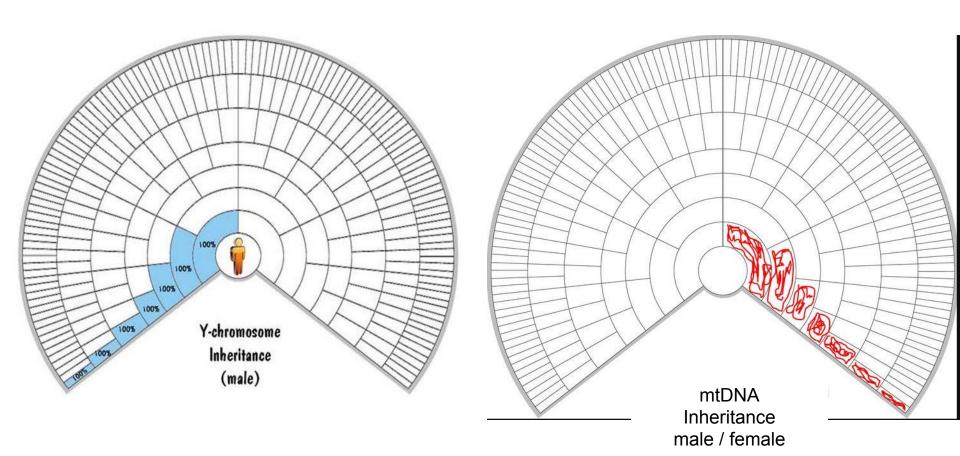


Grandmother



Y-DNA

mtDNA

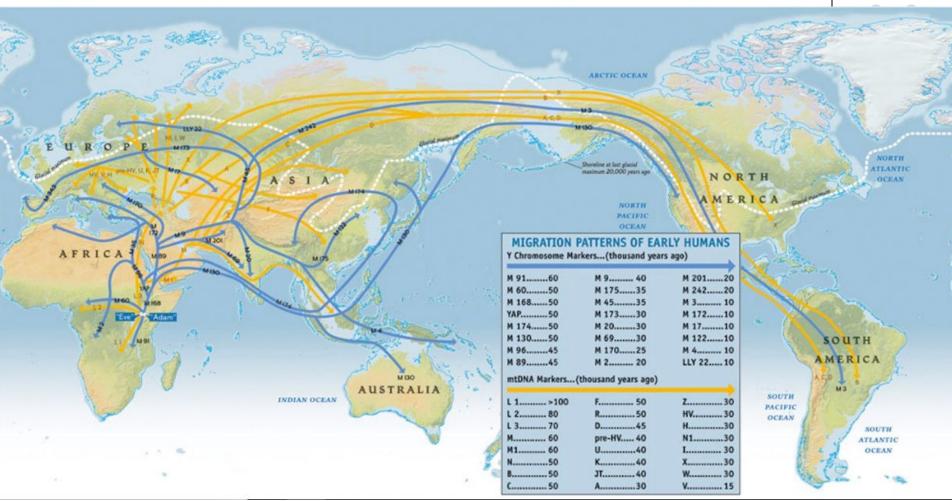




- Y-DNA genetic testing
 - The y chromosome is only passed down from a man to his son.
 - Every man has a y chromosome that has been passed down to him from thousands and thousands of generations of fathers to sons going back into the dawn of humanity (National Genographic Project).
 - Since the start of the use of surnames fathers have tended to pass on their surname along with a y chromosome
 - "Surname" projects have become very popular as people try to link together groups of men with a certain surname.

Genographic Project





- Y-DNA genetic testing
 - Matching y-dna results on top of a similar surname indicates that two individuals almost certainly share a common ancestor.
 - The number of mutations that exist between two "matching" individuals gives a measure of the amount of time going back until the two individuals link up to the same male ancestor whose y chromosome they both inherited.
 - Matching with a person who has a more "proven" family tree can give you a good idea on your own family line (focussed research)
 - Y-dna will only give info on your one direct line back along your father to father ancestry.

- Y-DNA genetic testing
 - Because there are different levels of tesing the y-dna, the number of differences between two individuals must be considered in conjunction with the amount of markers. The number of markers tested can be 12, 25, 37, 67, and 111 markers. For useful results at least 37 markers is recommended.

TMRCA at 90% Confidence for Y-chromosome Tests*

Number of .	is .	Genetic Distance										
Markers	0	1	2	3	4	5	6	7	8	9	10	11+
111	4	7	9	11	14	16	18	20	22	24	26	
67	6	10	13	17	20	23	26	29	32	34		
37	8	12	17	21	24	28						
25	18	29	39	49								
12	48	78	103									

^{*} Data derived using J. D. McDonald's TMRCA Calculator (http://dna-project.clan-donald-usa.org/tmrca.htm)

Probably not related Possibly related	Probably related	Related	Tightly related	Very tightly related
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Y-DNA TiP Report

In comparing Y-DNA 111 marker results, the probability that **Mr. I** a common ancestor within the last...

Porteous and Mr. J. Porteous share

COMPARISON CHART					
Generations	Percentage				
4	12.11%				
8	60.43%				
12	90.12%				
16	98.28%				
20	99.76%				
24	99.97%				

Refine your results with paper trail input

If traditional genealogical records indicate that a common ancestor between you and your match could not have lived in a certain number of past generations, your TiP results can be refined. Note, if you are not sure of this information, you should not change the value of "1" below.

Mr. I Porteous and Mr. J. Porteous did not share a common ancestor in the last 1 generation(s).

Markers 111 ▼ Display every 4 generations. ▼ RECALCULATE

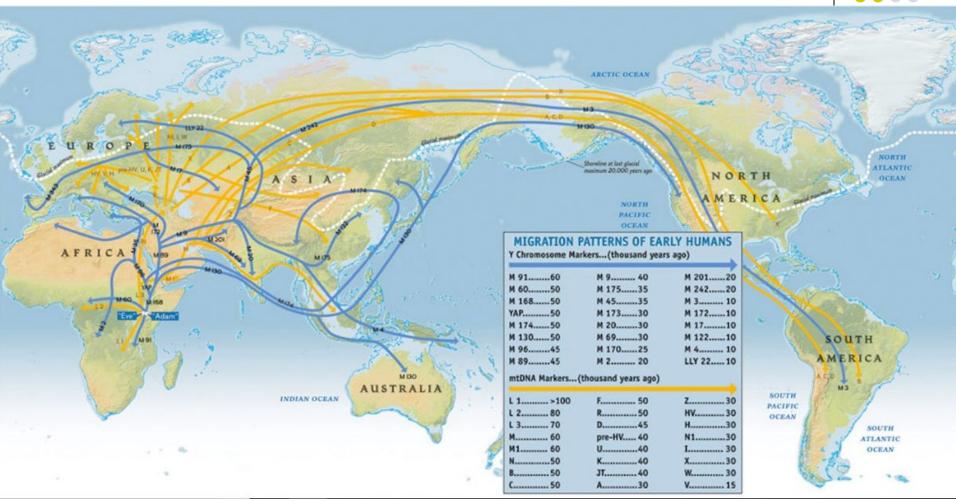
Since each marker has a different mutation rate, identical Genetic Distances will not necessarily yield the same probabilities. In other words, ever though **Mr. J. Porteous** has a Genetic Distance‡ of 6 from **Mr. I Porteous**, someone else with the same Genetic Distance may have different probabilities, because the distance of was prompted by mutations in different markers, with different mutation rates.



- mtDNA genetic testing
 - Mitochondria are part of the make up of every cell in our bodies and help in the generation of energy.
 - They possess their own dna or mtDNA
 - mtDNA is only inherited from your mother

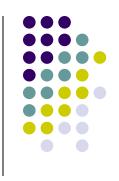
Genographic Project





- mtDNA genetic testing
 - This means that each of us has mtDNA that has been passed on unbroken going back along a mother to child for thousands of years
 - mtDNA projects, similar to y-dna surname projects, are being created to link together people possessing the same mtDNA.
 - The number of mutations between "matching" persons indicates the time back to a common ancestor.
 - Less common than surname projects as mtDNA "matches" will not share any common surname. Getting back to the common female ancestor requires figuring out the name of each successive generation of direct female ancestors.
 - Can be used to confirm "relatedness" if direct female descent from a common female ancestor is suspected.





- Autosomal DNA genetic
 - Unlike the previous two types of testing that can allow connections back hundreds of years in the past, autosomal dna testing is for trying to make a "recent" genetic connection
 - "matches" through this testing can find relatives out to ~ 5th cousins, which would be for people who share 4x great grandparents. For us that's back to ~1750-1800
 - very popular with non-genealogist for their "ethnicity" reports (AncestryDNA commercials). I don't spend much time on this AT ALL!

Genetic "Ethnicity" (Ancestry)



Ethnicity Estimate	Updates 🚯	^
 Ireland and Scotland 	36%	>
France	35%	>
 England, Wales & Northwestern Europe 	29%	>
Migrations		
Saint Lawrence River French Se From your regions: England, Wales & I		>
Montreal & Detroit French Settlers		
See other regions tested	3	350+

Ethnicity Estimate	pdates f	^	
 England, Wales & Northwestern Europe 	38%	>	^
• France	37%	>	
Ireland and Scotland	18%	>	
Germanic Europe	5%	>	
Sweden	2%	>	
Migrations			
Saint Lawrence River French Sett	lers	>	
From your regions: England, Wales & No	rthwes		
Montreal & Detroit French Settlers			

my brother

me

Genetic "Ethnicity" (FTDNA)



European	88% ^
British Isles	53%
West and Centra	l Europe 16%
Southeast Europ	e 12%
Scandinavia	7%
Middle Eastern	10% ~
Trace Results	D ~

European	94% ^
British Isles	54%
Southeast Europe	16%
Scandinavia	16%
East Europe	8%
Middle Eastern	7% ~
Show All	ĺ

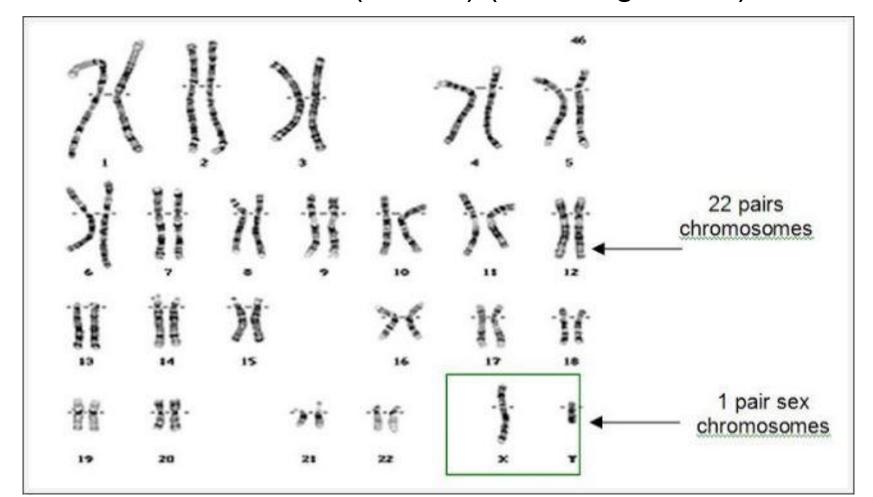
my sister

me

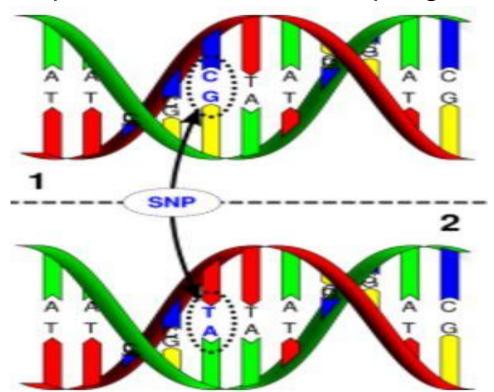




Autosomal DNA (atDNA) (including X-dna)

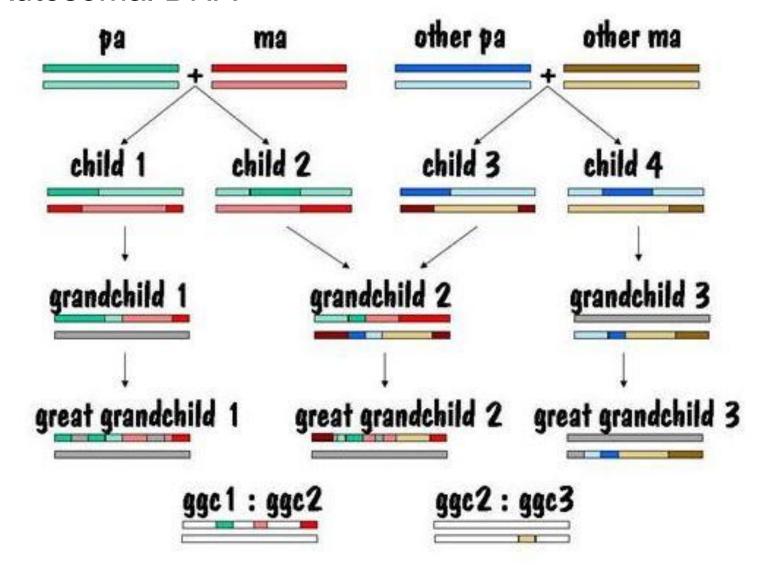


- Autosomal DNA
 - The vast majority of our DNA is identical, however when a mutation occurs or a Single Nucleotide Polymorphism or SNP, it makes that spot on our genome unique to us and to our offspring

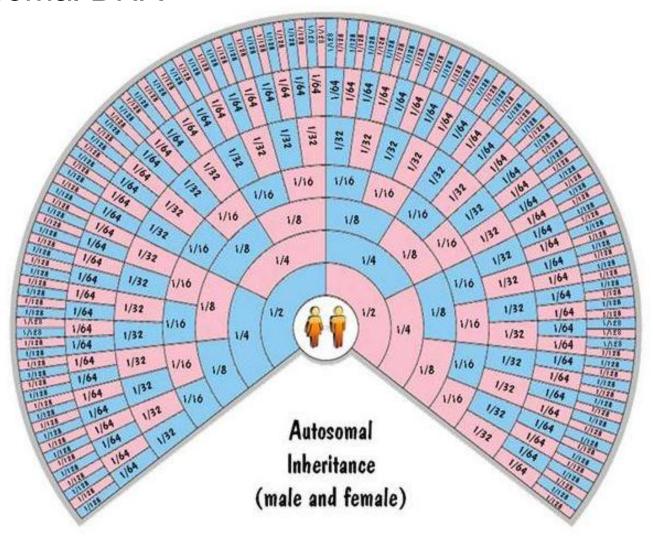




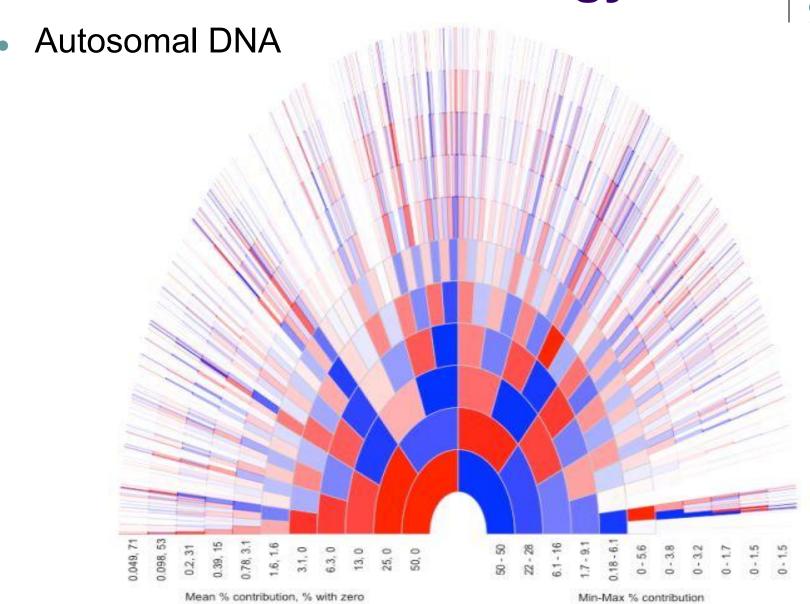
Autosomal DNA



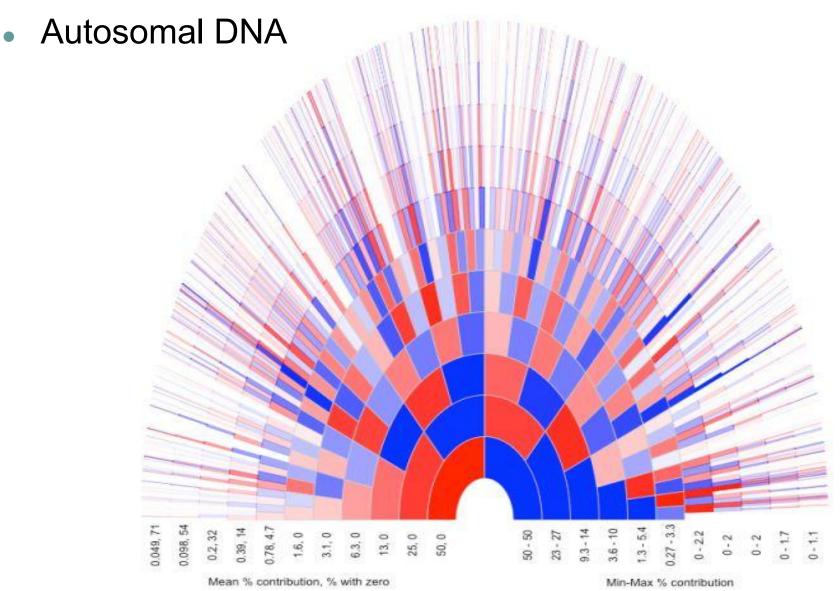
Autosomal DNA



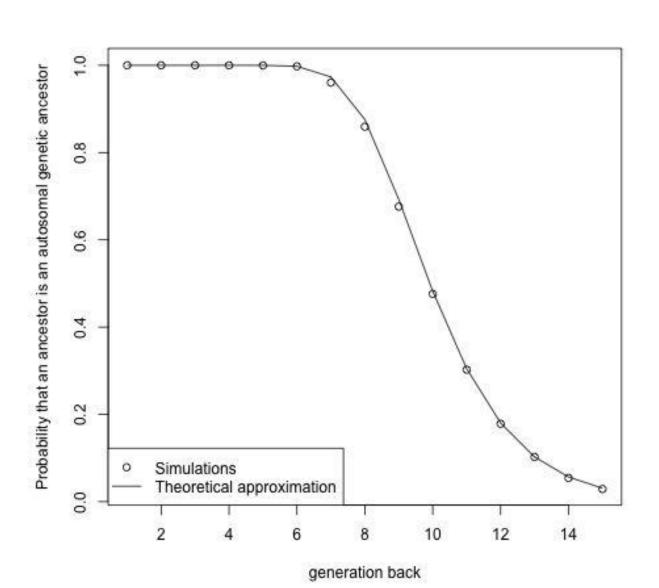






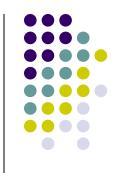


Genetic Genealogy Autosomal DNA





Autosomal DNA



So in any given generation in the past, there are a set of people in your genealogy who we can trace the various sections of your genome back to (lets call them your genetic ancestors). The probability we inherit any autosomal DNA from a **specific** ancestor from 12 generations ago is small (18%), because you have vast numbers of ancestors that far back (e.g. 12 generations ago you have 4096 ancestors) and your genetic ancestors are a very small subset of these people (on average around 700 people).

Reply

Autosomal DNA



Relationship	Match Probability
2 nd cousins or closer	> 99%
3 rd cousin	> 90%
4 th cousin	> 50%
5 th cousin	> 10%
6 th cousin and more distant	remote (typically less than a few percent)

Although it might appear that a genetic match with a distant cousin can't be made, the shear number of distant cousins that we have means that some will come up as a match. You might only have a 0.5% chance of matching an 8th cousins who we share atDNA with, but since we may have upwards of say 40,000 of these out there, that translates to 200 possible 8th cousin matches.

Genetic GenealogyHow we determine relatedness



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	Aunt/Uncle	1st great- aunt/uncle	2nd great- aunt/uncle	3rd great- aunt/uncle	4th great- aunt/uncle	5th great- aunt/uncle	6th great- aunt/uncle	7th great- aunt/uncle
grandparent	Aunt/Uncle	1st cousin	1st cousin once removed	1st cousin twice removed	1st cousin thrice removed	1st cousin four times removed	1st cousin five times removed	1st cousin six times removed	1st cousin seven times removed
1st great- grandparent	1 - 1 - 1 - 1 - 1 - 1 - 1 - 1 - 1 - 1 -	1st cousin once removed	2nd cousin	2nd cousin once removed	2nd cousin twice removed	2nd cousin thrice removed	2nd cousin four times removed	2nd cousin five times removed	2nd cousin six times removed
2nd great- grandparent		1st cousin twice removed	2nd cousin once removed	3rd cousin	3rd cousin once removed	3rd cousin twice removed	3rd cousin thrice removed	3rd cousin four times removed	3rd cousin five times removed
3rd great- grandparent	The second secon	1st cousin thrice removed	2nd cousin twice removed	3rd cousin once removed	4th cousin	4th cousin once removed	4th cousin twice removed	4th cousin thrice removed	4th cousin four times removed
4th great- grandparent		1st cousin four times removed	2nd cousin thrice removed	3rd cousin twice removed	4th cousin once removed	5th cousin	5th cousin once removed	5th cousin twice removed	5th cousin thrice removed
5th great- grandparent	**************************************	1st cousin five times removed	2nd cousin four times removed	3rd cousin thrice removed	4th cousin twice removed	5th cousin once removed	6th cousin	6th cousin once removed	6th cousin twice removed
	Control of The Control of Control	1st cousin six times removed	2nd cousin five times removed	3rd cousin four times removed	4th cousin thrice removed	5th cousin twice removed	6th cousin once removed	7th cousin	7th cousin once removed
7th great- grandparent		1st cousin seven times removed	2nd cousin six times removed	3rd cousin five times removed	4th cousin four times removed	5th cousin thrice removed	6th cousin twice removed	7th cousin once removed	8th cousin

Genetic Genealogy Predicted % of share atDNA

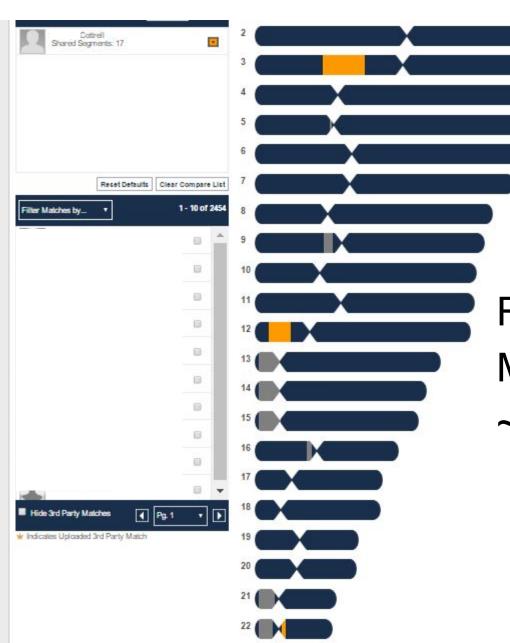


Start	parent	grandparent	1st great- grandparent	2nd great- grandparent	3rd great- grandparent	4th great- grandparent	5th great- grandparent	6th great- grandparent	7th great- grandparent
parent	Sibling 50.000%	25.000%	12.500%	6.250%	3.125%	1.563%	0.781%	0.391%	0.195%
grandparent	25.000%	1st Cousin 12.500%	6.250%	3.125%	1.563%	0.781%	0.391%	0.195%	0.098%
lst great- grandparent	12.500%	6.250%	2 nd Cousin 3.125%	1.563%	0.781%	0.391%	0.195%	0.098%	0.049%
2nd great- grandparent	6.250%	3.125%	1.563%	3rd Cousin 0.781%	0.391%	0.195%	0.098%	0.049%	0.024%
3rd great- grandparent	3.125%	1.563%	0.781%	0.391%	4th Cousin 0.195%	0.098%	0.049%	0.024%	0.012%
4th great- grandparent	1.563%	0.781%	0.391%	0.195%	0.098%	5th Cousin 0.049%	0.024%	0.012%	0.006%
5th great- grandparent	0.781%	0.391%	0.195%	0.098%	0.049%	0.024%	6th Cousin 0.012%	0.006%	0.003%
6th great- grandparent	0.391%	0.195%	0.098%	0.049%	0.024%	0.012%	0.006%	7th Cousin 0.003%	0.002%
7th great- raidparent	hai	red :	at D ₂ N	0.0248	~ 7:40	V. 06%	0.003%	0.002%	8 th Cousin

1



- Example of Genetic Match
 - R. Porteous and M. Cottrell
 - Suspected of being related through a common Lackey ancestor
 - Genetic testing indicated that the two individuals were genetic cousins which pointed to the common Lackey ancestor being in fact correct, making the two ladies 3rd Cousin's once removed.





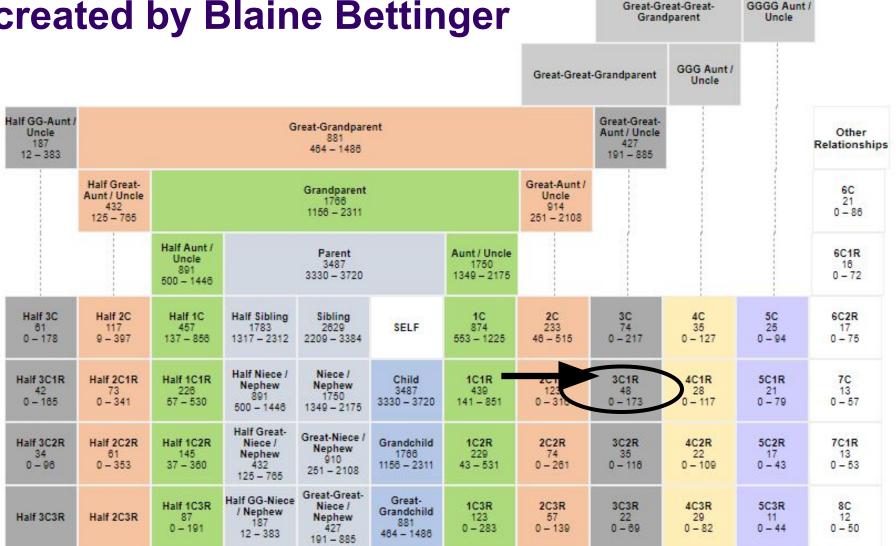
R. Porteous and

M. Cottrell share

~ 50cM and 3 segments of 5cM or greater

Predicted % of share atDNA The Shared CentiMorgan Project created by Blaine Bettinger







- Demonstration of the FTDNA site
 - Chromosome browser
 - "In Common With" tool
 - Surname search tool
 - x-DNA matches



- Demonstration of the Ancestry DNA site
 - searches
 - shared matches
 - trees
 - DNA circles
 - Chrome add ons
 - AncestryDNA Helper
 - DNA Match Labelling

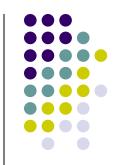
3rd Party Genetic Genealogy AnSites



- www.dnagedcom.com
- www.dnapainter.com
- www.gedmatch.com

Allows comparison between two people regardless of their testing company and regardless of amounts of share atDNA (FTDNA only shows those meeting their definition of "genetic cousins" ie. min 7cM largest segment and min total of 20cM shared).

Genome Mate Pro (GMP) (refer to Facebook group)
 Very powerful tool that uses a database of autosomal raw test results from any of the major test companies to analyse match information.



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